

Fossil Version Control

A User Guide

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Foreword

This book was started in 2010 using a Dual processor Mac 550 MHz PowerPC running Mac OSX 10.5. Current revisions are now done on MacBook Pro (Intel) running MacOS 10.12.1 at 2.5GHz. It was originally done using Fossil version 1.27 and now revisions to the book are using Fossil 1.36. Because of this some of the pictures here will look slightly different in your copy of Fossil but the maintainers have been careful to update the esthetics but not the functions of the various pages. So you will see different, better looking pages but the functions will be the same.

In November 2020, Fossil is up to version 2.14 and there have been extensive changes to the command set, so documentation is longer but follows the same pattern as in earlier versions. Edits now being done on Debian Linux x64 running on an Intel J1900 cpu at 2GHz with 4 processors and 4 GB RAM. Could be done on a Raspberry Pi or Chromebook.

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Chapter 1

Why source control

1.1 What is source control?

A source control system is software that manages the files in a project. A project (typically a software application, but also this book) usually has a number of files. These files in turn are managed by organizing them in directories and sub-directories. At any particular time this set of files in their present edited state make up a working copy of the project at the latest version. If other people are working on the same project you would give them your current working copy to start with. As they find and fix problems, their working copy will become different from the one that you have (it will be in a different state or version). For you to be able to continue where they left off, you will need some mechanism to update your working copy of the files to the latest version available in the team. As soon as you have updated your files, this new version of the project goes through the same cycle again. Most likely the current version will be identified with a code, this is why software has versions and books have editions.

Software developers on large projects with multiple developers could see this cycle and realized they needed a tool to control the changes. With multiple developers sometimes the same file would be edited by two different people changing it in different ways and records of what got changed would be lost. It was hard to bring out a new release of the software and be sure that all the work of all team members to fix bugs and write enhancements were included.

A tool called **Source Code Control System** (SCCS) was developed at Bell Labs in 1972 to track changes in files. It would remember each of the changes made to a file and store comments about why this was done. It also limited who could edit the file so conflicting edits would not be done.

This was important but developers could see more was needed. They needed

to be able to save the state of all the files in a project and give it a name (i.e., Release 3.14). As software projects mature you will have a released version of the software being used as well as bug reports written against it, while the next release of the software is being developed adding new features. The source control system would have to handle what are now called branches. One branch name for example “Version 1” is released but continues to have fixes added to create Version 1.1, 1.2, etc. At the same time you also have another branch which named “Version 2” with new features added under construction.

In 1986 the open source **Concurrent Version Control System** (CVS) was developed. This system could label groups of files and allow multiple branches (i.e. versions) simultaneously. There have been many other systems developed since then, some open source and some proprietary.

Fossil, which was originally released in 2006, is an easy to install version control system that also includes a ticketing system (see section 5.7), a Wiki (see section 7.4.5) and self hosted web server (see section 7.2.15).

1.2 Why version the files?

Why do you want to use a source control system? You won't be able to create files, delete files, or move files between directories at random. Making changes in your code becomes a check list of steps that must be followed carefully.

With all those hassles, why do it? One of the most horrible feelings as a developer is the “It worked yesterday” syndrome. That is, you had code that worked just fine and now it doesn't. If you work on only one document, it is conceivable that you saved a copy under a different name (perhaps with a date) that you could go back to. But if you did not, you doubtless feel helpless at your inability to get back to working code. With a source control system and careful adherence to procedures you can just go back in time and get yesterday's code, or the code from one hour ago, or from last month. After you have done this, starting from known good code, you can figure out what happened.

Having a source control system also gives you the freedom to experiment, “let's try that radical new technique”, and if it doesn't work then it's easy to go back to the previous state.

The rest of this book is a user manual for the Fossil version control system that does code management and much much more. It runs on multiple OS's and is free and open source software (FOSS). It is simple to install as it has only one executable and the repositories it creates are a single file that is easy to back up and are usually only 50 % the size of the original source.

1.2.1 How to get it

If this has interested you then you can get a copy of the Fossil executable here: www.fossil-scm.org/download.html. There are links to Linux, Mac, and Windows executable on this page. The source code is also available if you want or need to compile it yourself. This web site, containing the content you are reading, is self-hosted by Fossil.

1.3 Source control description

This next section is useful if you have not used source control systems before. I will define some of the vocabulary and explain the basic ideas of source control.

1.3.1 Check out systems

When describing the older source control systems, like SCCS, I said it managed the changes for a single file and also prevented multiple people from working on the same file at the same time. This is representative of a whole class of source control systems. In these you have the idea of “checking-out” a file so you can edit it, while becoming the current “owner”. At the same time, while other people using the system can see who is working on the file, they are prevented from touching it. They can get a read-only copy so they can say build software but only the “owner” can edit it. When done editing the “owner” checks it back in, after which anyone else could work on on it. At the same time the system has recorded who had it and the changes made to it.

This system works well in small groups with real time communication. A common problem is that a file is checked out by some one else and you have to make a change in it. In a small group setting, just a shout over the cube wall will solve the problem.

1.3.2 Merge systems

In systems represented by CVS or Subversion the barrier is not getting a file to work on but putting it back under version control. In these systems you pull the source code files to a working directory in your area. Then you edit these files, making necessary changes. When done you commit or check them back into the repository. At this point they are back under version control and the system knows the changes from the last version to this version.

This gets around the problem mentioned above when others are blocked from working on a file. You now have the opposite problem in that more than one person can edit the same file and make changes. This is handled by the check-in process. There only one person at a time may check in a file. That being the case, the system checks the file and if there are changes in the repository file that are *not* in the one to be checked in the check in process stops. The system will ask if

the user wants to merge these changes into his copy, after correcting manually for areas of overlapping change. Once that is done the new version of the file can be checked in.

This type of system is used on large projects like the Linux kernel or other systems where you have a large number of geographically distributed contributors.

1.3.3 Distributed systems

The representatives of two major systems we have described thus far are centralized. That is there is only one repository on a single server. When you get a copy of the files or check in files, it all goes to just one place. These work and can support many, many users. A distributed system is an extension of this where it allows the repositories to be duplicated and has mechanisms to synchronize them.

With a single server, users of the repository must be able to connect to it to get updates and to check in new code. If you are not connected to the server you are stuck and cannot continue working. Distributed systems allow you to have your own copy of the repository to continue working and, when back in communication, to synchronize with the server. This is very useful where people take their work home and cannot access the company network. Each person can have a copy of the repository, continue working, and re-sync upon return to the office.

1.3.4 Common terms

The following is a list of terms I will use when talking about version control or Fossil.

Repository This is the store where the version controlled files are kept. It will be managed by a source control system.

Source control system This is software that manages a group of files by keeping track of changes and allowing multiple users to modify them in a controlled fashion.

Commit in Fossil Store the current set of new and changed files into the repository.

Trunk The main line of code descent in a Fossil repository.

Branch A user defined split in the files served by an SCS. This allow multiple work points on the same repository. Older branches (versions) might have bug fixes applied and newer branches (versions) can have new features added.

Fork In Fossil, an involuntary split in the code path occurs when a file in the repository has changes not in a file to be committed.

Chapter 2

Single user

2.1 Introduction

If you have read this far and are at least persuaded to try, you will want to put one of your projects under software control using Fossil. This chapter is set up to lead you through that task and show you how to adapt your development to using this tool. The assumption is made in this section that you will be the only person using the repository, you are the designer, developer, and maintainer of this project. After you are comfortable using the tool, the next section will show how you use it when you have multiple people working on a project.

2.2 Creating a repository

2.2.1 Introduction

In the spirit of “eating one’s own dog food” we will use this book as the project we are going to manage with Fossil. The book is a directory of Markdown text files and the current working area looks like this:


```
fossilbook3
- content
  - book.md
  - image
    - book
      - fossil.png
  - index.md
  - introduction.md
- makefile
- outline.txt
- pandoc
```

```
- metadata-general.yaml
```

It took just an hour or so to start preliminary research and build the framework. Since that's about all I'm going to do today I want to build a repository and put all this stuff under Fossil control.

2.2.2 Create repository

I have a directory called FOSSIL in which I keep all my repositories, Fossil doesn't care but it helps me to keep them all in one place so I can back them up. First I need to create a new repository for the book. This is done using the command line after I move into the Fossil book directory called `fossilbook3`. (Here I have the `fossilbook3` directory side by side with the `FOSSIL` directory).



```
\$ cd fossilbook3 \$ fossil new ../FOSSIL/FossilBook.fossil
```

I create my repositories with the extension `.fossil`, this will be useful later with the `server` command (see section 7.2.15). When the repository is created, an initial password is assigned with an admin user of "jim" (i.e. me).

2.2.3 Connect repository

The repository is created but is empty and has no connection to the `fossilbook3` directory. The next step is to open the repository to the `fossilbook3` directory with the `open` command.



```
\$ fossil open ../FOSSIL/fossilbook3.fossil
```

This will register a connection to the repository and do an initial checkout into the current directory (which is `fossilbook3` in my case). The `open` command seemingly did nothing, as if no files were created. However, checking with the `status` command shows the repository, the directory it's linked to and that we are hooked to the trunk of the store.



```
\$ fossil status
```

```
repository: /home/jim/FOSSIL/fossilbook3.fossil local-root:
/home/jim/fossilbook3/ config-db: /home/jim/.config/fossil.db checkout:
21e43f604387da163ea84f0d2f73cbaefff0f681 2020-11-23 18:46:21 UTC parent:
2baae1d5fb4865028ba9f4b104f8fed2f9b44094 2020-11-23 12:45:17 UTC tags:
trunk comment: initial empty check-in (user: jim)
```

The `extra` command shows all the files in the directory that are *not* under control of Fossil. In this case that's all of them since we have not checked in anything.



```
\$ fossil extra
```

```
content/introduction.md
```

Note, the status message says `initial empty check-in`. So, Fossil provides us with some basis check-in that is empty. It is the root of your own later check-ins. Note also, you can open the repository more than once. You just need to do this in different working directories. For example, this can be used to work on different branches (see section 4) at the same time.

2.2.4 Add and initial commit

I must now add all the relevant files into the repository with the `add` command. The Fossil `add` is recursive so if I add the top level files it will automatically recurse into the subdirectories and get those files too. Before you do an `add` it pays to tidy up your directory so you don't accidentally add a bunch of transient files (like object files from a compile). It's easy to remove them later but a little tidying before hand can save you some work.



```
\$ fossil add .
```

I simply told fossil to do an `add` of the current directory (`.`) so it got all those files and all the files in the subdirectory. Note the file named `_FOSSIL_` or `.fslckout` (depending on your operating system) that wasn't added to the repository but is located among your working files on disk. This is the tag file that fossil keeps in a directory so it knows what repository it belongs to. Fossil won't add this file since it manages it separately, but everything else is fair game.

One final thing before I can quit for the day, these files have been added or rather they will be added to the repository when I commit them. That must be done and then we can relax and let Fossil manage things.



```
\$ fossil commit -m "Initial Commit"
```

I added a comment text to the commit ("Initial Commit") and it's a good idea to always do this. When later we see the timeline of the commits you will have notes to tell you what was done.

2.2.5 Fossil start up summary

fossil new <name> Creates a new fossil repository.

fossil open <repository> While in a source directory connects this directory to the fossil repository.

fossil add . Will add (recursively) all the files in the current directory and all subdirectories to the repository.

fossil commit -m "Initial Commit" Will put all the currently added files into the repository.

2.3 Set up user interface

One of the surprising features of Fossil is the webserver. This allows it to have a GUI type user interface with no operating system specific code, the GUI is the web browser supplied by your OS. In the previous steps I checked in my project to a Fossil repository, next I have to prepare the web interface for production use.



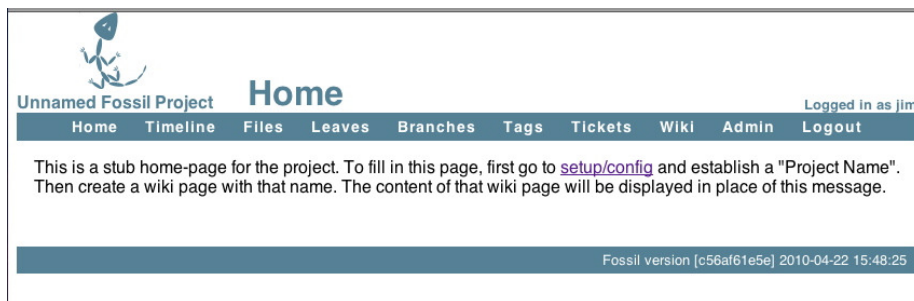
The Fossil web server uses the first available port starting at 8080 instead of the standard port 80 for all HTTP access. When run it will automatically start your Web browser and open the repository home page.

Fossil automatically finds an open port and will give a message on the command line telling you what port number is used. You can still use the `-port` option if you want to control the port number.



```
\$ fossil ui
```

This command shows how it's started. When I do this my browser starts if it is not running already and I am presented with the following home page:



Following the advice on the page I go to `setup/config`. I am going to do the minimum setup that you should do on all projects. As you get familiar with Fossil you will probably have many more things that you will customize for your taste but what follows are the only things you *have* to do.

I have entered in a project name and put in a description, the project name will be the name of the initial Wiki page (see section 7.4.5) and the description is useful for others to see what you are doing here. Then I go to the bottom of the page and pick the `Apply Changes` button.

Next I pick the `Admin` tab (you can see it in the header bar) and the pick `Users` from that page. I am presented with the users and will use this to set the password of the project. See section 6 for details on what else you can customize from the `Admin` page.

As you can see Fossil automatically configures a number of users beyond just the creator. The `anonymous` user you have already seen if you went to the Fossil web site to download the code. This user can view and get code but

cannot commit code. On the right side of the page are the many options you can give to a user, it's worth reading it all when you set up your repository. The important one is me (jim) which has "s" or "Super User Capabilities". This means I can do anything with the repository.

I will now edit the user "Jim" to make sure it has the settings I want. In this case you *must* set the password. Remember way back where Fossil set it during the creation of the repository (Figure[fig:Create-Repository]), it's a very good idea to change this to something you can remember rather than the original random one.

I have put in my contact information (e-mail address) and while you cannot see it I have typed in a password that I will remember. Then I applied the changes.

Now the repository is ready for further work, it's rather bare bones at this point but the most important things are set up.

2.3.1 User interface summary


fossil ui Run in the source directory will start a browser based user interface to fossil.

fossil ui -port <IP port #> Can be used if port 8080 is already in use on your system.

On the first run it is important to configure your project with a name and set the password for yourself.

2.4 Update repository

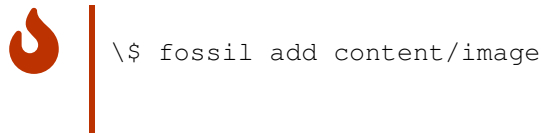
After writing the above section of the book I now have created a some new files and changed some of the existing files in the repository. Before quitting for the day I should add these new files into the repository and commit the changes saving this milestone in the project.



```
| \ $ fossil extra
```

I run `fossil extra` to see these new files. Choose the ones to add and note if there are perhaps any temporary files that should *not* be stored in the repository. I also ran `fossil status`. This shows changes to files that are already in the repository. The only files changed are `content/introduction.md` and `content/single-user.md`.

All I have to do now is add in the directory `content/image` which will add in the image files I want in the repository. Then I commit the changes to the repository and we can move on to other tasks of the day.


 A terminal window showing a red flame icon on the left, a vertical red line, and the command `\$ fossil add content/image` on the right.

After doing this commit I can bring up the Fossil ui (see section 7.2.15) and view the project `Timeline` by picking that tab on the Fossil header. We get this:

```
Timeline placeholder
```

You can see all my check-ins thus far and you can see after the check-in from Figure [fig:Update-for-new] I did another check-in because I missed some changes in the outline. The check-ins are labeled with the first 10 digits of their hash value and these are active links which you can click to view in detail what was changed in that version.

I clicked on the very last check-in (the LEAF) and the display is shown above. There are many things you can do at this point. From the list of changed files you can pick the `diff` link and it will show in text form the changes made in that particular file. The `Other Links` section has a very useful `ZIP Archive`. Clicking this will download a ZIP of this version to your browser. You will find this useful if you want to get a particular version, in fact this is normally how you get a new version of Fossil from <https://fossil-scm.org/>. The `edit` link will be used later to modify a leaf.

2.4.1 Update summary

fossil status will tell you the updated files before you commit.

fossil extra will list files not in the repository.

fossil commit -m "Commit comment" Commits a change (or changes). It is very important to have a descriptive comment on your commit.

2.5 Tickets

Besides managing your code Fossil has a trouble ticket system. This means you can create a ticket for a problem or feature you are going to add to your system then track your progress. Also you can tie the tickets to specific check-ins of your files. For software this is very useful for bug fixes and feature additions. For example you can look for a bug in the ticket list then have it take you to the change that fixed the problem. Then you know exactly what you did and not have to be confused by other changes you might have made.

When you click on `Tickets` in the menu it will bring up this window. You can create a new ticket, look at the list, or generate a new report. Keeping things simple I will just use the `All Tickets` list for now.

Initial Ticket Window placeholder

Picking `New Ticket` I get a form that I fill out like so:

Ticket Form placeholder

Pretty simple actually. You can put as much or as little detail in here as you wish, but remember this stuff might be really vital six weeks or six months from now so think of what you would want to know then.

Note, there is no `Submit` button now. I need to click on `Preview` in order to check how the ticket will be formatted before I can submit (and there are four different ways of formatting tickets). When everything is OK, I press `Submit`. I get this showing what I entered.


Viewing a Ticket placeholder

Finally picking `Tickets` then `All Tickets` I can see my new ticket in the list marked as “Open” and in a distinctive color.

Ticket List with open ticket placeholder

I try, in handling tickets, to have links from ticket to the commit that addressed the problem and a link from the commit back to the offending ticket. This way looking at the ticket I can get to the changes made and from the timeline I can get to the ticket and its resolution. To do this I will make sure and put the 10 digit hash label from the ticket into the check-in comment and put a link in the resolved ticket to the check-in.

Since I have now written the chapter and put in all these images of what to do I can now add in all the new images to the repository and check this in as another completed operation. And I do that like this:



```
 \$ fossil add Images/single-user
```

First I added in all the new image files. I am lazy and just told it to add in all the files in the `single-user` directory. I have previously added some of those like `config-initial.png` but Fossil is smart and knows this and won't add that one twice. Even though it shows it “ADDED”, it really didn't.

The commit line is very important, as you can see I put the 10 digit hash code for the ticket in brackets in the comment. As we will see in the Wiki section (section 7.4.5) this is a link to the ticket, so when viewing the comment in the

timeline or elsewhere you can click the bracketed item and you would go to the ticket.

Now that I have the items checked in I have to close the ticket. I do that by clicking on its link in the ticket list, that will go to the `View Ticket` window as shown in Figure [fig:viewticket]. From there I pick `edit` and fill it in as shown:

figure placeholder

I mark it as “Closed”. If you have code you can mark this as fixed, tested, or a number of other choices. Another very important step, I brought up the timeline and copied the link for the commit I had just done in Figure [fig:checkin]. By doing this my ticket is now cross linked with the commit and the commit has a link back to the ticket.

2.5.1 Ticket summary

- Tickets are a useful way of reminding you what needs done or bugs fixed
- When you commit a change that affects a ticket, put the 10 digit hash label of the ticket into the commit comment surrounded by brackets, e.g. [`<10 digit hash>`] so you can link to the ticket
- When you close the ticket put in the hash label of the commit that fixed it.



The ticket system is widely customizable. See section 6.1.

2.6 Wiki

As we saw in Figure [fig:Starting-Webserver] Fossil has a browser based user interface. In addition to the pages that are built in you can add pages to that web site via a wiki. This allows you to add code descriptions, user manuals, or other documentation. Fossil keeps all that stuff in one place under version control. A wiki is a web site where you can add pages and links from within your browser. You are given an initial page then if you type [`newpage`] this text will turn into a link and if clicked will take you to a new blank page. Remember in Figure [fig:Initial-Webserver-page] that is the initial page for your project and from there you can add new pages. These pages are automatically managed by the Fossil’s version control system. You don’t have to add or commit.

Since I did the setup on the repository (see Figure [fig:Initial-Configuration]) the home page has changed to this:

Home page placeholder

Not very helpful so the in rest of this chapter I will use the Wiki functions of Fossil to make this more useful. If I pick the Wiki item from the menu bar I get:

Wiki page placeholder

These are the controls that let you manage and modify the wiki. In essence, this is the ‘Help’ page of the wiki. Most important for now is the `Formatting rules` link. This link takes you to a page that describes what you can do to format a wiki page. If you just type text on a page it will appear but be formatted by your browser. You can type HTML commands to control this formatting. It’s worth your time to carefully read this page and note what you can and cannot do. The page just lists the valid HTML commands, and if you don’t know what they mean, I would suggest you find a page like this <https://web.stanford.edu/group/csp/cs21/htmlcheatsheet.pdf> and keep it handy.

Besides the HTML markup the most powerful command for the Wiki is `[page]` where it links to a new page. This is how you add pages and how you build your web site of documentation for the repository.



The wiki is customizable. E.g. you may want to show the list of wiki pages instead of the help page when picking the wiki from Fossil’s menu. See section 6.2 for this and for other details.

2.6.1 Wiki formatting

I now begin work. What I want to do is change the home page to be non-empty and also put a link on the home page to the PDF of this book. In Figure [fig:Wiki-controls] I click on the first item, the FossilBook home page. This takes me to the home page again but now I have an “Edit” option. We also have a “History” option so I could look at older versions of the page.

This shows my initial edit and a preview:

The page shows an edit window where I type things I want displayed and at the top is a row of tabs including a “Preview” tab showing how the page will look like. As you can see I typed some simple HTML to make a large and centered title. The body of the text I just typed and as you see the browser fits the text to the screen. You can have multiple paragraphs by just putting blank lines between your text. Next I wanted a bulleted list and this is done by typing two spaces, a `*` then two more spaces. On each of these lines I have a link to a new (not yet created page). If you look I put these in the form `[<new page> | <title>]`. This way I can have a long string that describes the link but have a nice short (no embedded spaces) page name.

OK, I will save my changes and then go to the new pages. I am doing some complicated things here. The first link is to the book PDF. This will be a file I create in the repository. The Pandoc program I’m using creates the PDF. I will do that, save it, and add it to the repository. But I don’t want to link to a static

file, that is I want the most current version of the PDF, the one I save each time I quit for the day. To do this we have to put in a link that looks like this:

```
[http:doc/tip/FossilBook.pdf | Book (pdf) ]
```

This is a special link the Fossil wiki understands, `doc` says this is documentation. `tip` says use the most current version; you could put a version link here. And finally since I am going to put the book PDF at the top level I only need the file name. If it was in a subdirectory I would have to say `doc/tip/subdir/filename`.

The second link is just to a regular page and I can just edit that one just like I did this the main page.

2.6.2 Wiki summary

- Format your text using HTML commands such as `<h1>Title</h1>` for page headings
- Create and link pages using `[<page> | <Link text>]`
- The page designation `http:doc/tip/\<document path relative to repository>` will display any document in the repository that your browser can handle (i.e. pdf, http etc)
- Never click on a link till *after* you have saved the page

Chapter 3

Multiple users

3.1 Introduction

In the previous chapter I went through using Fossil with just one user (me). In this chapter we will get into using it with multiple users. Thanks to Fossil's distributed design once the set up is done using it is not much different than the single user case with Fossil managing automatically the multiple user details.

3.2 Setup

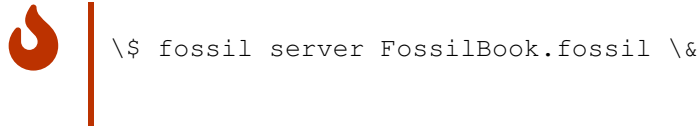
In the previous chapter the Fossil repository was a file on our system and we did commits to it and pulled copies of the source from it. Fossil is a distributed source control system. What this means is that there is a remote server repository in a place that all users can access. Each user has their own "cloned" copy of the repository and Fossil will automatically synchronize the users repository with the remote repository. From a each user's perspective you have your local repository and work with it using the same commands shown in chapter 2. It's just that now Fossil will keep your local repository in sync with the remote repository. This remote repository acts as a server.

3.2.1 Server

I have the FossilBook.fossil repository and now have to put it in place so multiple users can access it. There are two ways, the first is using fossil's built in webserver to host the file and the second is using the operating systems supported web server (if present) and a cgi type access.

3.2.1.1 Self hosted

A self-hosted server is quite simply the easiest way to do it. The downside is that you are responsible for keeping the machine available and the webserver up. That is, don't turn the machine off when you quit for the day or some other user is going to be upset. All I have to do is this:

A terminal window with a red flame icon on the left. The command entered is: `\$ fossil server FossilBook.fossil \&`

```
\$ fossil server FossilBook.fossil \&
```

This is on a UNIX system, the “&” at the end of the command line runs the fossil webserver in the background. If I know this machine has an IP address of 192.168.1.200 then from any other machine in the network I can type into my browser:

```
http://192.168.1.200:8081
```

and I can access the Fossil web server.

As you can see this is simple and works on any system that runs Fossil. As long as you carefully make sure it's always running and available for others this can be a very easy way to make the repository remotely available.

The problems with this method are:

1. If you have multiple repositories you have to use the server not the ui command, have all your repositories in the same directory, and have them all use the extension .fossil.
2. If the machine goes offline (i.e. for OS update) or other reason it might not automatically restart the Fossil servers.
3. Backup of the repositories might not be done.

This method does work, and if you only have one repository and a diligent owner of the remote machine, it will work and work well.

3.2.1.2 Server hosted

If you have a server type machine available (i.e., a Linux or UNIX box) that is running Apache or a Windows machine running IIS you can let it be the webserver for your repository. This has a number of advantages: this machine will be up all the time, it will probably be automatically backed up, and it can easily support multiple Fossil repositories.

I am not going into how to set up the webserver or how to enable CGI (Common Gateway Interface). See the following sites:

- For IIS see e.g. here <https://www.ibm.com/docs/en/cognos-analytics/11.1.0?topic=services-configuring-cgi-gateway-iis-version-7-later> and just ensure that your fossil.exe is in the path to be run by the cgi script.
- For Apache see here <http://httpd.apache.org/docs/2.4/howto/cgi.html> and ensure you know the directory where the CGI scripts are stored.

If you are not in control of the webserver you will need the help of the server admin to enable CGI and to copy your CGI scripts to the correct location.

3.2.1.2.1 CGI Script for hosted server If we assume an Apache server and, in my case, the cgi directory path is /Library/Webserver/CGI-Executables, then we have to write a script of the form:

```
#!/ <Fossil executable location> repository: <Fossil repository location>
```

and put it into the cgi script directory. I have put my Fossil executable into /usr/local/bin and I am putting my Fossil shared repository into /Users/Shared/FOSSIL. My script then becomes:

```
> #!/usr/local/bin/fossil # Put the book repository on the web repository: /Users/SH
```

After making the script I then copy it to the CGI directory and allow anyone to execute it.

```
sudo cp Book.cgi /Library/Webserver/CGI-Executables/Book.cgi
```

3.2.2 Test the setup

If all is in place then I should be able to access the webserver and get to this:

```
Web access to Fossil CGI hosted site - placeholder
```

3.3 User accounts

Serving a repository, either self hosting or the more complicated CGI method gets you to the same place as shown in Figure [fig:Web-access-to]. Now I have to set up user accounts for the other contributors to this book. Remember Fossil has automatically created an Anonymous user (see Figure [fig:User-Configuration]) thus others can access the site in a limited way, that is they can download the book but cannot commit changes. In this case I want to create a new account (Marilyn) that can make changes and commit changes as she is my editor.

To accomplish all this first I have to login by going to the log in page and entering my ID (jim) and my password. Now since I'm super-user I then go back to the User-Configuration page, Figure [fig:User-Configuration] and add a new user:

```
New Editor user - placeholderere
```

Since she is going to be an editor, this will be similar to a developer if we were doing code, so I picked the Developer privilege level. This way she can get the repository, check-in, check-out, and write and update tickets. I also added the attachments since she might need that to put on an image or other comment on actions she is doing. I also gave her a password so her access can be secured.

I could add other users at this point but don't need any others for this project, but you can see how easily this can be done. When you assign the user privileges just read carefully and don't give them any more than you think they need. If they have problems, you can easily modify their account in the future.

3.4 Multiple user operation

With the server set up and the user established the next thing to do is clone the repository. This means to make a copy of the webserver repository on my local machine. Once that is done this local repository uses the same commands and is very much like single the user use discussed in section 2. Fossil will synchronize your local repository with the one on the server.

3.4.1 Cloning

To clone a Fossil repository you have to know four things:

1. It's web address, for our repository. It is `https://fossil-scm.org/FossilBook`
2. Your account name, in my case it's jim
3. Your password (which I'm keeping to myself thank you...)
4. The local name of the repository, in this case I'm calling it `FossilBook.fossil`

You then go to where you want to keep the repository (in my case the FOSSIL directory) and use the clone command:



```
\$ fossil clone https://jim:@fossil-scm.org/FossilBook FossilBook.fossil
```

At this point I can go through the steps outlined in section 2 to set my user password and then open the Fossil repository on a working directory.

Now that I've moved everything to the new cloned repository I do a check in the end of the day which looks like this:



```
|\$ fossil commit -m "Moved to clone repository"
```

As you see the files were committed locally and then the local repository was automatically synchronized with the repository on the server.

3.4.2 Keeping the code in sync

After doing all the setup described above I now have a distributed source control system. My co-worker, Marilyn has also cloned the repository and begun work. She is editing the book correcting my clumsy phrasing and fixing spelling mistakes. She is working independently and on the same files I use. We must use Fossil to prevent us from both modifying the same files but in different ways. Remember Fossil has no file locking, there is nothing to prevent her from editing and changing a file while I work on it.

This is where we both must follow procedures to prevent this sort of problem. Even though she edits files I cannot see the changes till they are committed. Two different versions of the same file won't be a problem till I try to commit with my version and her version is in the current leaf.

There are two problems:

1. Before I do any work I must be sure I have the current versions of all the files.
2. When I commit I must make sure what I am committing has only my changes and is not stepping on changes she has done.

The first is pretty obvious. You make sure you have the latest version before you do anything. We do that with the `fossil update` command. In figure [fig:Cloned-repository-checkin] I had done my latest check in. Before starting any more work I should ensure that Marilyn hasn't checked in something else. I could check the timeline but instead I'll do an update to my repository and source files. When I do the update I specify it should be updated from the trunk. This ensures I get it from the latest and greatest ... and not some branch.



```
|\$ fossil update trunk
```

Ah ha! Marilyn has been at work and updated the book source and pdf. If I check the timeline from the webserver I see she has even documented it:

Now I know I have the current state of the world and I can proceed to add in new sections.



In the default configuration, Fossil operates in ‘auto-sync’ mode (configurable with the `fossil settings` command). This means that Fossil makes sure changes by others are synced from the Fossil server when the `fossil update` command is used. When doing a `commit`, Fossil will automatically push your changes back to the server you cloned from or most recently synced with. When doing an `update`, Fossil will first go to that same server and pull the recent changes to your local repository, then merge them into your local source tree.

3.4.3 Complications

The second problem described in section 3.4.2 is much harder. In this case I have diligently done my `fossil update` and started working. In the mean time Marilyn has also done her update and also started working. Now she is done and checks in her changes. I obviously don’t know this since I am happily working on my changes. What happens next ...



```
\$ fossil commit -m "Commit that might fork"
```


Ah ha, that very thing has happened and Fossil warned me that my copy of the file differs from the master copy. If I had provided the option `-force` to the `commit` then the repository would generate a fork and Marilyn’s future commits would be to her fork and my commits would be to mine. That would not be what we want since I want her to edit my copy of the book.

The next step would be to do as Fossil says and do an `update`. At this point you have to be careful since blindly updating the changed files could overwrite the stuff I’ve just done. So we do a trial update by using the `-n` and `-v` options to say “do a dry run” and show me the results_



```
\$ fossil update -n -v
```

That's a little more than I wanted as you can see almost everything is UNCHANGED but it shows that my file needs a MERGE and fossilbook.pdf needs an UPDATE. This is what I should expect, Marilyn has done edits to the ... file and so have I. So we have to merge the changes. But she has also updated the fossilbook.pdf which I have not. Before we go on if you are running on Linux or UNIX you can simplify this dry run by doing:



```
\$ fossil update -n -v \textbar{} grep -v UNCHANGED
```


By using the pipe and `grep` I can eliminate all those extra UNCHANGED lines.

3.4.4 Fixing the update file

First we fix the easy file, the fossilbook.pdf I can just update by itself so it matches the current repository. It doesn't need to be merged, so I just replace it. Before I do that I have to look at the repository time line

Placeholder for Timeline figure

I see that the current *Leaf* is ...[d44769cc23]... and it is tagged as *trunk*. I want to update the fossilbook.pdf from there. So I say:



```
\$ fossil update trunk fossilbook.pdf
```

and it's done.

3.4.5 Fixing the merge file

We can use the tools built into Fossil. In this case noticing that `commit` will cause a fork, Jim will use the `-force` option to cause the fork and will handle the merge later.



```
\begin{quote}
fossil commit -m "adding some changes of jim"
\end{quote}
```

Now the timeline looks like:

Placeholder for Timeline figure

To remove this fork (i.e. get the changes Marilyn did into the trunk) we use the Fossil `merge` command. We can use the `merge` because `...fossilbook.lyx...` is a text file and the merge markers are designed to work with text files. If it was a binary file we might have to use an external file or copy and paste between the two file versions using the handler program for the file.



```
\begin{quote}
fossil merge a91582b699
\end{quote}
```

Looking at the file (`fossilbook.lyx`) in a text editor (not LyX) we find:

```
> \>>>>>>> BEGIN MERGE CONFLICT
```

Placeholder for now

After the `commit` the timeline shows how the merge brought the fork back into the main *trunk*. Marilyn will then have to update to this new trunk. (See Section [sub:Updating-by-others])

3.5 Sharing other changes than code

3.5.1 Sharable content

When multiple users are working on the code, they may also want to use their local clone of the repository for tickets, the wiki and other stuff. So, while the local code changes are synchronized with the server using the fossil commands `update` and `commit` (or `pull` and `push` if you do it manually), other parts of the repository are not synchronized. If you want to see the current set of tickets or the wiki in your local copy of the repository as well, you need to synchronize those with the command `fossil sync`.

This command will synchronize all sharable content:

- check-ins
- wiki pages
- tickets
- forum posts
- tech notes
- unversioned content (using the additional option `-u`)

This is particularly useful if you cannot work on the remote server for a reason. You can start your local server with `fossil ui`, make changes to e.g. wiki pages and then sync the changes back to the remote server with `fossil sync`.

3.5.2 Synchronize the Fossil configuration

When cloning the remote repository, you also get the current configuration of the repository. This includes e.g. the skin and the ticket setup. When these settings change on the server, they will not automatically be synchronized with your local clone of the repository. So you may end up in a situation where you create a new ticket locally that does not reflect the structure of the tickets on the server because the server configuration has changed! So, if some important configuration changes on the server, you can (and should) pull those changes into your local clone. This is done using the `fossil configuration` command (see the section on the 7.5.4 command for all details). It can be used to change the configuration of the following areas:

- email
- interwiki
- project
- shun
- skin
- ticket
- user
- alias
- subscriber

So, to pull a changed ticket configuration into the local clone, you issue the following command:



```
\begin{quote}
fossil configuration pull ticket
\end{quote}
```

Using the `configuration` command, you can also import and export configurations via text files and even do a kind of merge operation. This makes it possible to edit more complex configurations in an editor of your choice (e.g. when you want to have some syntax highlighting). You can also push your local changes back to the server but that will require administration privileges on the remote server. Last but not least, you may want to do a 'real' sync, synchronizing configuration changes in the local repository with the remote repository. When you use the keyword 'all' for the area, all of the above areas will be synchronized:



```
\begin{quote}  
fossil configuration sync all  
\end{quote}
```

Chapter 4

Forks and branches

4.1 Introduction

This chapter will cover forking and branching in Fossil. **Forking** is where you *unintentionally* create two places to check into a repository. **Branching** is where you *intentionally* do this because you want to maintain two or more versions of the code in the same repository. We illustrated forking and its solutions in section 3.4.3. If, instead of fixing (merging) the file then doing the commit, we forced the commit, Fossil would fork the repository.

Forking is something to avoid because it creates two checkin paths for the code. Thus different users will be on different paths and can check in contradictory changes. Branches on the other hand are forks that you desire. These occur when you want to have two different versions of the code base in development at the same time. A typical scenario is having a production version of the code under maintenance and a development version both served from the same repository. In this case development changes should only be checked into the development branch. Maintenance changes might have to be checked into both.

Instead of using the book repository for these examples we will use a JSON parser program that has a number of files and documentation. This will make it simpler to illustrate branching and tagging.

There is a good discussion of these topics on the Fossil Web site: <https://fossil-scm.org/home/doc/trunk/www/branching.wiki>.

4.2 Forks, branch and merge

In this case the JSON code has just been placed in Fossil and two developers check out copies to work on. Jim wants to fix a number of compiler warnings that appear and Marilyn wants to fix the documentation. In both cases they

proceed as shown in chapter 3. The JSON code has been placed in a distributed location, each of them clones the repository, and opens a working copy of the code.

4.2.1 Marilyn's actions

She looks through the documentation and finds a number of problems and fixes them (the documentation uses LyX and PDFs). When she is satisfied with what she has done, she checks the current version of the documentation in:

4.2.2 Jim's actions

At the same time, Jim gets a working copy of version [6edba5fa8] of the code, puts in a ticket [d23bf4b] as shown in Figure [fig:Marilyn's-work]. After fixing the warnings, Jim is done and goes to commit. He does this *after* Marilyn has done her commit:



```
jim\$ fossil commit -m "[d23bf4b] Remove warnings"
```

At this point Fossil recognizes that Marilyn has changed the repository (she updated the documentation) but Jim does not have these changes because he checked out an earlier version of the code. Jim says he must get his changes in so he does a FORCE to force fossil to accept the commit:




```
jim\$ fossil\{\} commit -m "[d23bf4b] Remove warnings" -f
```

Looking at the timeline Jim sees this:

Not good, there are two Leafs and Marilyn would commit code to her fork and Jim would be committing code to his. So Jim must fix this by merging the code. Jim wants to merge versions [b72e96832e] of Marilyn and his [1beab85441].

4.2.3 Fixing the fork


So Jim who's checked out code is from Leaf [1beab85441] does a merge with Marilyn's leaf [b72e96832e] like so:



```
jim\$ fossil merge b72e96832e
```

As shown the two documentation files are updated, there are no merge conflicts as Jim didn't touch these files and Marilyn didn't touch the code files.


Next Jim does a commit to make this new merged set of files the new trunk. Remember doing the merge shown in Figure [fig:Merge-Operation] just updates your checked out code and does not change the repository till you check it in.



```
jim\$ fossil commit -m "after-merging-in-changes"
```

When we look at the timeline we have a single leaf for future code commits.

The only other thing remaining is that Marilyn does an update before proceeding so her checked out code matches the repository.



```
marilyn\$ fossil update
```

4.2.4 Commands used

fossil merge <fork> Used to merge a fork (specified by hash value) to current check out.

fossil update <version> Used to update current check out to specified version, if version not present use default tag for check out (see `fossil status`)

4.3 Merge without fork

In this case I will show how to merge in code changes from multiple users without causing a fork. In this case Marilyn has put in a BSD license text into all the code files while Jim is adding a help function to the code. In this case both of them put in tickets saying what they are doing but acting independently.


4.3.1 Check in attempt

Marilyn finished first and checks in her changes. Jim builds, tests and tries to check in his code and gets:

```
 | jim\> make
```

4.3.2 Update

The next action Jim takes is to do the update but without doing changes, using the `-n` flag which tells it to just show what's going to happen without making any file changes.

```
 | jim\> fossil update -n
```

This shows some files will be updated, i.e. be replaced by new text from the repository. The `main.c` file will be merged with the version from the repository. That is text from the repository will be mixed with the text from Jim's modified file. Note that it says MERGE meaning the two sets of text are a disjoint set. This means the merge can all be done by Fossil with no human intervention.

Jim can just do the update for real then commit the merged files to make a new leaf. So now we have Marilyn's and Jim changes combined in the latest version.

4.3.3 Commands used

fossil update -n Does a dry run of an update to show what files will be changed.

- UPDATE Implies file will be replaced by repository file - MERGE Implies file will be mixed text from repository and checked out

4.4 Branching

4.4.1 Introduction


We have discussed this before but branching is the intentional splitting of the code in the repository into multiple paths. This will usually be done with production

code where we have maintenance branch and a development branch. The maintenance branch is in use and would get bug fixes based on experience. The development branch would get those changes if applicable plus be modified to add features.

The JSON code parser has been tested and works so will be released to general use. Also we wish to modify it to add support for UTF-8 characters so it matches the JSON standard. The current version just works with ASCII 7 bit characters which is not standard. We wish to split the code into a VER-1.0 branch which is the current code in use and VER-2.0 branch which will add UTF-8 character support.


4.4.2 Branch the repository

Before proceeding we will make sure we have the current trunk code in our check out.



```
|\$ fossil status
```


Seeing that matches the latest leaf in the time line we can proceed to branch the code.



```
|\$ fossil branch new VER-1.0 trunk -bgcolor 0xFFC0FF
```

What was just done? We used the Fossil branch command to create two branches VER-1.0 and VER-2.0 and assigned each of them a color. We can see the timeline is now:

Instead of creating the branch explicitly, we could also have continued changing the code in the trunk. When doing the `commit` we can still decide that those new changes shouldn't go into trunk anyway but into a new branch. For this, we just need to specify the new branch with the `commit` command:



```
|\$ fossil commit -branch VER-1.0 -branchcolor 0xFFC0FF
```

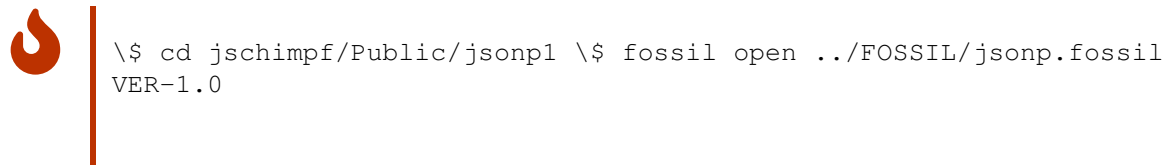
4.4.3 Color setup

As you see above the two branches have different colors in the timeline. This was due to the `-bgcolor` option added when we created each branch. (See Figure [fig:Branch-commands]). But we want this color to appear on subsequent checkins of each of these branches. To make that happen we have to set the options using the UI and picking a particular leaf on the timeline.

Under the 'Background Color' section I have checked 'Propagate color to descendants' so future checkins will have the same color.

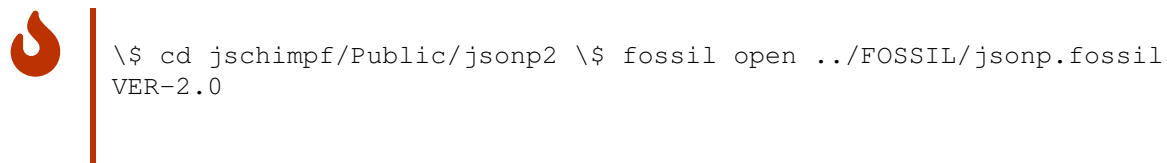
4.4.4 Check out the branches

Now the the repository is branched we can check out the two sets of code into different directories. We create `jsonp1` and `jsonp2` and proceed to open the different branches into them.



```
\$ cd jschimpf/Public/jsonp1 \$ fossil open ../FOSSIL/jsonp.fossil  
VER-1.0
```

Checking out VER-2.0 in the same way



```
\$ cd jschimpf/Public/jsonp2 \$ fossil open ../FOSSIL/jsonp.fossil  
VER-2.0
```

Notice on both of these the tags show which branch we are attached to.

4.4.5 Correcting errors in both

After doing this work I found that the `main.c` file had a warning about an unused variable. I wanted to correct this in both branches. At this point all the files in both branches are the same so correcting the file in either branch and copying it to the other is possible. I put in a ticket for the change and edit `main.c`. I copy it to both checkouts for the both branches and then check both in.



```
\$ cd jschimpf/Public/jsonp1 \$ fossil commit -m "{}2795e6c74d{} Fix
unused variable"
```

Now the timeline looks like this:

4.4.6 Private branches

Sometimes you may want to create a branch but keep the changes in that branch private. E.g. , this may be some experimental code you do not want to share with the other users (yet). To do this, you just create a branch and declare it 'private'. This means, the branch will stay in your local copy of the repository but it will not be synchronized with the remote repository and thus not distributed to others during a commit:



```
\$ fossil commit -branch myTests-123 -private"
```

Of course, you can do the same with the `fossil branch` command. It also has the `--private` option. If you at some time decide to make your branch public, you can do so with the 7.3.17 command.



You can associate wiki pages with branches using a name for the wiki page starting with `branch/`. Note though, although the code is private and not shared, the associated wiki page will be synchronized. Wiki pages are separate from all file-level content and do not take part in branching (or merging etc.). So, the wiki page is *associated* with that private branch but is not *a part of* that branch (in terms of the underlying artifact mechanism).

Source: Fossil forum post

4.4.7 Commands used

fossil branch Used to generate a branch of the repository. The command can optionally color the branch in the display.

fossil commit -branch Committing a new version into a new branch

Chapter 5

The Fossil user interface

Fossils graphical user interface is a web site directly served from the repository. It allows to inspect the history, documentation, tickets and other information on the repository. This interface is wah tusers will likely see in the first place when they want to learn about your software project.

5.1 fossil ui

{» ... about the 'fossil ui' command with reference to commands.md ... «}

5.2 Home

5.3 The timeline

The timeline presents a graphical overview of the development of your project. It can list all kinds of events that happened within the project:

- Check-ins
- Tickets (and a list of new tickets)
- Wiki
- Tech notes
- Tags

...

5.3.1 Check-ins

Each check-in is presented as one point on the timeline.

...

5.4 File

5.4.1 The timeline of a single file

You can also display the timeline of a single file (as opposed to a check-in with multiple files). There are three possibilities to get there.

1. Navigating to the file, starting from the 'File' menu.

Browse to desired file. Click on the file link. you will see the content of the file if it is a text file. Below the main menu (on top), you will see a line starting with the word 'File' and followed by the file name (as a link) and the check-in context. Click on that file name link. This brings you to the `/info` page for this file showing the timeline for this single file.

2. Starting from the full timeline

Click on the check-in link (the hash value) in the timeline. You see some metadata (a section called 'Overview'), the timeline context (a section called 'Context') and below the diffs for files affected by this check-in (a section called 'Changes'). In that last section you will see some lines saying 'Modified ...' with a linked file name. Click on that link and you will get to the page showing the timeline for this single file.

There also is the possibility to see the history of all files in a specific directory (recursively). For this, go to the 'File' section, ensure you are in 'Flat View' (not a tree view that can expand/collapse) Browse to desired directory and then click on the 'History' link on second-level menu. You will see a timeline of check-ins that affected any file within that directory. You will, however, not get a list of the files affected during each check-in. For this, you need to click on the individual check-ins and then look into the 'Changes' section of that check-in.

5.5 Branches

5.6 Tags

5.7 Tickets

5.8 Wiki

5.9 Admin

5.10 Login

Chapter 6

Fossil configuration

Fossil can be configured or customized to your own needs in a variety of ways. This section covers the configuration of some aspects you can change via the Fossil `Admin` menu in the browser interface of Fossil.

6.1 Ticket configuration

6.2 Wiki configuration

Picking the Wiki item from the Fossil menu bar shows the wiki help page by default. You can change that. You may want to show the list of existing wiki pages instead so you can go to every wiki page from here. For this, you need to modify the Fossil skin. Pick `Admin` in the Fossil menu bar and click on 'Skins'. Follow the outlined procedure to initialize the skin editing process (steps 1 to 3; see section 6.4 for details). Then, edit the 'Header' (in Step 4). This will bring up the HTML and TH1 code for the Fossil menu (see section 9 for details on the TH1 language). The only thing you need to do is to change `\wiki` to `\wcontent` in the part of the code that handles showing the wiki menu item. After applying this change, you will need to end the skin editing by doing steps 7-5.

6.3 Users configuration

6.4 Skins configuration

6.5 Configure email notifications

Chapter 7

Fossil commands

7.1 Introduction

This section will go through the various Fossil command line commands. This will be divided into sections, the first will detail the must know commands. These are the ones you will be using all the time and will probably have memorized in short order. The other commands will be divided into Maintenance, Advanced, and Miscellaneous. These you will probably be checking as reference before use.

The most important command is help. You can always type fossil help at the command line and it will list out all the commands it has. Then typing fossil help <command> will print out the detailed information on that command. You always have that as your reference. This section of the book will try to supplement the built in help with some examples and further explanation of what a command does. All of the commands will be placed in the index for easy searching

NOTE: Fossil is a moving target, commands might be added and others removed future versions. Type fossil help on your version to get the latest list. The following applies to the fossil I used when I wrote this and your version might be different.

7.2 Basic

7.2.1 help

This command is used to dump the current command set and version of Fossil. It can also be used in the form fossil help <command> to get further information on any command.

Actually this will give you only a subset of the help commands, limited to the commands that are used most often. If you want to see all commands available then issue the `fossil help -all` command. Between versions you will see changes as to what is included in the help sub-set.

An example of using the help function to get further information about a particular command:

```
$ fossil help
Usage: fossil help TOPIC
Try "fossil help help" or "fossil help -a" for more options
Frequently used commands:
add          cat          extras         merge         rm             ui
addremove   changes   finfo         mv            settings      undo
all         clean     gdiff        open          sql           unversioned
amend       clone     grep         pull          stash         update
annotate    commit    help         push          status        version
bisect      dbstat   info         rebuild       sync
blame       delete   init         remote        tag
branch      diff     ls           revert        timeline
This is fossil version 2.14 [939a13d94f] 2020-11-20 19:28:05 UTC
```

Adding the option `-a` (or `-all`) to the help command results in a listing of ALL the available commands:

```
$ fossil help -a
3-way-merge  clean          hook           rebuild        stash
add           clone          http           reconstruct    status
addremove    close          import         redo           sync
alerts       co             info           remote         tag
all          commit        init           remote-url     tarball
amend        configuration  interwiki     rename         ticket
annotate     dbstat        leaves        reparent       timeline
artifact     deconstruct   login-group   revert          tls-config
attachment   delete        ls            rm             touch
backoffice   descendants    md5sum        rss            ui
backup       diff          merge         scrub          undo
bisect       export        mv            search         unpublished
blame        extras        new           server         unset
branch       finfo         open          settings      unversioned
bundle       forget        pikchr        shasum         update
cache        fts-config    pop3d         sha3sum        user
cat          gdiff        praise        shell          uv
cgi          git           publish       smtpd          version
changes      grep         pull          sql            whatis
checkout     hash-policy   purge         sqlar          wiki
ci           help          push          sqlite3        zip
```


Use it to get further information about a particular command:

```
$ fossil help help
Usage: fossil help [OPTIONS] [TOPIC]
```

Display information on how to use TOPIC, which may be a command, webpage, or setting. Webpage names begin with "/". If TOPIC is omitted, a list of topics is returned.

The following options can be used when TOPIC is omitted:

```
-a|--all           List both common and auxiliary commands
-o|--options      List command-line options common to all commands
-s|--setting      List setting names
-t|--test         List unsupported "test" commands
-x|--aux          List only auxiliary commands
-w|--www          List all web pages
-f|--full         List full set of commands (including auxiliary
                  and unsupported "test" commands), options,
                  settings, and web pages
-e|--everything   List all help on all topics
```

These options can be used when TOPIC is present:

```
-h|--html          Format output as HTML rather than plain text
-c|--commands     Restrict TOPIC search to commands
```

As you can see from the description, `fossil help -a` doesn't even list all possibilities. With the `-f` option, you get even more!

7.2.2 add

The `add` command is used to add files into a repository. It is recursive and will pull in all files in subdirectories of the current. Fossil will not overwrite any of the files already present in the repository so it is safe to add all the files at any time. Only new files will be added.

```
$ fossil help add
Usage: fossil add ?OPTIONS? FILE1 ?FILE2 ...?
```

Make arrangements to add one or more files or directories to the current checkout at the next commit.

When adding files or directories recursively, filenames that begin with "." are excluded by default. To include such files, add the `--dotfiles` option to the command-line.

The `--ignore` and `--clean` options are comma-separated lists of glob patterns for files to be excluded. Example: `'*.o,*.obj,*.exe'` If the `--ignore` option does not appear on the command line then the "ignore-glob" setting is used. If the `--clean` option does not appear on the command line then the "clean-glob" setting is used.

If files are attempted to be added explicitly on the command line which match "ignore-glob", a confirmation is asked first. This can be prevented using the `-f|--force` option.

The `--case-sensitive` option determines whether or not filenames should be treated case sensitive or not. If the option is not given, the default depends on the global setting, or the operating system default, if not set.

Options:

<code>--case-sensitive</code> BOOL	Override the case-sensitive setting.
<code>--dotfiles</code>	include files beginning with a dot (".")
<code>-f --force</code>	Add files without prompting
<code>--ignore</code> CSG	Ignore unmanaged files matching patterns from the Comma Separated Glob (CSG) pattern list
<code>--clean</code> CSG	Also ignore files matching patterns from the Comma Separated Glob (CSG) list
<code>--reset</code>	Reset the ADDED state of a checkout, such that all newly-added (but not yet committed) files are no longer added. No flags other than <code>--verbose</code> and <code>--dry-run</code> may be used with <code>--reset</code> .

The following options are only valid with `--reset`:

<code>-v --verbose</code>	Outputs information about each <code>--reset</code> file.
<code>-n --dry-run</code>	Display instead of run actions.

See also: `addremove`, `rm`

Typing:

```
fossil add .
```

will add all files in the current directory and subdirectories.

Note none of these files are put in the repository until a commit is done.

7.2.3 `rm` or `del`

The `rm` command is used to remove files from the repository. The file is not deleted from the file system but it will be dropped from the repository on the

next commit. This file will still be available in earlier versions of the repository but not in later ones.

```
$ fossil help rm
Usage: fossil rm|delete|forget FILE1 ?FILE2 ...?
```

Remove one or more files or directories from the repository.

The 'rm' and 'delete' commands do NOT normally remove the files from disk. They just mark the files as no longer being part of the project. In other words, future changes to the named files will not be versioned. However, the default behavior of this command may be overridden via the command line options listed below and/or the 'mv-rm-files' setting.

The 'forget' command never removes files from disk, even when the command line options and/or the 'mv-rm-files' setting would otherwise require it to do so.

WARNING: If the "--hard" option is specified -OR- the "mv-rm-files" setting is non-zero, files WILL BE removed from disk as well. This does NOT apply to the 'forget' command.

Options:

--soft	Skip removing files from the checkout. This supersedes the --hard option.
--hard	Remove files from the checkout.
--case-sensitive BOOL	Override the case-sensitive setting.
-n --dry-run	If given, display instead of run actions.
--reset	Reset the DELETED state of a checkout, such that all newly-rm'd (but not yet committed) files are no longer removed. No flags other than --verbose or --dry-run may be used with --reset.
--verbose -v	Outputs information about each --reset file. Only usable with --reset.

See also: addremove, add

You can delete groups of files by using wild-cards in their names. Thus if I had a group of files like com_tr.c, com_rx.c and com_mgmt.c I could remove them all with:

```
fossil rm com_*.c
```

By running a "fossil status" you can see what files will be deleted on the next commit.

7.2.4 rename or mv

This command is used to rename a file in the repository. This does not rename files on disk so is usually used after you have renamed files on the disk then want to change this in the repository.

```
$ fossil help rename
Usage: fossil mv|rename OLDNAME NEWNAME
      or: fossil mv|rename OLDNAME... DIR
```

Move or rename one or more files or directories within the repository tree. You can either rename a file or directory or move it to another subdirectory.

The 'mv' command does NOT normally rename or move the files on disk. This command merely records the fact that file names have changed so that appropriate notations can be made at the next commit. However, the default behavior of this command may be overridden via command line options listed below and/or the 'mv-rm-files' setting.

The 'rename' command never renames or moves files on disk, even when the command line options and/or the 'mv-rm-files' setting would otherwise require it to do so.

WARNING: If the "--hard" option is specified -OR- the "mv-rm-files" setting is non-zero, files WILL BE renamed or moved on disk as well. This does NOT apply to the 'rename' command.

Options:

--soft	Skip moving files within the checkout. This supersedes the --hard option.
--hard	Move files within the checkout.
--case-sensitive BOOL	Override the case-sensitive setting.
-n --dry-run	If given, display instead of run actions.

See also: changes, status

Just like add or rm you can use wild cards in the names and rename groups of files. Like them "fossil status" will show you the current state.

7.2.5 status

The status command is used to show you the current state of your files relative to the repository. It will show you files added, deleted, and changed. This is only the condition of files that are already in the repository or under control of fossil. It also shows from where in the timeline you are checked out and where your repository is kept.

```
$ fossil help status
Usage: fossil changes|status ?OPTIONS? ?PATHS ...?
```

Report the change status of files in the current checkout. If one or more PATHS are specified, only changes among the named files and directories are reported. Directories are searched recursively.

... [snipped]

See also: extras, ls

```
$ fossil status
repository:  /home/chris/repos/fossilbook.fossil
local-root:  /home/chris/fossilbook/
config-db:   /home/chris/.config/fossil.db
checkout:    516e309ealc86132e3a9be313e8331ec91f255ee 2020-11-21 16:26:12 UTC
parent:      030a38fe4b45b7088faa635982037d2eb05143b6 2020-11-21 16:23:34 UTC
tags:        trunk
comment:     Use .md file extension in book.md links. (user: chris)
EDITED      content/introduction.md
EDITED      content/rights.md
```

The listing above shows where my cloned copy of the repository is kept, where I am working, and the tags show me that I am checked out of the trunk. Finally it shows the status of the files: I am working on two of them.

7.2.6 changes

This lists the changed files like status but shows other information that status does not.

```
$ fossil help changes
Usage: fossil changes|status ?OPTIONS? ?PATHS ...?
```

Report the change status of files in the current checkout. If one or more PATHS are specified, only changes among the named files and directories are reported. Directories are searched recursively.

The status command is similar to the changes command, except it lacks several of the options supported by changes and it has its own header and footer information. The header information is a subset of that shown by the info command, and the footer shows if there are any forks. Change type classification is always enabled for the status command.

Each line of output is the name of a changed file, with paths shown according to the "relative-paths" setting, unless overridden by the --abs-paths or --rel-paths options.

By default, all changed files are selected for display. This behavior can be overridden by using one or more filter options (listed below), in which case only files with the specified change type(s) are shown. As a special case, the `--no-merge` option does not inhibit this default. This default shows exactly the set of changes that would be checked in by the `commit` command.

If no filter options are used, or if the `--merge` option is used, the artifact hash of each merge contributor check-in version is displayed at the end of the report. The `--no-merge` option is useful to display the default set of changed files without the merge contributors.

If change type classification is enabled, each output line starts with a code describing the file's change type, e.g. `EDITED` or `RENAMED`. It is enabled by default unless exactly one change type is selected. For the purposes of determining the default, `--changed` counts as selecting one change type. The default can be overridden by the `--classify` or `--no-classify` options.

`--edited` and `--updated` produce disjoint sets. `--updated` shows a file only when it is identical to that of its merge contributor, and the change type classification is `UPDATED_BY_MERGE` or `UPDATED_BY_INTEGRATE`. If the file had to be merged with any other changes, it is considered to be merged or conflicted and therefore will be shown by `--edited`, not `--updated`, with types `EDITED` or `CONFLICT`. The `--changed` option can be used to display the union of `--edited` and `--updated`.

`--differ` is so named because it lists all the differences between the checked-out version and the checkout directory. In addition to the default changes (excluding `--merge`), it lists extra files which (if `ignore-glob` is set correctly) may be worth adding. Prior to doing a commit, it is good practice to check `--differ` to see not only which changes would be committed but also if any files should be added.

If both `--merge` and `--no-merge` are used, `--no-merge` has priority. The same is true of `--classify` and `--no-classify`.

The `"fossil changes --extra"` command is equivalent to `"fossil extras"`.

General options:

<code>--abs-paths</code>	Display absolute pathnames.
<code>--rel-paths</code>	Display pathnames relative to the current working directory.
<code>--hash</code>	Verify file status using hashing rather than relying on file <code>mtimes</code> .

```

--case-sensitive BOOL  Override case-sensitive setting.
--dotfiles            Include unmanaged files beginning with a dot.
--ignore <CSG>       Ignore unmanaged files matching CSG glob patterns.

```

Options specific to the changes command:

```

--header             Identify the repository if report is non-empty.
-v|--verbose        Say "(none)" if the change report is empty.
--classify          Start each line with the file's change type.
--no-classify       Do not print file change types.

```

Filter options:

```

--edited            Display edited, merged, and conflicted files.
--updated          Display files updated by merge/integrate.
--changed          Combination of the above two options.
--missing          Display missing files.
--added            Display added files.
--deleted          Display deleted files.
--renamed          Display renamed files.
--conflict         Display files having merge conflicts.
--meta            Display files with metadata changes.
--unchanged        Display unchanged files.
--all              Display all managed files, i.e. all of the above.
--extra           Display unmanaged files.
--differ           Display modified and extra files.
--merge           Display merge contributors.
--no-merge        Do not display merge contributors.

```

See also: extras, ls

7.2.7 extras

The extras command is used to find files you have added to your working directory but are not yet under Fossil control. This is important because if you move your working directory or others attempt to use the repository they won't have these files.

```

$ fossil help extras
Usage: fossil extras ?OPTIONS? ?PATH1 ...?

```

Print a list of all files in the source tree that are not part of the current checkout. See also the "clean" command. If paths are specified, only files in the given directories will be listed.

Files and subdirectories whose names begin with "." are normally ignored but can be included by adding the --dotfiles option.

Files whose names match any of the glob patterns in the "ignore-glob" setting are ignored. This setting can be overridden by the `--ignore` option, whose CSG argument is a comma-separated list of glob patterns.

Pathnames are displayed according to the "relative-paths" setting, unless overridden by the `--abs-paths` or `--rel-paths` options.

Options:

<code>--abs-paths</code>	Display absolute pathnames.
<code>--case-sensitive</code> BOOL	Override case-sensitive setting
<code>--dotfiles</code>	Include files beginning with a dot (".")
<code>--header</code>	Identify the repository if there are extras
<code>--ignore</code> CSG	Ignore files matching patterns from the argument
<code>--rel-paths</code>	Display pathnames relative to the current working directory.

See also: `changes`, `clean`, `status`

The `--dotfiles` option shows you any files starting with "." that are not under Fossil control. This would be important if you need those files in your repository. The last option `--ignore` allows you to ignore certain files you know don't belong in the repository. In the earlier version of this repository, when Lyx was the editor of choice, there was a file called `fossilbook.lyx~` that is a LyX backup file that was not wanted, as it is temporary. So one can say

```
fossil extra -{}-ignore \*.lyx~
```

and only get:

```
$ fossil extra -{}-ignore \*.lyx\~
```

```
image/basic/help.png
```

instead of:

```
$ fossil extra
```

```
image/basic/help1.png
```

```
fossilbook.lyx~
```

7.2.8 revert

The `revert` command is used to take a file back to the value in the repository. This is useful when you make a error in editing or other mistake.

```
$ fossil help revert
```

```
Usage: fossil revert ?OPTIONS? ?FILE ...?
```

Revert to the current repository version of FILE, or to

the baseline VERSION specified with -r flag.

If FILE was part of a rename operation, both the original file and the renamed file are reverted.

Using a directory name for any of the FILE arguments is the same as using every subdirectory and file beneath that directory.

Revert all files if no file name is provided.

If a file is reverted accidentally, it can be restored using the "fossil undo" command.

Options:

```
-r|--revision VERSION    Revert given FILE(s) back to given
                        VERSION
```

See also: redo, undo, checkout, update

With no parameters it will revert the file to the current revision, see Figure [fig:status-run]. The -r option allows you to pick any revision from the time line.

7.2.9 update

The update option will update a file or files to match the repository. With multiple users it should be done before you start working on any files. This ensures you have the latest version of all the files.

```
$ fossil help update
```

```
Usage: fossil update ?OPTIONS? ?VERSION? ?FILES...?
```

Change the version of the current checkout to VERSION. Any uncommitted changes are retained and applied to the new checkout.

The VERSION argument can be a specific version or tag or branch name. If the VERSION argument is omitted, then the leaf of the subtree that begins at the current version is used, if there is only a single leaf. VERSION can also be "current" to select the leaf of the current version or "latest" to select the most recent check-in.

If one or more FILES are listed after the VERSION then only the named files are candidates to be updated, and any updates to them will be treated as edits to the current version. Using a directory name for one of the FILES arguments is the same as using every subdirectory and file beneath that directory.

If FILES is omitted, all files in the current checkout are subject to being updated and the version of the current checkout is changed to VERSION. Any uncommitted changes are retained and applied to the new checkout.

The `-n` or `--dry-run` option causes this command to do a "dry run". It prints out what would have happened but does not actually make any changes to the current checkout or the repository.

The `-v` or `--verbose` option prints status information about unchanged files in addition to those file that actually do change.

Options:

<code>--case-sensitive</code> BOOL	Override case-sensitive setting
<code>--debug</code>	Print debug information on stdout
<code>--latest</code>	Acceptable in place of VERSION, update to latest version
<code>--force-missing</code>	Force update if missing content after sync
<code>-n --dry-run</code>	If given, display instead of run actions
<code>-v --verbose</code>	Print status information about all files
<code>-W --width</code> WIDTH	Width of lines (default is to auto-detect). Must be more than 20 or 0 (= no limit, resulting in a single line per entry).
<code>--setmtime</code>	Set timestamps of all files to match their SCM-side times (the timestamp of the last checkin which modified them).
<code>-K --keep-merge-files</code>	On merge conflict, retain the temporary files used for merging, named <code>*-baseline</code> , <code>*-original</code> , and <code>*-merge</code> .

See also: `revert`

Update has a number of options, first you can tie the update to a particular version, if not picked then it just uses the latest. Second it can work on a single files or many files at once. That is you could say

```
$ fossil update \*.c
```

and it would update all C files.

Since this is a rather large set of changes it has a special "dry run" mode. If you add `-n` on the command it will just print out what will be done but not do it. This is very useful to do this trial if you are unsure what might happen. The `-v` command (which can be used with `-n` or alone) prints out the action for each file even if it does nothing.



The `update` command can be used to do a kind of cherrypicking. When you have a branch (or just some other version) and want to only merge a single file of that branch/version into the current checkout, you can do `fossil update other_branch other_file`. This will bring a copy of that other file into the current branch/version. However, this will not leave any trace in the timeline. Further, you will have to `fossil add` that file specifically yourself so it gets committed in the next `fossil commit`.

7.2.10 checkout or co

This command is similar to `update`.

```
$ fossil help checkout
Usage: fossil checkout ?VERSION | --latest? ?OPTIONS?
      or: fossil co ?VERSION | --latest? ?OPTIONS?
```

Check out a version specified on the command-line. This command will abort if there are edited files in the current checkout unless the `--force` option appears on the command-line. The `--keep` option leaves files on disk unchanged, except the manifest and manifest.uuid files.

The `--latest` flag can be used in place of `VERSION` to checkout the latest version in the repository.

Options:

<code>--force</code>	Ignore edited files in the current checkout
<code>--keep</code>	Only update the manifest and manifest.uuid files
<code>--force-missing</code>	Force checkout even if content is missing
<code>--setmtime</code>	Set timestamps of all files to match their SCM-side times (the timestamp of the last checkin which modified them).

See also: `update`

7.2.11 undo

This is used to undo the last `update`, `merge`, or `revert` operation.

```
$ fossil help undo
Usage: fossil undo ?OPTIONS? ?FILENAME...?
      or: fossil redo ?OPTIONS? ?FILENAME...?
```

The `undo` command reverts the changes caused by the previous command

if the previous command is one of the following:

- * fossil update
- * fossil merge
- * fossil revert
- * fossil stash pop
- * fossil stash apply
- * fossil stash drop
- * fossil stash goto
- * fossil clean (*see note below*)

Note: The "fossil clean" command only saves state for files less than 10MiB in size and so if fossil clean deleted files larger than that, then "fossil undo" will not recover the larger files.

If FILENAME is specified then restore the content of the named file(s) but otherwise leave the update or merge or revert in effect. The redo command undoes the effect of the most recent undo.

If the `-n|--dry-run` option is present, no changes are made and instead the undo or redo command explains what actions the undo or redo would have done had the `-n|--dry-run` been omitted.

If the most recent command is not one of those listed as undoable, then the undo command might try to restore the state to be what it was prior to the last undoable command, or it might be a no-op. If in doubt about what the undo command will do, first run it with the `-n` option.

A single level of undo/redo is supported. The undo/redo stack is cleared by the commit and checkout commands. Other commands may or may not clear the undo stack.

Future versions of Fossil might add new commands to the set of commands that are undoable.

Options:

- `-n|--dry-run` do not make changes but show what would be done

See also: commit, status

It acts on a single file or files if specified, otherwise if no file given, it undoes all of the last changes.

7.2.12 diff

The diff command is used to produce a text listing of the difference of a file in the working directory and that same file in the repository. If you don't specify a file it will show the differences between all the changed files in the working directory vs the repository. If you use the `-from` and `-to` options you can specify which versions to check and to compare between two different versions in the repository. Not using the `-to` means compare with the working directory.

If you have configured an external diff program it will be used unless you use the `-i` option which uses the diff built into Fossil.

```
$ fossil help diff
Usage: fossil diff|gdiff ?OPTIONS? ?FILE1? ?FILE2 ...?
```

Show the difference between the current version of each of the FILES specified (as they exist on disk) and that same file as it was checked out. Or if the FILE arguments are omitted, show the unsaved changes currently in the working check-out.

If the `--from VERSION` or `-r VERSION` option is used it specifies the source check-in for the diff operation. If not specified, the source check-in is the base check-in for the current check-out.

If the `--to VERSION` option appears, it specifies the check-in from which the second version of the file or files is taken. If there is no `--to` option then the (possibly edited) files in the current check-out are used.

The `--checkin VERSION` option shows the changes made by check-in VERSION relative to its primary parent.

The `-i` command-line option forces the use of the internal diff logic rather than any external diff program that might be configured using the `setting` command. If no external diff program is configured, then the `-i` option is a no-op. The `-i` option converts `gdiff` into `diff`.

The `-N` or `--new-file` option causes the complete text of added or deleted files to be displayed.

The `--diff-binary` option enables or disables the inclusion of binary files when using an external diff program.

The `--binary` option causes files matching the glob PATTERN to be treated as binary when considering if they should be used with external diff program. This option overrides the `binary-glob` setting.

Options:

<code>--binary PATTERN</code>	Treat files that match the glob PATTERN as binary
<code>--branch BRANCH</code>	Show diff of all changes on BRANCH
<code>--brief</code>	Show filenames only
<code>--checkin VERSION</code>	Show diff of all changes in VERSION
<code>--command PROG</code>	External diff program. Overrides "diff-command"
<code>--context -c N</code>	Use N lines of context
<code>--diff-binary BOOL</code>	Include binary files with external commands
<code>--exec-abs-paths</code>	Force absolute path names on external commands
<code>--exec-rel-paths</code>	Force relative path names on external commands
<code>--from -r VERSION</code>	Select VERSION as source for the diff
<code>--internal -i</code>	Use internal diff logic
<code>--new-file -N</code>	Show complete text of added and deleted files
<code>--numstat</code>	Show only the number of lines delete and added
<code>--side-by-side -y</code>	Side-by-side diff
<code>--strip-trailing-cr</code>	Strip trailing CR
<code>--tclsh PATH</code>	Tcl/Tk used for --tk (default: "tclsh")
<code>--tk</code>	Launch a Tcl/Tk GUI for display
<code>--to VERSION</code>	Select VERSION as target for the diff
<code>--undo</code>	Diff against the "undo" buffer
<code>--unified</code>	Unified diff
<code>-v --verbose</code>	Output complete text of added or deleted files
<code>-w --ignore-all-space</code>	Ignore white space when comparing lines
<code>-W --width N</code>	Width of lines in side-by-side diff
<code>-Z --ignore-trailing-space</code>	Ignore changes to end-of-line whitespace

7.2.13 gdiff

This is the same as the diff command but uses (if configured) a graphical diff program you have on your system. See the settings command for details on how to set the graphical diff program.

```
$ fossil help gdiff
```

```
Usage: fossil diff|gdiff ?OPTIONS? ?FILE1? ?FILE2 ...?
```

Show the difference between the current version of each of the FILES specified (as they exist on disk) and that same file as it was checked out. Or if the FILE arguments are omitted, show the unsaved changes currently in the working check-out.

If the "--from VERSION" or "-r VERSION" option is used it specifies the source check-in for the diff operation. If not specified, the source check-in is the base check-in for the current check-out.

If the "--to VERSION" option appears, it specifies the check-in from

which the second version of the file or files is taken. If there is no "--to" option then the (possibly edited) files in the current check-out are used.

The "--checkin VERSION" option shows the changes made by check-in VERSION relative to its primary parent.

The "-i" command-line option forces the use of the internal diff logic rather than any external diff program that might be configured using the "setting" command. If no external diff program is configured, then the "-i" option is a no-op. The "-i" option converts "gdiff" into "diff".

The "-N" or "--new-file" option causes the complete text of added or deleted files to be displayed.

The "--diff-binary" option enables or disables the inclusion of binary files when using an external diff program.

The "--binary" option causes files matching the glob PATTERN to be treated as binary when considering if they should be used with external diff program. This option overrides the "binary-glob" setting.

Options:

--binary PATTERN	Treat files that match the glob PATTERN as binary
--branch BRANCH	Show diff of all changes on BRANCH
--brief	Show filenames only
--checkin VERSION	Show diff of all changes in VERSION
--command PROG	External diff program. Overrides "diff-command"
--context -c N	Use N lines of context
--diff-binary BOOL	Include binary files with external commands
--exec-abs-paths	Force absolute path names on external commands
--exec-rel-paths	Force relative path names on external commands
--from -r VERSION	Select VERSION as source for the diff
--internal -i	Use internal diff logic
--new-file -N	Show complete text of added and deleted files
--numstat	Show only the number of lines delete and added
--side-by-side -y	Side-by-side diff
--strip-trailing-cr	Strip trailing CR
--tclsh PATH	Tcl/Tk used for --tk (default: "tclsh")
--tk	Launch a Tcl/Tk GUI for display
--to VERSION	Select VERSION as target for the diff
--undo	Diff against the "undo" buffer
--unified	Unified diff
-v --verbose	Output complete text of added or deleted files
-w --ignore-all-space	Ignore white space when comparing lines

```
-W|--width N           Width of lines in side-by-side diff
-Z|--ignore-trailing-space Ignore changes to end-of-line whitespace
```

7.2.14 ui

The `ui` command is used to start Fossil in a local webserver. The `-port` option is used to specify the port it uses, by default it uses 8080. It should automatically start the system's web browser and it will come up with the repository web page. If run within a working directory it will bring up the web page for that repository. If run outside the working directory you can specify the repository on the command line.

```
$ fossil help ui
Usage: fossil server ?OPTIONS? ?REPOSITORY?
       or: fossil ui ?OPTIONS? ?REPOSITORY?
```

Open a socket and begin listening and responding to HTTP requests on TCP port 8080, or on any other TCP port defined by the `-P` or `--port` option. The optional argument is the name of the repository. The repository argument may be omitted if the working directory is within an open checkout.

The `"ui"` command automatically starts a web browser after initializing the web server. The `"ui"` command also binds to 127.0.0.1 and so will only process HTTP traffic from the local machine.

The REPOSITORY can be a directory (aka folder) that contains one or more repositories with names ending in `".fossil"`. In this case, a prefix of the URL pathname is used to search the directory for an appropriate repository. To thwart mischief, the pathname in the URL must contain only alphanumerics, `"_"`, `"/"`, `"-"`, and `"."`, and no `"-"` may occur after `"/"`, and every `"."` must be surrounded on both sides by alphanumerics. Any pathname that does not satisfy these constraints results in a 404 error. Files in REPOSITORY that match the comma-separated list of glob patterns given by `--files` and that have known suffixes such as `".txt"` or `".html"` or `".jpeg"` and do not match the pattern `"*.fossil*"` will be served as static content. With the `"ui"` command, the REPOSITORY can only be a directory if the `--notfound` option is also present.

For the special case REPOSITORY name of `"/"`, the list global configuration database is consulted for a list of all known repositories. The `--replist` option is implied by this special case. See also the `"fossil all ui"` command.

By default, the `"ui"` command provides full administrative access without

having to log in. This can be disabled by turning off the "localauth" setting. Automatic login for the "server" command is available if the --localauth option is present and the "localauth" setting is off and the connection is from localhost. The "ui" command also enables --repolist by default.

Options:

--baseurl URL	Use URL as the base (useful for reverse proxies)
--create	Create a new REPOSITORY if it does not already exist
--extroot DIR	Document root for the /ext extension mechanism
--files GLOBLIST	Comma-separated list of glob patterns for static files
--localauth	enable automatic login for requests from localhost
--localhost	listen on 127.0.0.1 only (always true for "ui")
--https	Indicates that the input is coming through a reverse proxy that has already translated HTTPS into HTTP.
--jsmode MODE	Determine how JavaScript is delivered with pages. Mode can be one of: <ul style="list-style-type: none"> inline All JavaScript is inserted inline at the end of the HTML file. separate Separate HTTP requests are made for each JavaScript file. bundled One single separate HTTP fetches all JavaScript concatenated together. Depending on the needs of any given page, inline and bundled modes might result in a single amalgamated script or several, but both approaches result in fewer HTTP requests than the separate mode.
--max-latency N	Do not let any single HTTP request run for more than N seconds (only works on unix)
--nocompress	Do not compress HTTP replies
--nojail	Drop root privileges but do not enter the chroot jail
--nossll	signal that no SSL connections are available (Always set by default for the "ui" command)
--notfound URL	Redirect
--page PAGE	Start "ui" on PAGE. ex: --page "timeline?y=ci"
-P --port TCPPOINT	listen to request on port TCPPOINT
--th-trace	trace TH1 execution (for debugging purposes)
--repolist	If REPOSITORY is dir, URL "/" lists repos.
--scgi	Accept SCGI rather than HTTP
--skin LABEL	Use override skin LABEL
--usepidkey	Use saved encryption key from parent process. This is only necessary when using SEE on Windows.

See also: cgi, http, winsrv

7.2.15 server

This is a more powerful version of the `ui` command. This allows you to have multiple repositories supported by a single running Fossil webserver. This way you start the server and instead of a particular repository you specify a directory where a number of repositories reside (all having the extension `.fossil`) then you can open and use any of them.

```
$ fossil help server
Usage: fossil server ?OPTIONS? ?REPOSITORY?
      or: fossil ui ?OPTIONS? ?REPOSITORY?
```

Open a socket and begin listening and responding to HTTP requests on TCP port 8080, or on any other TCP port defined by the `-P` or `--port` option. The optional argument is the name of the repository. The repository argument may be omitted if the working directory is within an open checkout.

The `"ui"` command automatically starts a web browser after initializing the web server. The `"ui"` command also binds to 127.0.0.1 and so will only process HTTP traffic from the local machine.

The REPOSITORY can be a directory (aka folder) that contains one or more repositories with names ending in `".fossil"`. In this case, a prefix of the URL pathname is used to search the directory for an appropriate repository. To thwart mischief, the pathname in the URL must contain only alphanumerics, `"_"`, `"/"`, `"-"`, and `"."`, and no `"-"` may occur after `"/"`, and every `"."` must be surrounded on both sides by alphanumerics. Any pathname that does not satisfy these constraints results in a 404 error. Files in REPOSITORY that match the comma-separated list of glob patterns given by `--files` and that have known suffixes such as `".txt"` or `".html"` or `".jpeg"` and do not match the pattern `"*.fossil*"` will be served as static content. With the `"ui"` command, the REPOSITORY can only be a directory if the `--notfound` option is also present.

For the special case REPOSITORY name of `"/"`, the list global configuration database is consulted for a list of all known repositories. The `--replist` option is implied by this special case. See also the `"fossil all ui"` command.

By default, the `"ui"` command provides full administrative access without having to log in. This can be disabled by turning off the `"localauth"` setting. Automatic login for the `"server"` command is available if the `--localauth` option is present and the `"localauth"` setting is off and the connection is from localhost. The `"ui"` command also enables `--replist`

by default.

Options:

<code>--baseurl URL</code>	Use URL as the base (useful for reverse proxies)
<code>--create</code>	Create a new REPOSITORY if it does not already exist
<code>--extroot DIR</code>	Document root for the /ext extension mechanism
<code>--files GLOBLIST</code>	Comma-separated list of glob patterns for static files
<code>--localauth</code>	enable automatic login for requests from localhost
<code>--localhost</code>	listen on 127.0.0.1 only (always true for "ui")
<code>--https</code>	Indicates that the input is coming through a reverse proxy that has already translated HTTPS into HTTP.
<code>--jsmode MODE</code>	Determine how JavaScript is delivered with pages. Mode can be one of: <ul style="list-style-type: none"> <code>inline</code> All JavaScript is inserted inline at the end of the HTML file. <code>separate</code> Separate HTTP requests are made for each JavaScript file. <code>bundled</code> One single separate HTTP fetches all JavaScript concatenated together. Depending on the needs of any given page, inline and bundled modes might result in a single amalgamated script or several, but both approaches result in fewer HTTP requests than the separate mode.
<code>--max-latency N</code>	Do not let any single HTTP request run for more than N seconds (only works on unix)
<code>--nocompress</code>	Do not compress HTTP replies
<code>--nojail</code>	Drop root privileges but do not enter the chroot jail
<code>--nossll</code>	signal that no SSL connections are available (Always set by default for the "ui" command)
<code>--notfound URL</code>	Redirect
<code>--page PAGE</code>	Start "ui" on PAGE. ex: <code>--page "timeline?y=ci"</code>
<code>-P --port TCPPOINT</code>	listen to request on port TCPPOINT
<code>--th-trace</code>	trace TH1 execution (for debugging purposes)
<code>--repolist</code>	If REPOSITORY is dir, URL "/" lists repos.
<code>--scgi</code>	Accept SCGI rather than HTTP
<code>--skin LABEL</code>	Use override skin LABEL
<code>--usepidkey</code>	Use saved encryption key from parent process. This is only necessary when using SEE on Windows.

See also: `cgi`, `http`, `winsrv`

7.2.16 `commit` or `ci`

This is the command used to put the current changes in the working directory into the repository, giving this a new version and updating the timeline.

```
$ fossil help commit
Usage: fossil commit ?OPTIONS? ?FILE...?
       or: fossil ci ?OPTIONS? ?FILE...?
```

Create a new version containing all of the changes in the current checkout. You will be prompted to enter a check-in comment unless the comment has been specified on the command-line using "-m" or a file containing the comment using -M. The editor defined in the "editor" fossil option (see fossil help set) will be used, or from the "VISUAL" or "EDITOR" environment variables (in that order) if no editor is set.

All files that have changed will be committed unless some subset of files is specified on the command line.

The --branch option followed by a branch name causes the new check-in to be placed in a newly-created branch with the name passed to the --branch option.

Use the --branchcolor option followed by a color name (ex: '#ffc0c0') to specify the background color of entries in the new branch when shown in the web timeline interface. The use of the --branchcolor option is not recommended. Instead, let Fossil choose the branch color automatically.

The --bgcolor option works like --branchcolor but only sets the background color for a single check-in. Subsequent check-ins revert to the default color.

A check-in is not permitted to fork unless the --allow-fork option appears. An empty check-in (i.e. with nothing changed) is not allowed unless the --allow-empty option appears. A check-in may not be older than its ancestor unless the --allow-older option appears. If any of files in the check-in appear to contain unresolved merge conflicts, the check-in will not be allowed unless the --allow-conflict option is present. In addition, the entire check-in process may be aborted if a file contains content that appears to be binary, Unicode text, or text with CR/LF line endings unless the interactive user chooses to proceed. If there is no interactive user or these warnings should be skipped for some other reason, the --no-warnings option may be used. A check-in is not allowed against a closed leaf.

If a commit message is blank, you will be prompted: ("continue (y/N)?") to confirm you really want to commit with a blank commit message. The default value is "N", do not commit.

The `--private` option creates a private check-in that is never synced. Children of private check-ins are automatically private.

The `--tag` option applies the symbolic tag name to the check-in.

The `--hash` option detects edited files by computing each file's artifact hash rather than just checking for changes to its size or mtime.

Options:

<code>--allow-conflict</code>	allow unresolved merge conflicts
<code>--allow-empty</code>	allow a commit with no changes
<code>--allow-fork</code>	allow the commit to fork
<code>--allow-older</code>	allow a commit older than its ancestor
<code>--baseline</code>	use a baseline manifest in the commit process
<code>--bgcolor COLOR</code>	apply COLOR to this one check-in only
<code>--branch NEW-BRANCH-NAME</code>	check in to this new branch
<code>--branchcolor COLOR</code>	apply given COLOR to the branch
<code>--close</code>	close the branch being committed
<code>--date-override DATETIME</code>	DATE to use instead of 'now'
<code>--delta</code>	use a delta manifest in the commit process
<code>--hash</code>	verify file status using hashing rather than relying on file mtimes
<code>--integrate</code>	close all merged-in branches
<code>-m --comment COMMENT-TEXT</code>	use COMMENT-TEXT as commit comment
<code>-M --message-file FILE</code>	read the commit comment from given file
<code>--mimetype MIMETYPE</code>	mimetype of check-in comment
<code>-n --dry-run</code>	If given, display instead of run actions
<code>--no-prompt</code>	This option disables prompting the user for input and assumes an answer of 'No' for every question.
<code>--no-warnings</code>	omit all warnings about file contents
<code>--no-verify</code>	do not run before-commit hooks
<code>--nosign</code>	do not attempt to sign this commit with gpg
<code>--override-lock</code>	allow a check-in even though parent is locked
<code>--private</code>	do not sync changes and their descendants
<code>--tag TAG-NAME</code>	assign given tag TAG-NAME to the check-in
<code>--trace</code>	debug tracing.
<code>--user-override USER</code>	USER to use instead of the current default

DATETIME may be "now" or "YYYY-MM-DDTHH:MM:SS.SSS". If in year-month-day form, it may be truncated, the "T" may be replaced by a space, and it may also name a timezone offset from UTC as "-HH:MM" (westward) or "+HH:MM" (eastward). Either no timezone suffix or "Z" means UTC.

See also: `branch`, `changes`, `update`, `extras`, `sync`

It's a very good idea to always put a comment (`-comment` or `-m`) text on any commit. This way you get documentation in the timeline.

7.3 Maintenance

These commands you will probably use less often since the actions they perform are not needed in normal operation. You will have to use them and referring here or to `fossil help` will probably be required before use. Some of them like `new` or `clone` are only needed when you start a repository. Others like `rebuild` or `reconstruct` are only needed to fix or update a repository.

7.3.1 new

This command is used to create a new repository.

```
$ fossil help new
Usage: fossil new ?OPTIONS? FILENAME
       or: fossil init ?OPTIONS? FILENAME
```

Create a repository for a new project in the file named `FILENAME`. This command is distinct from `clone`. The `clone` command makes a copy of an existing project. This command starts a new project.

By default, your current login name is used to create the default admin user. This can be overridden using the `-A|--admin-user` parameter.

By default, all settings will be initialized to their default values. This can be overridden using the `--template` parameter to specify a repository file from which to copy the initial settings. When a template repository is used, almost all of the settings accessible from the setup page, either directly or indirectly, will be copied. Normal users and their associated permissions will not be copied; however, the system default users `"anonymous"`, `"nobody"`, `"reader"`, `"developer"`, and their associated permissions will be copied.

Options:

<code>--template</code>	<code>FILE</code>	Copy settings from repository file
<code>--admin-user -A</code>	<code>USERNAME</code>	Select given <code>USERNAME</code> as admin user
<code>--date-override</code>	<code>DATETIME</code>	Use <code>DATETIME</code> as time of the initial check-in
<code>--shal</code>		Use an initial hash policy of <code>"shal"</code>

`DATETIME` may be `"now"` or `"YYYY-MM-DDTHH:MM:SS.SSS"`. If in year-month-day form, it may be truncated, the `"T"` may be replaced by

a space, and it may also name a timezone offset from UTC as "-HH:MM" (westward) or "+HH:MM" (eastward). Either no timezone suffix or "Z" means UTC.

See also: clone

The file name specifies the new repository name. The options provided allow you to specify the admin user name if you want it to be different than your current login and the starting date if you want it to be different than now.

7.3.2 clone

The clone command is used to create your own local version of the master repository. If you are supporting multiple users via a network accessible version of the original repository (see Section[*sub:Server-Setup*]), then this command will copy that repository to your machine. Also it will make a link between your copy and the master, so that changes made in your copy will be propagated to the master.

```
$ fossil help clone
```

```
Usage: fossil clone ?OPTIONS? URI ?FILENAME?
```

Make a clone of a repository specified by URI in the local file named FILENAME. If FILENAME is omitted, then an appropriate filename is deduced from last element of the path in the URL.

URI may be one of the following forms ([...] denotes optional elements):

- * HTTP/HTTPS protocol:

```
http[s]://[userid[:password]@]host[:port][/path]
```

- * SSH protocol:

```
ssh://[userid@]host[:port]/path/to/repo.fossil[?fossil=path/fossil.exe]
```

- * Filesystem:

```
[file://]path/to/repo.fossil
```

For ssh and filesystem, path must have an extra leading '/' to use an absolute path.

Use %HH escapes for special characters in the userid and password. For example "%40" in place of "@", "%2f" in place of "/", and "%3a" in place of ":".

Note that in Fossil (in contrast to some other DVCSes) a repository is distinct from a checkout. Cloning a repository is not the same thing as opening a repository. This command always clones the repository. This command might also open the repository, but only if the `--no-open` option is omitted and either the `--workdir` option is included or the `FILENAME` argument is omitted. Use the separate `open` command to open a repository that was previously cloned and already exists on the local machine.

By default, the current login name is used to create the default admin user for the new clone. This can be overridden using the `-A|--admin-user` parameter.

Options:

<code>--admin-user -A USERNAME</code>	Make USERNAME the administrator
<code>--httpauth -B USER:PASS</code>	Add HTTP Basic Authorization to requests
<code>--nocompress</code>	Omit extra delta compression
<code>--no-open</code>	Clone only. Do not open a check-out.
<code>--once</code>	Don't remember the URI.
<code>--private</code>	Also clone private branches
<code>--save-http-password</code>	Remember the HTTP password without asking
<code>--ssh-command -c SSH</code>	Use SSH as the "ssh" command
<code>--ssl-identity FILENAME</code>	Use the SSL identity if requested by the server
<code>-u --unversioned</code>	Also sync unversioned content
<code>-v --verbose</code>	Show more statistics in output
<code>--workdir DIR</code>	Also open a checkout in DIR

See also: `init`, `open`

As with `create`, you can specify the admin user for this clone with an option. The URL for the master repository is of the form:

```
https://user:password@domain
```

Where `user` and `password` are for a valid user of the selected repository. It is best to check the path with a browser before doing the clone. Make sure you can reach it, for example the repository for this book is:

```
https://www.fossil-scm.org/schimpf-book/home
```

Putting that into a browser should get you the home page for this book. (See Figure [fig:Web-access-to]). After you have verified that, then running the clone command should work.

Don't forget (as I always do) to put in the file name for the local repository, (see `FILENAME` above)

7.3.3 open

The open command is used to copy the files in a repository to a working directory. Doing this allows you to build or modify the product. The command also links this working directory to the repository so commits will go into the repository.

```
$ fossil help open
Usage: fossil open REPOSITORY ?VERSION? ?OPTIONS?
```

Open a new connection to the repository name REPOSITORY. A checkout for the repository is created with its root at the current working directory, or in DIR if the "--workdir DIR" is used. If VERSION is specified then that version is checked out. Otherwise the most recent check-in on the main branch (usually "trunk") is used.

REPOSITORY can be the filename for a repository that already exists on the local machine or it can be a URI for a remote repository. If REPOSITORY is a URI in one of the formats recognized by the clone command, then remote repo is first cloned, then the clone is opened. The clone will be stored in the current directory, or in DIR if the "--reporidir DIR" option is used. The name of the clone will be taken from the last term of the URI. For "http:" and "https:" URIs, you can append an extra term to the end of the URI to get any repository name you like. For example:

```
fossil open https://fossil-scm.org/home/new-name
```

The base URI for cloning is "https://fossil-scm.org/home". The extra "new-name" term means that the cloned repository will be called "new-name.fossil".

Options:

--empty	Initialize checkout as being empty, but still connected with the local repository. If you commit this checkout, it will become a new "initial" commit in the repository.
-f --force	Continue with the open even if the working directory is not empty.
--force-missing	Force opening a repository with missing content
--keep	Only modify the manifest and manifest.uuid files
--nested	Allow opening a repository inside an opened checkout
--reporidir DIR	If REPOSITORY is a URI that will be cloned, store the clone in DIR rather than in "."
--setmtime	Set timestamps of all files to match their SCM-side times (the timestamp of the last checkin which modified them).
--workdir DIR	Use DIR as the working directory instead of ".". The DIR

directory is created if it does not exist.

See also: `close`, `clone`

If you have multiple users or have a branched repository then it is probably wise to specify the particular version you want. When you run this it will create all the files and directories in the repository in your work area. In addition the files *FOSSIL*, *manifest* and *manifest.uuid* will be created by Fossil.

You can have multiple open connections to the same repository at the same time. Every open connection just needs to be in its own separate working directory. This is very useul when you have many branches and do not want to commit your work just to switch over to another branch and continue or check something there. Just create a new open connection in another directory and you can `cd` between them and work on multiple branches at the same time. If you want to know where all the open checkouts of your repository are located, use `fossil info -v`.

7.3.4 close

This is the opposite of `open`, in that it breaks the connection between this working directory and the Fossil repository.

```
$ fossil help close
Usage: fossil close ?OPTIONS?
```

The opposite of "open". Close the current database connection. Require a `-f` or `--force` flag if there are unsaved changes in the current check-out or if there is non-empty stash.

Options:

```
--force|-f necessary to close a check out with uncommitted changes
```

See also: `open`

This is useful if you need to abandon the current working directory. Fossil will not let you do this if there are changes between the current directory and the repository. With the force flag you can explicitly cut the connection even if there are changes.

7.3.5 version

This command is used to show the current version of fossil.

```
$ fossil help version
Usage: fossil version ?-verbose|-v?
```

Print the source code version number for the fossil executable.

If the verbose option is specified, additional details will be output about what optional features this binary was compiled with

```
$ fossil version
This is fossil version 2.14 [939a13d94f] 2020-11-20 19:28:05 UTC
```

The above figure shows the help and example of running the command. When you have problems with fossil it is very important to have this version information. You can then inquire of the Fossil news group about this problem and with the version information they can easily tell you if the problem is fixed already or is new.

7.3.6 rebuild

If you update your copy of Fossil you will want to run this command against all the repositories you have. This will automatically update them to the new version of Fossil.

```
$ fossil help rebuild
Usage: fossil rebuild ?REPOSITORY? ?OPTIONS?
```

Reconstruct the named repository database from the core records. Run this command after updating the fossil executable in a way that changes the database schema.

Options:

<code>--analyze</code>	Run ANALYZE on the database after rebuilding
<code>--cluster</code>	Compute clusters for unclustered artifacts
<code>--compress</code>	Strive to make the database as small as possible
<code>--compress-only</code>	Skip the rebuilding step. Do <code>--compress</code> only
<code>--deanalyze</code>	Remove ANALYZE tables from the database
<code>--force</code>	Force the rebuild to complete even if errors are seen
<code>--ifneeded</code>	Only do the rebuild if it would change the schema version
<code>--index</code>	Always add in the full-text search index
<code>--noverify</code>	Skip the verification of changes to the BLOB table
<code>--noindex</code>	Always omit the full-text search index
<code>--pagesize N</code>	Set the database pagesize to N. (512..65536 and power of 2)
<code>--quiet</code>	Only show output if there are errors
<code>--randomize</code>	Scan artifacts in a random order
<code>--stats</code>	Show artifact statistics after rebuilding
<code>--vacuum</code>	Run VACUUM on the database after rebuilding
<code>--wal</code>	Set Write-Ahead-Log journalling mode on the database

7.3.7 all

This command is actually a modifier and when used before certain commands will run them on all the repositories.

```
$ fossil help all
Usage: fossil all SUBCOMMAND ...
```

The `~/.fossil` file records the location of all repositories for a user. This command performs certain operations on all repositories that can be useful before or after a period of disconnected operation.

On Win32 systems, the file is named `"_fossil"` and is located in `%LOCALAPPDATA%`, `%APPDATA%` or `%HOMEPATH%`.

Available operations are:

backup	Backup all repositories. The argument must be the name of a directory into which all backup repositories are written.
cache	Manages the cache used for potentially expensive web pages. Any additional arguments are passed on verbatim to the cache command.
changes	Shows all local checkouts that have uncommitted changes. This operation has no additional options.
clean	Delete all "extra" files in all local checkouts. Extreme caution should be exercised with this command because its effects cannot be undone. Use of the <code>--dry-run</code> option to carefully review the local checkouts to be operated upon and the <code>--whatif</code> option to carefully review the files to be deleted beforehand is highly recommended. The command line options supported by the clean command itself, if any are present, are passed along verbatim.
config	Only the "config pull AREA" command works.
dbstat	Run the "dbstat" command on all repositories.
extras	Shows "extra" files from all local checkouts. The command line options supported by the extra command itself, if any are present, are passed along verbatim.
fts-config	Run the "fts-config" command on all repositories.

info	Run the "info" command on all repositories.
pull	Run a "pull" operation on all repositories. Only the --verbose option is supported.
push	Run a "push" on all repositories. Only the --verbose option is supported.
rebuild	Rebuild on all repositories. The command line options supported by the rebuild command itself, if any are present, are passed along verbatim. The --force and --randomize options are not supported.
sync	Run a "sync" on all repositories. Only the --verbose and --unversioned options are supported.
set unset	Run the "setting", "set", or "unset" commands on all repositories. These command are particularly useful in conjunction with the "max-loadavg" setting which cannot otherwise be set globally.
server ui	Run the "ui" or "server" commands on all repositories. The root URI gives a listing of all repos.

In addition, the following maintenance operations are supported:

add	Add all the repositories named to the set of repositories tracked by Fossil. Normally Fossil is able to keep up with this list by itself, but sometimes it can benefit from this hint if you rename repositories.
ignore	Arguments are repositories that should be ignored by subsequent clean, extras, list, pull, push, rebuild, and sync operations. The -c --ckout option causes the listed local checkouts to be ignored instead.
list ls	Display the location of all repositories. The -c --ckout option causes all local checkouts to be listed instead.

Repositories are automatically added to the set of known repositories when one of the following commands are run against the repository: clone, info, pull, push, or sync. Even previously ignored repositories are added back to the list of repositories by these commands.

Options:

```

--dry-run          If given, display instead of run actions.
--showfile        Show the repository or checkout being operated upon.
--stop-on-error    Halt immediately if any subprocess fails.

```

7.3.8 push

This command will push changes in the local repository to the master or remote repository.

```

$ fossil help push
Usage: fossil push ?URL? ?options?

```

Push all sharable changes from the local repository to a remote repository. Sharable changes include public check-ins, edits to wiki pages, tickets, and tech-notes, as well as forum content. Use `--private` to also push private branches. Use the "configuration push" command to push website configuration details.

If URL is not specified, then the URL from the most recent clone, push, pull, remote, or sync command is used. See "fossil help clone" for details on the URL formats.

Options:

```

-B|--httpauth USER:PASS  Credentials for the simple HTTP auth protocol,
                           if required by the remote website
--ipv4                    Use only IPv4, not IPv6
--once                    Do not remember URL for subsequent syncs
--proxy PROXY             Use the specified HTTP proxy
--private                 Push private branches too
-R|--repository REPO     Local repository to push from
--ssl-identity FILE       Local SSL credentials, if requested by remote
--ssh-command SSH        Use SSH as the "ssh" command
-v|--verbose              Additional (debugging) output
--verify                  Exchange extra information with the remote
                           to ensure no content is overlooked

```

See also: clone, config, pull, remote, sync

7.3.9 pull

This command will copy changes from the remote repository to the local repository. You could then use update to apply these changes to checked out files.

```

$ fossil help pull
Usage: fossil pull ?URL? ?options?

```

Pull all sharable changes from a remote repository into the local repository. Sharable changes include public check-ins, edits to wiki pages, tickets, and tech-notes, as well as forum content. Add the `--private` option to pull private branches. Use the "configuration pull" command to pull website configuration details.

If URL is not specified, then the URL from the most recent clone, push, pull, remote, or sync command is used. See "fossil help clone" for details on the URL formats.

Options:

<code>-B --httpauth USER:PASS</code>	Credentials for the simple HTTP auth protocol, if required by the remote website
<code>--from-parent-project</code>	Pull content from the parent project
<code>--ipv4</code>	Use only IPv4, not IPv6
<code>--once</code>	Do not remember URL for subsequent syncs
<code>--private</code>	Pull private branches too
<code>--project-code CODE</code>	Use CODE as the project code
<code>--proxy PROXY</code>	Use the specified HTTP proxy
<code>-R --repository REPO</code>	Local repository to pull into
<code>--ssl-identity FILE</code>	Local SSL credentials, if requested by remote
<code>--ssh-command SSH</code>	Use SSH as the "ssh" command
<code>-v --verbose</code>	Additional (debugging) output
<code>--verily</code>	Exchange extra information with the remote to ensure no content is overlooked

See also: clone, config, push, remote, sync

7.3.10 sync

This command is used to sync a remote copy with the original copy of the repository, it does both a push and pull. This can also be used to switch a local repository to a different main repository by specifying the URL of a remote repository. If you want to run the update command with `-n` where it does a dry run, this does not do a sync first so doing `fossil sync` then `fossil update -n` will do that for you.

```
$ fossil help sync
Usage: fossil sync ?URL? ?options?
```

Synchronize all sharable changes between the local repository and a remote repository. Sharable changes include public check-ins and edits to wiki pages, tickets, and technical notes.

If URL is not specified, then the URL from the most recent clone, push,

pull, remote, or sync command is used. See "fossil help clone" for details on the URL formats.

Options:

-B --httpauth USER:PASS	Credentials for the simple HTTP auth protocol, if required by the remote website
--ipv4	Use only IPv4, not IPv6
--once	Do not remember URL for subsequent syncs
--proxy PROXY	Use the specified HTTP proxy
--private	Sync private branches too
-R --repository REPO	Local repository to sync with
--ssl-identity FILE	Local SSL credentials, if requested by remote
--ssh-command SSH	Use SSH as the "ssh" command
-u --unversioned	Also sync unversioned content
-v --verbose	Additional (debugging) output
--verily	Exchange extra information with the remote to ensure no content is overlooked

See also: clone, pull, push, remote

7.3.11 clean

This call can be used to remove all the "extra" files in a source tree. This is useful if you wish to tidy up a source tree or to do a clean build.

```
$ fossil help clean
Usage: fossil clean ?OPTIONS? ?PATH ...?
```

Delete all "extra" files in the source tree. "Extra" files are files that are not officially part of the checkout. If one or more PATH arguments appear, then only the files named, or files contained with directories named, will be removed.

If the --prompt option is used, prompts are issued to confirm the permanent removal of each file. Otherwise, files are backed up to the undo buffer prior to removal, and prompts are issued only for files whose removal cannot be undone due to their large size or due to --disable-undo being used.

The --force option treats all prompts as having been answered yes, whereas --no-prompt treats them as having been answered no.

Files matching any glob pattern specified by the --clean option are deleted without prompting, and the removal cannot be undone.

No file that matches glob patterns specified by `--ignore` or `--keep` will ever be deleted. Files and subdirectories whose names begin with "." are automatically ignored unless the `--dotfiles` option is used.

The default values for `--clean`, `--ignore`, and `--keep` are determined by the (versionable) `clean-glob`, `ignore-glob`, and `keep-glob` settings.

The `--verily` option ignores the `keep-glob` and `ignore-glob` settings and turns on `--force`, `--emptydirs`, `--dotfiles`, and `--disable-undo`. Use the `--verily` option when you really want to clean up everything. Extreme care should be exercised when using the `--verily` option.

Options:

<code>--allckouts</code>	Check for empty directories within any checkouts that may be nested within the current one. This option should be used with great care because the <code>empty-dirs</code> setting (and other applicable settings) belonging to the other repositories, if any, will not be checked.
<code>--case-sensitive</code> <i>BOOL</i>	Override case-sensitive setting
<code>--dirsonly</code>	Only remove empty directories. No files will be removed. Using this option will automatically enable the <code>--emptydirs</code> option as well.
<code>--disable-undo</code>	WARNING: This option disables use of the undo mechanism for this clean operation and should be used with extreme caution.
<code>--dotfiles</code>	Include files beginning with a dot (".").
<code>--emptydirs</code>	Remove any empty directories that are not explicitly exempted via the <code>empty-dirs</code> setting or another applicable setting or command line argument. Matching files, if any, are removed prior to checking for any empty directories; therefore, directories that contain only files that were removed will be removed as well.
<code>-f --force</code>	Remove files without prompting.
<code>-i --prompt</code>	Prompt before removing each file. This option implies the <code>--disable-undo</code> option.
<code>-x --verily</code>	WARNING: Removes everything that is not a managed file or the repository itself. This option implies the <code>--force</code> , <code>--emptydirs</code> , <code>--dotfiles</code> , and <code>--disable-undo</code> options. Furthermore, it completely disregards the <code>keep-glob</code> and <code>ignore-glob</code> settings. However, it does honor the <code>--ignore</code> and <code>--keep</code> options.
<code>--clean</code> <i>CSG</i>	WARNING: Never prompt to delete any files matching this comma separated list of glob patterns. Also,

	deletions of any files matching this pattern list cannot be undone.
--ignore CSG	Ignore files matching patterns from the comma separated list of glob patterns.
--keep <CSG>	Keep files matching this comma separated list of glob patterns.
-n --dry-run	Delete nothing, but display what would have been deleted.
--no-prompt	This option disables prompting the user for input and assumes an answer of 'No' for every question.
--temp	Remove only Fossil-generated temporary files.
-v --verbose	Show all files as they are removed.

See also: `addrremove`, `extras`, `status`

7.3.12 branch

This command is used if you want to create or list branches in a repository. Previously we discussed forks (See Section [sub:Complications]); branches are the same idea but under user control. This would be where you have version 1.0 of something but want to branch off version 2.0 to add new features but want to keep a 1.0 branch for maintenance.

```
$ fossil help branch
```

```
Usage: fossil branch SUBCOMMAND ... ?OPTIONS?
```

Run various subcommands to manage branches of the open repository or of the repository identified by the `-R` or `--repository` option.

```
fossil branch current
```

Print the name of the branch for the current check-out

```
fossil branch info BRANCH-NAME
```

Print information about a branch

```
fossil branch list|ls ?OPTIONS?
```

List all branches. Options:

-a --all	List all branches. Default show only open branches
-c --closed	List closed branches.
-r	Reverse the sort order
-t	Show recently changed branches first

```
fossil branch new BRANCH-NAME BASIS ?OPTIONS?
```

Create a new branch `BRANCH-NAME` off of check-in `BASIS`. Supported options for this subcommand include:

```
--private           branch is private (i.e., remains local)
--bgcolor COLOR     use COLOR instead of automatic background
--nosign           do not sign contents on this branch
--date-override DATE DATE to use instead of 'now'
--user-override USER USER to use instead of the current default
```

`DATE` may be "now" or "YYYY-MM-DDTHH:MM:SS.SSS". If in year-month-day form, it may be truncated, the "T" may be replaced by a space, and it may also name a timezone offset from UTC as "-HH:MM" (westward) or "+HH:MM" (eastward). Either no timezone suffix or "Z" means UTC.

Options valid for all subcommands:

```
-R|--repository FILE      Run commands on repository FILE
```

7.3.13 merge

This command does the opposite of `branch`, it brings two branches together.

```
$ fossil help merge
Usage: fossil merge ?OPTIONS? ?VERSION?
```

The argument `VERSION` is a version that should be merged into the current checkout. All changes from `VERSION` back to the nearest common ancestor are merged. Except, if either of the `--cherrypick` or `--backout` options are used only the changes associated with the single check-in `VERSION` are merged. The `--backout` option causes the changes associated with `VERSION` to be removed from the current checkout rather than added.

If the `VERSION` argument is omitted, then Fossil attempts to find a recent fork on the current branch to merge.

Only file content is merged. The result continues to use the file and directory names from the current checkout even if those names might have been changed in the branch being merged in.

Options:

```
--backout           Do a reverse cherrypick merge against VERSION.
                    In other words, back out the changes that were
                    added by VERSION.
```

<code>--baseline BASELINE</code>	Use BASELINE as the "pivot" of the merge instead of the nearest common ancestor. This allows a sequence of changes in a branch to be merged without having to merge the entire branch.
<code>--binary GLOBPATTERN</code>	Treat files that match GLOBPATTERN as binary and do not try to merge parallel changes. This option overrides the "binary-glob" setting.
<code>--case-sensitive BOOL</code>	Override the case-sensitive setting. If false, files whose names differ only in case are taken to be the same file.
<code>--cherrypick</code>	Do a cherrypick merge VERSION into the current checkout. A cherrypick merge pulls in the changes of the single check-in VERSION, rather than all changes back to the nearest common ancestor.
<code>-f --force</code>	Force the merge even if it would be a no-op.
<code>--force-missing</code>	Force the merge even if there is missing content.
<code>--integrate</code>	Merged branch will be closed when committing.
<code>-K --keep-merge-files</code>	On merge conflict, retain the temporary files used for merging, named *-baseline, *-original, and *-merge.
<code>-n --dry-run</code>	If given, display instead of run actions
<code>-v --verbose</code>	Show additional details of the merge

7.3.14 tag

This command can be used to control "tags" which are attributes added to any entry in the time line. You can also add/delete/control these tags from the UI by going into the timeline, picking an entry then doing an edit. See Figure [fig:Remove-Leaf].

```
$ fossil help tag
Usage: fossil tag SUBCOMMAND ...
```

Run various subcommands to control tags and properties.

```
fossil tag add ?OPTIONS? TAGNAME CHECK-IN ?VALUE?
```

Add a new tag or property to CHECK-IN. The tag will be usable instead of a CHECK-IN in commands such as update and merge. If the `--propagate` flag is present, the tag value propagates to all descendants of CHECK-IN

Options:

<code>--raw</code>	Raw tag name.
<code>--propagate</code>	Propagating tag.
<code>--date-override DATETIME</code>	Set date and time added.
<code>--user-override USER</code>	Name USER when adding the tag.
<code>--dryrun -n</code>	Display the tag text, but do not actually insert it into the database.

The `--date-override` and `--user-override` options support importing history from other SCM systems. DATETIME has the form 'YYYY-MMM-DD HH:MM:SS'.

`fossil tag cancel ?--raw? TAGNAME CHECK-IN`

Remove the tag TAGNAME from CHECK-IN, and also remove the propagation of the tag to any descendants. Use the `--dryrun` or `-n` options to see what would have happened.

Options:

<code>--raw</code>	Raw tag name.
<code>--date-override DATETIME</code>	Set date and time deleted.
<code>--user-override USER</code>	Name USER when deleting the tag.
<code>--dryrun -n</code>	Display the control artifact, but do not insert it into the database.

`fossil tag find ?OPTIONS? TAGNAME`

List all objects that use TAGNAME. TYPE can be "ci" for check-ins or "e" for events. The limit option limits the number of results to the given value.

Options:

<code>--raw</code>	Raw tag name.
<code>-t --type TYPE</code>	One of "ci", or "e".
<code>-n --limit N</code>	Limit to N results.

`fossil tag list|ls ?OPTIONS? ?CHECK-IN?`

List all tags, or if CHECK-IN is supplied, list all tags and their values for CHECK-IN. The tagtype option

takes one of: propagated, singleton, cancel.

Options:

```
--raw           List tags raw names of tags
--tagtype TYPE  List only tags of type TYPE
-v|--inverse    Inverse the meaning of --tagtype TYPE.
```

The option `--raw` allows the manipulation of all types of tags used for various internal purposes in fossil. It also shows "cancel" tags for the "find" and "list" subcommands. You should not use this option to make changes unless you are sure what you are doing.

If you need to use a tagname that might be confused with a hexadecimal baseline or artifact ID, you can explicitly disambiguate it by prefixing it with "tag:". For instance:

```
fossil update decaf
```

will be taken as an artifact or baseline ID and fossil will probably complain that no such revision was found. However

```
fossil update tag:decaf
```

will assume that "decaf" is a tag/branch name.

7.3.15 settings

This command is used to set or unset a number of properties for fossil.

```
$ fossil help settings
```

```
Usage: fossil settings ?SETTING? ?VALUE? ?OPTIONS?
```

```
or: fossil unset SETTING ?OPTIONS?
```

The "settings" command with no arguments lists all settings and their values. With just a SETTING name it shows the current value of that setting. With a VALUE argument it changes the property for the current repository.

Settings marked as versionable are overridden by the contents of the file named `.fossil-settings/PROPERTY` in the check-out root, if that file exists.

The "unset" command clears a setting.

Settings can have both a "local" repository-only value and "global" value that applies to all repositories. The local values are stored in the

"config" table of the repository and the global values are stored in the configuration database. If both a local and a global value exists for a setting, the local value takes precedence. This command normally operates on the local settings. Use the `--global` option to change global settings.

Options:

- `--global` set or unset the given property globally instead of setting or unsetting it for the open repository only.
- `--exact` only consider exact name matches.

See also: configuration

7.3.16 info

```
fossil info ?VERSION | REPOSITORY_FILENAME? ?OPTIONS?
```

This command show various information on the repository, a bit like the `status` command does. Using the command without any arguments, it shows information on the current checkout:

- name of the project and its code (hash value)
- path to the repository
- path to Fossil's global configuration database for the user (typically `~/fossil`)
- path to the local open checkout
- id and time stamp of the last checkout, its parent and child
- tags
- last commit message

When you do not want information on the last check-in, but on other versions, you just supply the version on the command line. You can also explicitly provide the repository file with the `-R` option. Then you will only get the project name and code and the number of chek-ins.

The `-v` option is helpful since it provides not only the above information on the current checkout but also tells you the locations of all other known checkouts there are (if you have multiple open checkouts of the same repository in different directories). It also lists the possible access URLs that have been used over time (for the ui).

7.3.17 publish

```
fossil publish ?--only? TAGS...
```

This command causes artifacts identified by the given `TAGS . . .` (these can be more than one tag) to be published (i.e. made non-private). This can be used

(for example) to convert a private branch into a public branch, or to publish a bundle that was imported privately.

If any of TAGS names a branch, then all check-ins on the most recent instance of that branch are included, not just the most recent check-in.

If any of TAGS name check-ins then all files and tags associated with those check-ins are also published automatically. If the `--only` option is used, then only the specific artifacts identified by TAGS are published.

If a TAG is already public, this command is a harmless no-op.

7.4 Miscellaneous

These are commands that don't seem to fit in any category but are useful.

7.4.1 zip

You can do what this command does from the web based user interface. In Figure [fig:Timeline-Detail] you can download a ZIP archive of the particular version of the files. This command lets you do it from the command line.

```
$ fossil help zip
Usage: fossil zip VERSION OUTPUTFILE [OPTIONS]
```

Generate a ZIP archive for a check-in. If the `--name` option is used, its argument becomes the name of the top-level directory in the resulting ZIP archive. If `--name` is omitted, the top-level directory name is derived from the project name, the check-in date and time, and the artifact ID of the check-in.

The GLOBLIST argument to `--exclude` and `--include` can be a comma-separated list of glob patterns, where each glob pattern may optionally be enclosed in `"..."` or `'...'` so that it may contain commas. If a file matches both `--include` and `--exclude` then it is excluded.

Options:

<code>-X --exclude GLOBLIST</code>	Comma-separated list of GLOBs of files to exclude
<code>--include GLOBLIST</code>	Comma-separated list of GLOBs of files to include
<code>--name DIRECTORYNAME</code>	The name of the top-level directory in the archive
<code>-R REPOSITORY</code>	Specify a Fossil repository

7.4.2 user

This command lets you modify user information. Again this is a command line duplication of what you can do from the user interface in the browser, see Figure [fig:New-Editor-user].


```
$ fossil help user
```

```
Usage: fossil user SUBCOMMAND ... ?-R|--repository FILE?
```

Run various subcommands on users of the open repository or of the repository identified by the `-R` or `--repository` option.

```
fossil user capabilities USERNAME ?STRING?
```

Query or set the capabilities for user `USERNAME`

```
fossil user contact USERNAME ?CONTACT-INFO?
```

Query or set contact information for user `USERNAME`

```
fossil user default ?USERNAME?
```

Query or set the default user. The default user is the user for command-line interaction.

```
fossil user list
```

```
fossil user ls
```

List all users known to the repository

```
fossil user new ?USERNAME? ?CONTACT-INFO? ?PASSWORD?
```

Create a new user in the repository. Users can never be deleted. They can be denied all access but they must continue to exist in the database.

```
fossil user password USERNAME ?PASSWORD?
```

Change the web access password for a user.

7.4.3 **finfo**

This command will print the history of any particular file. This can be useful if you need this history in some other system. You can pass this text file to the other system which can then parse and use the data.

```
$ fossil help finfo
```

```
Usage: fossil finfo ?OPTIONS? FILENAME
```

Print the complete change history for a single file going backwards in time. The default mode is `-l`.

For the `-l|--log` mode: If `"-b|--brief"` is specified one line per revision is printed, otherwise the full comment is printed. The `"-n|--limit N"` and `"--offset P"` options limits the output to the first N changes after skipping P changes.

In the `-s` mode prints the status as `<status> <revision>`. This is a quick status and does not check for up-to-date-ness of the file.

In the `-p` mode, there's an optional flag `"-r|--revision REVISION"`. The specified version (or the latest checked out version) is printed to stdout. The `-p` mode is another form of the `"cat"` command.

Options:

<code>-b --brief</code>	display a brief (one line / revision) summary
<code>--case-sensitive B</code>	Enable or disable case-sensitive filenames. B is a boolean: "yes", "no", "true", "false", etc.
<code>-l --log</code>	select log mode (the default)
<code>-n --limit N</code>	Display the first N changes (default unlimited). N less than 0 means no limit.
<code>--offset P</code>	skip P changes
<code>-p --print</code>	select print mode
<code>-r --revision R</code>	print the given revision (or ckout, if none is given) to stdout (only in print mode)
<code>-s --status</code>	select status mode (print a status indicator for FILE)
<code>-W --width N</code>	Width of lines (default is to auto-detect). Must be more than 22 or else 0 to indicate no limit.

See also: `artifact`, `cat`, `descendants`, `info`, `leaves`

An example would be to run it on the `outline.txt` file in the earlier version of our book directory:

```
$ fossil finfo outline.txt

History of outline.txt

2010-05-17 [0272dc0169] Finished maintenance commands (user: jim, artifact:
{} [25b6e38e97])

2010-05-12 [5e5c0f7d55] End of day commit (user: jim, artifact: [d1a1d31
2010-05-10 [e924ca3525] End of day update (user: jim, artifact: [7cd1907
2010-05-09 [0abb95b046] Intermediate commit, not done with basic command
{} (user: jim, artifact: [6f7bcd48b9])
```

```

2010-05-07 [6921e453cd] Update outline & book corrections (user: jim,
{}          artifact: [4eff85c793])
2010-05-03 [158492516c] Moved to clone repository (user: jim, artifact:
{}          [23b729cb66])
2010-05-03 [1a403c87fc] Update before moving to server (user: jim, artifact:
{}          [706a9d394d])
2010-04-30 [fa5b9247bd] Working on chapter 1 (user: jim, artifact:
{}          [7bb188f0c6])
2010-04-29 [51be6423a3] Update outline (user: jim, artifact: [7cd39dfa06])
2010-04-27 [39bc728527] [1665c78d94] Ticket Use (user: jim, artifact:
{}          [1f82aaf41c])
2010-04-26 [497b93858f] Update to catch changes in outline (user: jim,
{}          artifact: [b870231e48])
2010-04-25 [8fa0708186] Initial Commit (user: jim, artifact: [34a460a468])

```

7.4.4 timeline

This prints out the timeline of the project in various ways. The command would be useful if you were building a GUI front end for Fossil and wanted to display the timeline. You could issue this command and get the result back and display it in your UI. There are a number of options in the command to control the listing.

```

$ fossil help timeline
Usage: fossil timeline ?WHEN? ?CHECKIN|DATETIME? ?OPTIONS?

```

Print a summary of activity going backwards in date and time specified or from the current date and time if no arguments are given. The WHEN argument can be any unique abbreviation of one of these keywords:

```

before

```

```

after
descendants | children
ancestors | parents

```

The CHECKIN can be any unique prefix of 4 characters or more. You can also say "current" for the current version.

DATETIME may be "now" or "YYYY-MM-DDTHH:MM:SS.SSS". If in year-month-day form, it may be truncated, the "T" may be replaced by a space, and it may also name a timezone offset from UTC as "-HH:MM" (westward) or "+HH:MM" (eastward). Either no timezone suffix or "Z" means UTC.

Options:

<code>-n --limit N</code>	If N is positive, output the first N entries. If N is negative, output the first -N lines. If N is zero, no limit. Default is -20 meaning 20 lines.
<code>-p --path PATH</code>	Output items affecting PATH only. PATH can be a file or a sub directory.
<code>--offset P</code>	skip P changes
<code>--sql</code>	Show the SQL used to generate the timeline
<code>-t --type TYPE</code>	Output items from the given types only, such as: ci = file commits only e = technical notes only t = tickets only w = wiki commits only
<code>-v --verbose</code>	Output the list of files changed by each commit and the type of each change (edited, deleted, etc.) after the check-in comment.
<code>-W --width N</code>	Width of lines (default is to auto-detect). N must be either greater than 20 or it must be zero 0 to indicate no limit, resulting in a single line per entry.
<code>-R REPO_FILE</code>	Specifies the repository db to use. Default is the current checkout's repository.

7.4.5 wiki

This command allows you to have command line control of the wiki. Again this is useful if you were writing a shell to control Fossil or wanted to add a number of computer generated pages to the Wiki.

```

$ fossil help wiki
Usage: fossil wiki (export|create|commit|list) WikiName

```

Run various subcommands to work with wiki entries or tech notes.

```
fossil wiki export ?OPTIONS? PAGENAME ?FILE?
fossil wiki export ?OPTIONS? -t|--technote DATETIME|TECHNOTE-ID ?FILE?
```

Sends the latest version of either a wiki page or of a tech note to the given file or standard output. A filename of "-" writes the output to standard output. The directory parts of the output filename are created if needed.

If PAGENAME is provided, the named wiki page will be output.

Options:

```
--technote|-t DATETIME|TECHNOTE-ID
    Specifies that a technote, rather than a wiki page,
    will be exported. If DATETIME is used, the most
    recently modified tech note with that DATETIME will
    output.
-h|--html
    The body (only) is rendered in HTML form, without
    any page header/footer or HTML/BODY tag wrappers.
-H|--HTML
    Works like -h|--html but wraps the output in
    <html><body>...</body></html>.
-p|--pre
    If -h|--H is used and the page or technote has
    the text/plain mimetype, its HTML-escaped output
    will be wrapped in <pre>...</pre>.
```

```
fossil wiki (create|commit) PAGENAME ?FILE? ?OPTIONS?
```

Create a new or commit changes to an existing wiki page or technote from FILE or from standard input. PAGENAME is the name of the wiki entry or the timeline comment of the technote.

Options:

```
-M|--mimetype TEXT-FORMAT
    The mime type of the update.
    Defaults to the type used by
    the previous version of the
    page, or text/x-fossil-wiki.
    Valid values are: text/x-fossil-wiki,
    text/x-markdown and text/plain. fossil,
    markdown or plain can be specified as
    synonyms of these values.
-t|--technote DATETIME
    Specifies the timestamp of
    the technote to be created or
    updated. When updating a tech note
    the most recently modified tech note
    with the specified timestamp will be
```

	updated.
<code>-t --technote TECHNOTE-ID</code>	Specifies the technote to be updated by its technote id.
<code>--technote-tags TAGS</code>	The set of tags for a technote.
<code>--technote-bgcolor COLOR</code>	The color used for the technote on the timeline.

```
fossil wiki list ?OPTIONS?
fossil wiki ls ?OPTIONS?
```

Lists all wiki entries, one per line, ordered case-insensitively by name.

Options:

<code>-t --technote</code>	Technotes will be listed instead of pages. The technotes will be in order of timestamp with the most recent first.
<code>-s --show-technote-ids</code>	The id of the tech note will be listed along side the timestamp. The tech note id will be the first word on each line. This option only applies if the <code>--technote</code> option is also specified.

DATE_{TIME} may be "now" or "YYYY-MM-DDTHH:MM:SS.SSS". If in year-month-day form, it may be truncated, the "T" may be replaced by a space, and it may also name a timezone offset from UTC as "-HH:MM" (westward) or "+HH:MM" (eastward). Either no timezone suffix or "Z" means UTC.

The "Sandbox" wiki pseudo-page is a special case. Its name is checked case-insensitively and either "create" or "commit" may be used to update its contents.

7.5 Advanced

These are commands that you will rarely have to use. These are functions that are needed to do very complicated things with Fossil. If you have to use these you are probably way beyond the audience for this book.

7.5.1 scrub

This is used to removed sensitive information like passwords from a repository. This allows you to then send the whole repository to someone else for their use.

```
$ fossil help scrub
Usage: fossil scrub ?OPTIONS? ?REPOSITORY?
```

The command removes sensitive information (such as passwords) from a repository so that the repository can be sent to an untrusted reader.

By default, only passwords are removed. However, if the `--verily` option is added, then private branches, concealed email addresses, IP addresses of correspondents, and similar privacy-sensitive fields are also purged. If the `--private` option is used, then only private branches are removed and all other information is left intact.

This command permanently deletes the scrubbed information. THE EFFECTS OF THIS COMMAND ARE NOT REVERSIBLE. USE WITH CAUTION!

The user is prompted to confirm the scrub unless the `--force` option is used.

Options:

```
--force      do not prompt for confirmation
--private    only private branches are removed from the repository
--verily     scrub real thoroughly (see above)
```

7.5.2 search

This is used to search the timeline entries for a pattern. This can also be done in your browser on the timeline page.

```
$ fossil help search
Usage: fossil search [-all|-a] [-limit|-n #] [-width|-W #] pattern...
```

Search for timeline entries matching all words provided on the command line. Whole-word matches scope more highly than partial matches.

Note: The command only search the EVENT table. So it will only display check-in comments or other comments that appear on an unaugmented timeline. It does not search document text or forum messages.

Outputs, by default, some top-N fraction of the results. The `-all` option can be used to output all matches, regardless of their search score. The `-limit` option can be used to limit the number of entries returned. The `-width` option can be used to set the output width used when printing matches.

Options:

```
-a|--all           Output all matches, not just best matches.
-n|--limit N      Limit output to N matches.
-W|--width WIDTH  Set display width to WIDTH columns, 0 for
                  unlimited. Defaults the terminal's width.
```

7.5.3 sha3sum

This can compute the sha1 value for a particular file. These sums are the labels that Fossil uses on all objects and should be unique for any file.

```
$ fossil help sha3sum
Usage: fossil sha3sum FILE...
```

Compute an SHA3 checksum of all files named on the command-line. If a file is named "-" then take its content from standard input.

To be clear: The official NIST FIPS-202 implementation of SHA3 with the added 01 padding is used, not the original Keccak submission.

Options:

```
--224             Compute a SHA3-224 hash
--256             Compute a SHA3-256 hash (the default)
--384             Compute a SHA3-384 hash
--512             Compute a SHA3-512 hash
--size N         An N-bit hash. N must be a multiple of 32 between
                  128 and 512.
-h, --dereference If FILE is a symbolic link, compute the hash on
                  the object pointed to, not on the link itself.
```

See also: md5sum, shasum

7.5.4 configuration

This command allows you to save or load a custom configuration of Fossil.

```
$ fossil help configuration
Usage: fossil configuration METHOD ... ?OPTIONS?
```

Where METHOD is one of: export import merge pull push reset. All methods accept the -R or --repository option to specify a repository.

```
fossil configuration export AREA FILENAME
```

Write to FILENAME exported configuration information for AREA.

AREA can be one of:

```
all email interwiki project shun skin
ticket user alias subscriber
```

`fossil configuration import FILENAME`

Read a configuration from FILENAME, overwriting the current configuration.

`fossil configuration merge FILENAME`

Read a configuration from FILENAME and merge its values into the current configuration. Existing values take priority over values read from FILENAME.

`fossil configuration pull AREA ?URL?`

Pull and install the configuration from a different server identified by URL. If no URL is specified, then the default server is used. Use the `--overwrite` flag to completely replace local settings with content received from URL.

`fossil configuration push AREA ?URL?`

Push the local configuration into the remote server identified by URL. Admin privilege is required on the remote server for this to work. When the same record exists both locally and on the remote end, the one that was most recently changed wins.

`fossil configuration reset AREA`

Restore the configuration to the default. AREA as above.

`fossil configuration sync AREA ?URL?`

Synchronize configuration changes in the local repository with the remote repository at URL.

Options:

```
-R|--repository FILE          Extract info from repository FILE
```

See also: `settings`, `unset`

7.5.5 descendants

This is used to find where the checked out files are in the time line.

```
$ fossil help descendants
```

```
Usage: fossil descendants ?CHECKIN? ?OPTIONS?
```

Find all leaf descendants of the check-in specified or if the argument is omitted, of the check-in currently checked out.

Options:

-R --repository FILE	Extract info from repository FILE
-W --width N	Width of lines (default is to auto-detect). Must be greater than 20 or else 0 for no limit, resulting in a one line per entry.

See also: `finfo`, `info`, `leaves`

Chapter 8

Pikchr

This section needs completion.

Chapter 9

TH1 Scripting language

TH1 is designed to be the scripting language of Fossil. It is used in several places within Fossil. As a typical user, you will probably first meet TH1 when customizing the ticket system (see 5.7 for an overview of the ticket system and 6.1 on how to customize the ticket system). You can customize the behaviour of the ticket web pages that show

- the form for creating new tickets
- existing tickets
- the form for editing existing tickets
- the list of ticket reports

If you are changing the overall look and feel of Fossil's web pages, you can use TH1 when designing the skin. Both the "Content Header" and the "Content Footer" can contain TH1 code (also the optional Javascript for the footer). This includes the possibility to change Fossil's default "Content Security Policy" (CSP) which can be overridden by custom TH1 code.

Further, 8 scripts can include TH1 scripts. However, this is restricted to the Fossil command `pikchr` on the command line. TH1 scripts are not enabled within Fossil's web pages (also including `/pikchrshow`).

Finally, there are two places where TH1 can be used when Fossil is explicitly enabled to do so at compile-time:

- when compiled with the `--with-th1-docs` option, Fossil's embedded documentation facility via the `/doc` web pages can contain TH1 scripts. It is a compile-time option since it opens Fossil scripting to anyone who has commit capabilities on the repository and thus can commit documentation files containing arbitrary TH1 code. This can be dangerous and must therefore be enabled already at compile-time. In addition, this option must be enabled in Fossil's settings (option `th1-docs` in the Admin menu under `Settings`; see section 6). Note, that TH1 scripts only work in

embedded documentation files having the mime type “application/x-th1”. The ideal situation is to use a file having a file name ending in `.th1` and containing HTML. The TH1 code is then embedded into the special `<th1>` tag.

- when compiled with the `--with-th1-hooks` option, TH1 scripts can be used to monitor, create, alter, or cancel the execution of Fossil commands and web pages (additionally, the option `th1-hooks` must be enabled at runtime (in the `Admin` menu under `Settings`)). Hooks can then be set to run before and/or after Fossil command execution, and before and/or after the rendering of a web page.

9.1 Introduction to TH1

9.1.1 TH1 is a Tcl-like language

TH1 is a string-based command language closely based on the Tcl language. The language has only a few fundamental constructs and relatively little syntax which is meant to be simple. TH1 is an interpreted language and it is parsed, compiled and executed when the script runs. In Fossil, TH1 scripts are typically run to build web pages, often in response to web form submissions.

The basic mechanisms of TH1 are all related to strings and string substitutions. The TH1/Tcl way of doing things is a little different from some other programming languages with which you may already be familiar, so it is worth making sure you understand the basic concepts.

9.2 The Hello world program

The traditional starting place for a language introduction is the classic “Hello, World” program. In TH1 this program has only a single line:

```
puts "Hello, world\n"
```

The command to output a string in TH1 is the `puts` command (the same command name as in Tcl). A single unit of text after the `puts` command will be printed to the output stream. If the string has more than one word, you must enclose the string in double quotes or curly brackets. A set of words enclosed in quotes or curly brackets is treated as a single unit, while words separated by white space are treated as multiple arguments to the command.

9.3 TH1 structure and syntax

9.3.1 Datatypes

TH1 has at its core only a single data type which is string. All values in TH1 are strings, variables hold strings and the procedures return strings. Strings in TH1 consist of single byte characters and are zero terminated. Characters outside of the ASCII range, i.e. characters in the 0x80-0xff range have no TH1 meaning at all: they are not considered digits or letters, nor are they considered white space.

Depending on context, TH1 can interpret a string in four different ways. First of all, strings can be just that: text consisting of arbitrary sequences of characters. Second, a string can be considered a list, an ordered sequence of words separated by white space. Third, a string can be a command. A command is a list where the first word is interpreted as the name of a command, optionally followed by further argument words. Fourth, and last, a string can be interpreted as an expression.

The latter three interpretations of strings are discussed in more detail below.

9.3.2 Lists

A list in TH1 is an ordered sequence of items, or words separated by white space. In TH1 the following characters constitute white space:

```
' '      0x20
'\t'     0x09
'\n'     0x0A
'\v'     0x0B
'\f'     0x0C
'\r'     0x0D
```

A word can be any sequence of characters delimited by white space. It is not necessary that a word is alphanumeric at all: ".%*" is a valid word in TH1. If a word needs to contain embedded white space characters, it needs to be quoted, with either a double quotes or with opening/closing curly brackets. Quoting has the effect of grouping the quoted content into a single list element.

Words cannot start with one of the TH1 special characters { } [] \ ; and ". Note that a single quote ` is not a special character in TH1. To use one of these characters to start a word it must be escaped, which is discussed further later on.

TH1 offers several built-in commands for working with lists, such as counting the number of words in a list, retrieving individual words from the list by index and appending new items to a list. These commands are discussed in the section "Working with lists".

9.3.3 Commands

TH1 casts everything into the mold of a command, even programming constructs like variable assignment and procedure definition. TH1 adds a tiny amount of syntax needed to properly invoke commands, and then it leaves all the hard work up to the command implementation.

Commands are a special form of list. The basic syntax for a TH1 command is:

```
command arg1 arg2 arg3 ...
```

The command is either the name of a built-in command or a TH1 procedure.

White space is used to separate the command name and its arguments, and a newline character or semicolon is used to terminate a command. TH1 comments are lines with a # character at the beginning of the line, or with a # character after the semicolon terminating a command.

9.3.4 Grouping & substitution

TH1 does not interpret the arguments to the commands except to perform grouping, which allows multiple words in one argument, and substitution, which is used to deal with special characters, to fetch variables and to perform nested command calls. Hence, the behavior of the TH1 command processor can be summarized in three basic steps:

1. Argument grouping.
2. Value substitution of backslash escapes, variables and nested commands
3. Command invocation.

Note that there is no step to evaluate the arguments to a command. After substitution, arguments are passed verbatim to the command and it is up to the command to evaluate its arguments as needed.

9.3.5 Argument grouping

TH1 has two mechanisms for grouping multiple words into a single item:

- Double quotes, " "
- Curly brackets, { }

Whilst both have the effect of grouping a set of words, they have different impact on the next phase of substitution. In brief, double quotes only group their content and curly brackets group and prevent all substitution on their content.

Grouping proceeds from left to right in the string and is not affected by the subsequent substitution. If a substitution leads to a string which would be grouped differently, it has no effect, as the grouping has already been decided in the preceding grouping phase.

9.3.6 Value substitutions

TH1 performs three different substitutions (see the `th.c/thSubstWord` code for details)

- Backslash escape substitution
- Variable substitution
- Nested command substitution

Like grouping, substitution proceeds from left to right and is performed only once: if a substitution leads to a string which could again be substituted such this not happen.

9.3.7 Backslash escape substitution

In general, the backslash (`\`) disables substitution for the single character immediately following the backslash. Any character immediately following the backslash will stand as literal. This is useful to escape the special meaning of the characters `{ }` `[]` `\` `;` and `"`.

There are two specific strings which are replaced by specific values during the substitution phase. A backslash followed by the letter `n` gets replaced by the newline character, as in `C`. A backslash followed by the letter `x` and two hexadecimal digits gets replaced by the character with that value, i.e. writing `\x20` is the same as writing a space. Note that the `\x` substitution does not “keep going” as long as it has hex digits as in `Tcl`, but insists on two digits. The word `\x2121` is not a single exclamation mark, but the 3 letter word `!21`.

9.3.8 Variable substitution

Like any programming language, TH1 has a concept of variables. TH1 variables are named containers that hold string values. Variables are discussed in more detail later in this document, for now we limit ourselves to variable substitution.

The dollar sign (`$`) may be used as a special shorthand form for substituting variable values. If `$` appears in an argument that isn't enclosed in curly brackets then variable substitution will occur. The characters after the `$`, up to the first character that isn't a number, letter, or underscore, are taken as a variable name and the string value of that variable is substituted for the name. For example, if variable `foo` has the value `test`, then the command `puts $foo.c` is equivalent to the command:

```
puts test.c
```

There are two special forms for variable substitution. If the next character after the name of the variable is an open parenthesis, then the variable is assumed to be an array name, and all of the characters between the open parenthesis and the next close parenthesis are taken as an index into the array. Command substitutions and variable substitutions are performed on the information between

the parentheses before it is used as an index. For example, if the variable `x` is an array with one element named `first` and value `87` and another element named `14` and value `more`, then the command

```
puts xyz$x(first)zyx
```

is equivalent to the command

```
puts xyz87zyx
```

If the variable `index` has the value `'14'`, then the command

```
puts xyz$x($index)zyx
```

is equivalent to the command

```
puts xyzmorezyx
```

See the section `Variables and arrays` below for more information on arrays.

The second special form for variables occurs when the dollar sign is followed by an open curly bracket. In this case the variable name consists of all the characters up to the next curly bracket. Array references are not possible in this form: the name between curly brackets is assumed to refer to a scalar variable. For example, if variable `foo` has the value `'test'`, then the command

```
set a abc${foo}bar
```

is equivalent to the command

```
set a abctestbar
```

A dollar sign followed by something other than a letter, digit, underscore, or left parenthesis is treated as a literal dollar sign. The following prints a single character `$`.

```
puts x $
```

9.3.9 Command substitution

The last form of substitution is command substitution. A nested command is delimited by square brackets, `[]`. The TH1 interpreter takes everything between the brackets and evaluates it as a command. It rewrites the outer command by replacing the square brackets and everything between them with the result of the nested command.

Example:

```
puts `[string length foobar]
```

```
=> 6
```

In the example, the nested command is: `string length foobar`. This command returns the length of the string `foobar`. The nested command runs first. Then, command substitution causes the outer command to be rewritten as if it were:

```
puts 6
```

If there are several cases of command substitution within a single command, the interpreter processes them from left to right. As each right bracket is encountered, the command it delimits is evaluated. This results in a sensible ordering in which nested commands are evaluated first so that their result can be used in arguments to the outer command.

9.3.10 Argument grouping revisited

During the substitution phase of command evaluation, the two grouping operators, the curly bracket and the double quote are treated differently by the TH1 interpreter.

Grouping words with double quotes allows substitutions to occur within the double quotes. A double quote is only used for grouping when it comes after white space. The string `a"b"` is a normal 4 character string, and not the two character string `ab`.

```
puts a"b"
```

```
=> a"b"
```

Grouping words within curly brackets disables substitution within the brackets. Again, A opening curly bracket is only used for grouping when it comes after white space. Characters within curly brackets are passed to a command exactly as written, and not even backslash escapes are processed.

Note that curly brackets have this effect only when they are used for grouping (i.e. at the beginning and end of a sequence of words). If a string is already grouped, either with double quotes or curly brackets, and the curly brackets occur in the middle of the grouped string (e.g. `"foo{bar}"`), then the curly brackets are treated as regular characters with no special meaning. If the string is grouped with double quotes, substitutions will occur within the quoted string, even between the brackets.

The square bracket syntax used for command substitution does not provide grouping. Instead, a nested command is considered part of the current group. In the command below, the double quotes group the last argument, and the nested command is just part of that group:

```
puts "The length of $s is [string length $s]."
```

If an argument is made up of only a nested command, you do not need to group it with double-quotes because the TH1 parser treats the whole nested command as part of the group. A nested command is treated as an unbroken

sequence of characters, regardless of its internal structure. It is included with the surrounding group of characters when collecting arguments for the main command.

9.3.11 Summary

The following rules summarize the fundamental mechanisms of grouping and substitution that are performed by the TH1 interpreter before it invokes a command:

- Command arguments are separated by white space, unless arguments are grouped with curly brackets or double quotes as described below.
- Grouping with curly brackets, `{ }`, prevents substitutions. Curly brackets nest. The interpreter includes all characters between the matching left and right brace in the group, including newlines, semicolons, and nested curly brackets. The enclosing (i.e., outermost) curly brackets are not included in the group's value.
- Grouping with double quotes, `" "`, allows substitutions. The interpreter groups everything until another double quote is found, including newlines and semicolons. The enclosing quotes are not included in the group of characters. A double-quote character can be included in the group by quoting it with a backslash, (i.e. `\"`).
- Grouping decisions are made before substitutions are performed, which means that the values of variables or command results do not affect grouping.
- A dollar sign, `$`, causes variable substitution. Variable names can be any length, and case is significant. If variable references are embedded into other strings, or if they include characters other than letters, digits, and the underscore, they can be distinguished with the `${varname}` syntax.
- Square brackets, `[]`, cause command substitution. Everything between the brackets is treated as a command, and everything including the brackets is replaced with the result of the command. Nesting is allowed.
- The backslash character, `\`, is used to quote special characters. You can think of this as another form of substitution in which the backslash and the next character or group of characters is replaced with a new character.
- Substitutions can occur anywhere unless prevented by curly bracket grouping. Part of a group can be a constant string, and other parts of it can be the result of substitutions. Even the command name can be affected by substitutions.
- A single round of substitutions is performed before command invocation. The result of a substitution is not interpreted a second time. This rule is important if you have a variable value or a command result that

contains special characters such as spaces, dollar signs, square brackets, or curly brackets. Because only a single round of substitution is done, you do not have to worry about special characters in values causing extra substitutions.

9.3.12 Caveats

- A common error is to forget a space between arguments when grouping with curly brackets or quotes. This is because white space is used as the separator, while the curly brackets or quotes only provide grouping. If you forget the space, you will get syntax errors about the wrong number of arguments being applied. The following is an error because of the missing space between } and {:

```
if {$x > 1}{ puts "x = $x" }
```

- When double quotes are used for grouping, the special effect of curly brackets is turned off. Substitutions occur everywhere inside a group formed with double quotes. In the next command, the variables are still substituted:

```
set x xvalue

set y "foo {$x} bar"

=> foo {xvalue} bar
```

- Spaces are not required around the square brackets used for command substitution. For the purposes of grouping, the interpreter considers everything between the square brackets as part of the current group.

9.4 TH1 expressions

The TH1 interpreter itself does not evaluate math expressions. TH1 just does grouping, substitutions and command invocations. However, several built-in commands see one or more of their arguments as expressions and request the interpreter to calculate the value of such expressions.

The `expr` command is the simplest such command and is used to parse and evaluate expressions:

```
puts [expr 7.4/2]

=> 3.7
```

Note that an expression can contain white space, but if it does it must be grouped in order to be recognized as a single argument.

Within the context of expression evaluation TH1 works with three datatypes: two types of number, integer and floating point, and string. Integer values are promoted to floating point values as needed. The Boolean values True and False are represented by the integer values 1 and 0 respectively. The implementation of expr is careful to preserve accurate numeric values and avoid unnecessary conversions between numbers and strings.

Before expression evaluation takes place, both variable and command substitution is performed on the expression string. Hence, you can include variable references and nested commands in math expressions, even if the expression string was originally quoted with curly brackets. Note that backslash escape substitution is not performed by the expression evaluator.

A TH1 expression consists of a combination of operands, operators, and parentheses. White space may be used between the operands and operators and parentheses; it is ignored by the expression processor. Where possible, operands are interpreted as integer values. If an operand is not in integer format, then it is treated as a floating-point number if that is possible. Floating-point numbers may be specified in any of the ways accepted by an ANSI-compliant C compiler. For example, all of the following are valid floating-point numbers: 2.1, 3., 6e4, 7.91e+16. If no numeric interpretation is possible, then an operand is left as a string (and only a limited set of operators may be applied to it).

Operands may be specified in any of the following ways:

- As a numeric value, either integer or floating-point.
- As a string enclosed in curly brackets. The characters between the opening bracket and matching closing bracket are used as the operand without any substitutions.
- As a string enclosed in double quotes. The expression parser performs variable and command substitutions on the information between the quotes, and uses the resulting value as the operand.
- As a TH1 variable, using standard \$ notation. The variable's value is used as the operand.
- As a TH1 command enclosed in square brackets. The command will be executed and its result will be used as the operand.

Where substitutions occur above (e.g. inside quoted strings), they are performed by the expression processor. However, an additional layer of substitution may already have been performed by the command parser before the expression processor was called. As discussed below, it is usually best to enclose expressions in curly brackets to prevent the command parser from performing substitutions on the contents.

The valid operators are listed below, grouped in decreasing order of precedence:

All of the binary operators group left-to-right within the same precedence level. For example, the expression `4*2 < 7` evaluates to 0.

All internal computations involving integers are done with the C type `int`, and all internal computations involving floating-point are done with the C type `double`. Conversion among internal representations for integer, floating-point, and string operands is done automatically as needed. For arithmetic computations, integers are used until some floating-point number is introduced, after which floating-point is used.

String values may be used as operands of the comparison operators, although the expression evaluator tries to do comparisons as integer or floating-point when it can. If one of the operands of a comparison is a string and the other has a numeric value, the numeric operand is converted back to a string.

9.5 TH1 variables

Like almost all programming languages TH1 has the concept of variables. TH1 variables bind a name to a string value. A variable name must be unique in its scope, either the global scope or a local scope. TH1 supports two types of variables: scalars and arrays.

TH1 allows the definition of variables and the use of their values either through `$`-style variable substitution, the `set` command, or a few other mechanisms. Variables need not be declared: a new variable will automatically be created each time a new variable name is used.

9.5.1 Working with variables

TH1 has two key commands for working with variables, `set` and `unset`:

```
set varname ?value?
```

```
unset varname
```

```
info exists varname
```

The `set` command returns the value of variable `varname`. If the variable does not exist, then an error is thrown. If the optional argument `value` is specified, then the `set` command sets the value of `varname` to `value`, creating a new variable in the current scope if one does not already exist, and returns its value.

The `unset` command removes a variable from its scope. The argument `varname` is a variable name. If `varname` refers to an element of an array, then that element is removed without affecting the rest of the array. If `varname` consists of an array name with no parenthesized index, then the entire array is deleted. The `unset` command returns an empty string as result. An error occurs if the variable doesn't exist.

The `info exists` command returns 1 if the variable named `varname` exists in the current scope, either the global scope or the current local scope, and returns 0 otherwise.

9.5.2 Scalar variables and array variables

TH1 supports two types of variables: scalars and arrays. A scalar variable has a single value, whereas an array variable can have any number of elements, each with a name (called its “index”) and a value. TH1 arrays are one-dimensional associative arrays, i.e. the index can be any single string.

If `varname` contains an open parenthesis and ends with a close parenthesis, then it refers to an array element: the characters before the open parenthesis are the name of the array, and the characters between the parentheses are the index within the array. Otherwise `varname` refers to a scalar variable.

For example, the command `set x(first) 44` will modify the element of `x` whose index is `first` so that its new value is 44. Two-dimensional arrays can be simulated in TH1 by using indices that contain multiple concatenated values. For example, the commands

```
set a(2,3) 1
```

```
set a(3,6) 2
```

set the elements of `a` whose indices are 2,3 and 3,6.

In general, array elements may be used anywhere in TH1 that scalar variables may be used. If an array is defined with a particular name, then there may not be a scalar variable with the same name. Similarly, if there is a scalar variable with a particular name then it is not possible to make array references to the variable. To convert a scalar variable to an array or vice versa, remove the existing variable with the `unset` command.

9.5.3 Variable scope

Variables exist in a scope. The TH1 interpreter maintains a global scope that is available to and shared by all commands executed by it. Each invocation of a user defined command creates a new local scope. This local scope holds the arguments and local variables of that user command invocation and only exists as long as the user command is executing.

If not in the body of user command, then references to `varname` refer to a global variable, i.e. a variable in the global scope. In contrast, in the body of a user defined command references to `varname` refer to a parameter or local variable of the command. However, in the body of a user defined command, a global variable can be explicitly referred to by preceding its name by `::`.

TH1 offers a special command to access variables not in the local scope of the current command but in the local scope of the call chain of commands that leads to the current command. This is the `upvar` command:

```
upvar ?frame? othervar myvar ?othervar myvar ...?
```

The `upvar` command arranges for one or more local variables in the current procedure to refer to variables in an enclosing procedure call, or to global variables. If `frame` is an integer, then it gives a distance (up the command calling stack) to move. The argument `frame` may be omitted if `othervar` is not an integer (`frame` then defaults to 1). For each `othervar` argument, the `upvar` command makes the variable by that name in the local scope identified by the `frame` offset accessible in the current procedure by the name given in the corresponding `myvar` argument. The variable named by `othervar` need not exist at the time of the call; it will be created the first time `myvar` is referenced, just like an ordinary variable. The `upvar` command is only meaningful from within user defined command bodies. Neither `othervar` nor `myvar` may refer to an element of an array. The `upvar` command returns an empty string. The `upvar` command simplifies the implementation of call-by-name procedure calling and also makes it easier to build new control constructs as TH1 commands. For example, consider the following procedure:

```
proc incr {name} {
    upvar $name x
    set x [expr $x+1]
}
```

`incr` is invoked with an argument giving the name of a variable, and it adds one to the value of that variable.

9.6 TH1 commands, scripts and program flow

In TH1 there is actually no distinction between commands (often known as ‘statements’ and ‘functions’ in other languages) and “syntax”. There are no reserved words (like `if` and `while`) as exist in C, Java, Python, Perl, etc. When the TH1 interpreter starts up there is a list of built-in, known commands that the interpreter uses to parse a line. These commands include `for`, `set`, `puts`, and so on. They are, however, still just regular TH1 commands that obey the same syntax rules as all TH1 commands, both built-in, and those that you create yourself with the `proc` command.

9.6.1 Commands revisited

Like Tcl, TH1 is build up around commands. A command does something for you, like outputting a string, computing a math expression, or generating HTML to display a widget on the screen.

Commands are a special form of list. The basic syntax for a TH1 command is:

```
command arg1 arg2 arg3 ...
```

The command is either the name of a built-in command or a TH1 procedure.

White space is used to separate the command name and its arguments, and a newline character or semicolon is used to terminate a command. TH1 comments are lines with a # character at the beginning of the line, or with a # character after the semicolon terminating a command.

9.6.2 Scripts

Normally, control in TH1 flows from one command to the next. The next command is either in the same list (if the current command is terminated with a semicolon) or in the next input line. A TH1 program is thus a TH1 list of commands.

Such a list of commands is referred to as a “script”. A script is hence a self contained code fragment containing one or more commands. The commands in a script are analogous to statements in other programming languages.

Some commands take one or more scripts as arguments and run those scripts zero or more times depending on the other arguments. For example, the if command executes either the then script or the else script once, depending on the if expression being true or false. The command that takes a script will perform the normal grouping and substitution as part of executing the script.

Note that the script always needs to be enclosed in curly brackets to prevent substitution taking place twice: once as part of the execution of the top level command and once again when preparing the script. Forgetting to enclose a script argument in curly brackets is common source of errors.

A few commands (return, error, break and continue) immediately stop execution of the current script instead of passing control to the next command in the list. Control is instead returned to the command that initiated the execution of the current script.

9.6.3 Command result codes

Each command produces two results: a result code and a string. The code indicates whether the command completed successfully or not, and the string gives additional information. The valid codes are defined in `th.h`, and are:

TH1 programmers do not normally need to think about return codes, since `TH1_OK` is almost always returned. If anything else is returned by a command, then the TH1 interpreter immediately stops processing commands and returns to its caller. If there are several nested invocations of the TH1 interpreter in progress, then each nested command will usually return the error to its caller,

until eventually the error is reported to the top-level application code. The application will then display the error message for the user.

In a few cases, some commands will handle certain “error” conditions themselves and not return them upwards. For example, the `for` command checks for the `TH1_BREAK` code; if it occurs, `for` stops executing the body of the loop and returns `TH1_OK` to its caller. The `for` command also handles `TH1_CONTINUE` codes and the procedure interpreter handles `TH1_RETURN` codes. The `catch` command allows TH1 programs to catch errors and handle them without aborting command interpretation any further.

9.6.4 Flow control commands

The flow control commands in TH1 are:

```
if expr1 body1 ?elseif expr2 body2? ? ?else? bodyN?
```

```
for init condition incr script
```

```
break    ?value?
```

```
continue ?value?
```

```
error    ?value?
```

```
catch script ?varname?
```

Below each command is discussed in turn

The `if` command has the following syntax:

```
if expr1 body1 ?elseif expr2 body2? ? ?else? bodyN?
```

The `expr` arguments are expressions, the `body` arguments are scripts and the `elseif` and `else` arguments are keyword constant strings. The `if` command optionally executes one of its `body` scripts.

The `expr` arguments must evaluate to an integer value. If it evaluates to a non-zero value the following `body` script is executed and upon return from that script processing continues with the command following the `if` command. If an `expr` argument evaluates to zero, its `body` script is skipped and the next option is tried. When there are no more options to try, processing also continues with the next command.

The `if` command returns the value of the executed script, or “0” when no script was executed.

The `for` command has the following syntax:

```
for init condition incr body
```

The `init`, `incr` and `body` arguments are all scripts. The `condition` argument is an expression yielding an integer result. The `for` command is a looping command, similar in structure to the C `for` statement.

The `for` command first invokes the TH1 interpreter to execute `init`. Then it repeatedly evaluates `condition` as an expression; if the result is non-zero it invokes the TH1 interpreter on `body`, then invokes the TH1 interpreter on `incr`, then repeats the loop. The command terminates when `test` evaluates to zero.

If a `continue` command is invoked within execution of the `body` script then any remaining commands in the current execution of `body` are skipped; processing continues by invoking the TH1 interpreter on `incr`, then evaluating `condition`, and so on. If a `break` command is invoked within `body` or `next`, then the `for` command will return immediately. The operation of `break` and `continue` are similar to the corresponding statements in C.

The `for` command returns an empty string.

The `break` command has the following syntax:

```
break ?value?
```

The `break` command returns immediately from the current procedure (or top-level command), with `value` as the return value and `TH1_BREAK` as the result code. If `value` is not specified, the string "break" will be returned as result.

The `continue` command has the following syntax:

```
continue ?value?
```

The `continue` command returns immediately from the current procedure (or top-level command), with `value` as the return value and `TH1_CONTINUE` as the result code. If `value` is not specified, the string "continue" will be returned as result.

The `error` command has the following syntax:

```
error ?value?
```

The `error` command returns immediately from the current procedure (or top-level command), with `value` as the return value and `TH1_ERROR` as the result code. If `value` is not specified, the string "error" will be returned as result.

The `catch` command has the following syntax:

```
catch script ?varname?
```

The `catch` command may be used to prevent errors from aborting command interpretation. The `catch` command calls the TH1 interpreter recursively to execute `script`, and always returns a `TH1_OK` code, regardless of any errors that might occur while executing `script`.

The return value from `catch` is a decimal string giving the code returned by the TH1 interpreter after executing `script`. This will be 0(`TH1_OK`) if there were

no errors in command; otherwise it will have a non-zero value corresponding to one of the exceptional result codes. If the varname argument is given, then it gives the name of a variable; catch sets the value of the variable to the string returned from running script (either a result or an error message).

9.6.5 Creating user defined commands

The `proc` command creates a new command. The syntax for the `proc` command is:

```
proc name args body
```

The `proc` command creates a new TH1 command procedure, name, replacing any existing command there may have been by that name. Whenever the new command is invoked, the contents of body will be executed by the TH1 interpreter.

The parameter `args` specifies the formal arguments to the procedure. It consists of a list, possibly empty, each of whose elements specifies one argument. Each argument specifier is also a list with either one or two fields. If there is only a single field in the specifier, then it is the name of the argument; if there are two fields, then the first is the argument name and the second is its default value. Curly brackets and backslashes may be used in the usual way to specify complex default values.

The `proc` command returns the null string.

9.6.6 Execution of user defined commands

If a command is a user defined command (i.e. a command created with the `proc` command), then the TH1 interpreter creates a new local variable context, binds the formal arguments to their actual values (i.e. TH1 uses call by value exclusively) and loads the body script. Execution then proceeds with the first command in that script. Execution ends when the last command has been executed or when one of the returning commands is executed. When the script ends, the local variable context is deleted and processing continues with the next command after the user defined command.

More in detail, when a user defined command is invoked, a local variable is created for each of the formal arguments to the procedure; its value is the value of corresponding argument in the invoking command or the argument's default value. Arguments with default values need not be specified in a procedure invocation. However, there must be enough actual arguments for all the formal arguments that don't have defaults, and there must not be any extra actual arguments.

There is one special case to permit procedures with variable numbers of arguments. If the last formal argument has the name `args`, then a call to the procedure may contain more actual arguments than the procedure has formals. In this case,

all of the actual arguments starting at the one that would be assigned to `args` are combined into a list (as if the `list` command had been used); this combined value is assigned to the local variable `args`.

When `body` is being executed, variable names normally refer to local variables, which are created automatically when referenced and deleted when the procedure returns. One local variable is automatically created for each of the procedure's arguments. Global variables can be accessed by using the `::` syntax.

When a procedure is invoked, the procedure's return value is the value specified in a `return` command. If the procedure doesn't execute an explicit `return`, then its return value is the value of the last command executed in the procedure's body. If an error occurs while executing the procedure body, then the procedure as a whole will return that same error.

The syntax for the `return` command is:

```
return ?-code code? ?value?
```

The optional argument pair `-code code` allows to change the return status code from the default of `TH1_OK` to another status code. This code has to be specified with its numeric value.

9.6.7 Special commands

TH1 includes three core commands that assist with working with commands. They are:

```
breakpoint args
```

```
rename oldcmd newcmd
```

```
uplevel ?level? script
```

The `breakpoint` command does nothing. It is used as placeholder to place breakpoints during debugging.

The `rename` command renames a user defined or a built-in command. The old name is removed and the new name is inserted in the interpreter's command table.

The `uplevel` command executes a command in the variable scope of a command higher up in the call chain. The `script` argument is evaluated in the variable scope indicated by `level`. The `uplevel` command returns the result of that evaluation. If `level` is an integer, then it gives a distance (up the procedure calling stack) to move before executing the command. If `level` is omitted then it defaults to '1'.

For example, suppose that procedure `a` was invoked from top-level, and that it called `b`, and that `b` called `c`. Suppose that `c` invokes the `uplevel` command. If `level` is `1` or omitted, then the command will be executed in the variable context

of `b`. If level is 2 then the command will be executed in the variable context of `a`. If level is 3 then the command will be executed at top-level (i.e. only global variables will be visible).

The `uplevel` command causes the invoking procedure to disappear from the procedure calling stack while the command is being executed. In the above example, suppose `c` invokes the command

```
uplevel 1 {set x 43; d}
```

where `d` is another TH1 procedure. The `set` command will modify the variable `x` in the context of `b`, and `d` will execute at level 3, as if called from `b`. If it in turn executes the command

```
uplevel {set x 42}
```

then the `set` command will modify the same variable `x` in the context of `b` context: the procedure `c` does not appear to be on the call stack when `d` is executing.

9.7 Working with strings

TH1 provides the `string` command to facilitate working with strings. The `string` command is a single command with seven subcommands, identified by the first argument. The first argument serves no purpose other than to identify the subcommand. If the first argument does not match a subcommand, an error is thrown.

The seven `string` subcommands are:

```
string length string
```

```
string compare string1 string2
```

```
string first needle haystack ?startindex?
```

```
string last needle haystack ?startindex?
```

```
string range string first last
```

```
string repeat string count
```

```
string is alnum string
```

The `string length` subcommand takes one parameter, which is a string. It returns the decimal string with the length of the string. As TH1 uses a single byte character encoding the string size is both the size in characters and in bytes.

The `string compare` subcommand performs a character-by-character comparison of argument strings `string1` and `string2` in the same way as the C `strcmp`

procedure. It returns a decimal string with value -1, 0, or 1, depending on whether `string1` is lexicographically less than, equal to, or greater than `string2`.

The `string first` subcommand searches argument `haystack` for a sequence of characters that exactly match the characters in argument `needle`. If found, it returns a decimal string with the index of the first character in the first such match within `haystack`. If not found, it returns return -1. The optional integer argument `startindex` specifies the position where the search begins; the default value is 0, i.e. the first character in `haystack`.

The `string last` subcommand searches argument `haystack` for a sequence of characters that exactly match the characters in argument `needle`. If found, it returns a decimal string with the index of the first character in the last such match within `haystack`. If not found, it returns return -1. The optional integer argument `startindex` specifies the position where the search begins; the default value is 0, i.e. the first character in `haystack`.

The `string range` subcommand returns a range of consecutive characters from argument `string`, starting with the character whose index is `first` and ending with the character whose index is `last`. An index of zero refers to the first character of the string. `last` may be `end` to refer to the last character of the string. If `first` is less than zero then it is treated as if it was zero, and if `last` is greater than or equal to the length of the string then it is treated as if it were `end`. If `first` is greater than `last` then an empty string is returned.

The `string repeat` subcommand returns a string that is formed by repeating the argument string for `count` times. The argument count must be an integer. If count is zero or less the empty string is returned.

The `string is alnum` subcommand tests whether the argument string is an alphanumeric string, i.e. a string with only alphanumeric characters. It returns a decimal string with value 1 if the string is alphanumeric, and with value 0 if it is not.

9.8 Working with lists

The list is the basic TH1 data structure. A list is simply an ordered collection of items, numbers, words, strings, or other lists. For instance, the following string is a list with four items. The third item is a sub-list with two items:

```
{first second {a b} fourth}
```

TH1 has three core commands to work with lists:

```
list ?arg1 ?arg2? ...?
```

```
lindex list index
```

```
llength list
```


The `list` command returns a list comprising all the args. Braces and backslashes get added as necessary, so that the `lindex` command may be used on the result to re-extract the original arguments. For example, the command

```
list a b {c d e} {f {g h}}
```

will return

```
a b {c d e} {f {g h}}
```

The `lindex` command treats argument list as a TH1 list and returns the element with index number `index` from it. The argument `index` must be an integer number and zero refers to the first element of the list. In extracting the element, the `lindex` command observes the same rules concerning braces and quotes and backslashes as the TH1 command interpreter; however, variable substitution and command substitution do not occur. If `index` is negative or greater than or equal to the number of elements in value, an empty string is returned.

The `llength` command treats argument list as a list and returns a decimal string giving the number of elements in it.

Chapter 10

Chiselapp

Chiselapp is a website that is like github but hosts Fossil repositories. This way you can have your repository on a internet accessible host. This works like the Apache hosted repositories described in [par:Server-hosted] but Chiselapp supplies the server and the host is on the internet not a local area network.

After you set up a FREE account you can then push your repository to them and zap you are on the internet at:

`https://chiselapp.com/user/<your account>/repository/<Project>`

10.1 Create an account

Your first step is to create an account. The Chiselapp home page is:

Fill out the form with your information in my case I used my name and my Gmail account to set it up and my account is jschimpf.

10.2 Repositories

You can create repositories on the site and then copy one of your local repositories there. You have the choice of making public or private repositories. Public are visible to anyone visiting the site and private are visible only to you. In addition you do the standard Fossil assignment of users and privileges so once someone accesses the repository they only can do what you allow.(
Figure:[fig:User-Configuration])

The rest of this section will show how I am putting the repository NULLMODEM <http://chiselapp.com/user/jschimpf/repository/NULMODEM/index> on to ChiselApp.

10.2.1 Create Repository

The first step is to pick the option Create New Repository on your login page. This will give you the following screen:

So I fill in the name as NULLMODEM and I put in my repository password but what is Project Code ? Here you have to run Fossil to extract this information from your repository as follows:

```
500 FOSSIL> fossil info -R NULMODEM.fossil
```

The form is now filled in and we can create the repository and you get this:

10.2.2 Moving data

The next step is moving the repository on my disk to Chiselapp. This is done via a push command in Fossil. I am doing this command in the directory where NULMODEM.fossil lives so I don't need to type a path. Note the command is complete but I'm hiding my password when you do this type you password in full where I have .

```
501 FOSSIL> fossil push https://jschimpf:<passwd>@chiselapp.com/user/jschimpf
```

10.3 Fixing Data

When you go to your new repository things are a bit messed up. You get:

Whoa where's all my nice formatting and pointers to my documentation ? They are hidden and you have to get them back:

Go to the timeline view:

And see the top checkin that is the initial empty check-in this is an artifact of how Chiselapp creates your repository and you have to SHUN it

You will then be taken to another page where it will ask you if you really want to do this and pick Shun again.

Not quite there yet, you have to log into the project (Remember your name and password from Figure:[\[fig:Filled-in-form\]](#)) log in with this information and go to the Admin->Configuration page. Put in the same information you had on your local repository and ZAP your home page is back.

10.4 Final Fixes

The home page is now restored and we are ready to go.

The only problem now is the System Manual link doesn't work. The original was:

The fix is to change the link to just `/doc/tip/DOCS/NULMODEM.pdf`

10.5 Syncing

When you created the repository on ChiselApp you used this command:

```
fossil push https://jschimpf:<passwd>@chiselapp.com/user/jschimpf/repository/NULMODEM
```

when you did that the repository was created but it did not sync with your local one. This is probably not a good idea as you want the ChiselApp repository to stay up to date. If you leave off the `-once` then it will sync locally. If it isn't syncing now and you want to reverse this at any time just type:

```
fossil push https://jschimpf:<passwd>@chiselapp.com/user/jschimpf/repository/NULMODEM
```

i.e. the command without the `-once` and you are syncing again.

10.6 Final Result

Now you can go to <https://chiselapp.com/user/jschimpf/repository/NULMODEM> and view the repository and do what an anonymous user can do.

Chapter 11

Advanced uses

Fossil can be used in various ways. It is flexible and can be adapted and customized. Here are some examples for advanced usage.

11.1 Additional tables in the repository

It is possible to add more tables to the sqlite database that Fossil uses to store the repository into. If you add tables having names starting with `fx_` they will not be touched and also not deleted during a rebuild of the database.

However, these tables won't be synched and not be copied when cloning the database. They are only meant for local data. Still, you could add those tables in a server database and use the data to serve some specific content of the repository to all users accessing the repository over the web frontend.

Chapter 12

What's next ?

12.1 Learning more

This book so far has covered how to use the many features of Fossil and has, I hope, interested you in using it. The question “what’s next” now comes up. First go to the Fossil website. While there you can go to the Docs link and view the list pages. There are all sorts of topics covered in depth. If that still doesn’t help, you can join the Fossil forum and ask a question. I have found the forum contributors, including members of the development team, to be very helpful and have had my questions asked very quickly.

In the Fossil forum you will see suggestions for changes to be made to Fossil, some of these are accepted very quickly and will appear within hours in the Fossil source code. Others engender long discussions and it is interesting to read the pros and cons of suggested changes.

Fossil is an evolving program but if you get a version that has all the features you need you can stick with that version as long as you like. Going to a new version though is simple and just requires a rebuild of your current repositories. The developers have been very careful to preserve the basic structure so it is easy and safe to switch versions.

12.2 Contributing

Finally if you wish to contribute to the project there are many things to do, often triggered by forum comments.

Chapter 13

Revision history

Use the following table to track a history of this document's revisions. An entry should be made into this table for each version of the document.

Version	Author	Description	Issue Date
0.1	js	Initial Version	24-Apr-2010
0.2	js	Finishing up Single User Chapter	27-Apr-2010
0.3	js	Working on Introduction Chapter	30-Apr-2010
0.4	js	Adding multiuser chapter	1-May-2010
0.5	mn	Adding editorial corrections [ebf40b842a]	4-May-2010
0.6	js	Adding Command sections [e11399d575]	8-May-2010
0.7	js	English & spelling corrections	19-May-2010
0.8	js	Spelling fixes	30-May-2010
0.9	ws	Using Fossil merge [db6c734300]	2-Jun-2010
1.0	js/ws	Put Fossil merge first in handling fork	3-Jun-2010
1.1	mn	Fixes in multiple user chapter [e8770d3172]	4-Jun-2010
1.2	js	Start advanced use chapter [2abc23dae5]	4-Jun-2010
1.3	mn	English corrections Chapter 1 [8b324dc900]	5-Jun-2010
1.4	mn	Sections 2.1 & 2.2 corrections [0b34cb6f04]	7-Jun-2010
1.5	js	Move close leaf to adv use [2abc23dae5]	7-Jun-2010
1.6	js	Convert Advanced chapter to forks and branching	13-Jun-2010
1.7	js/tr	Add note about IP port usage [a62efa8eba]	8-Jul-2010
1.71	javelin	Check on misspelling section 1.1 [637d974f62]	15-Sep-2011
1.72	anon	Fix absolute path in image regs [d54868853b]	15-Sep-2011
1.73	anon	Fix fossil create section 2.2.5 [36772d90a5]	15-Sep-2011
1.74	anon	Push/Pull described incorrectly [1b930fced6]	15-Sep-2011
1.75	arnel	Commands might be changed [4aaf1f78bb]	15-Sep-2011
2.0	FvD	Updated and matched to fossil 1.25	March 2013
3.0	ctp	Migrated from Lyx to Markdown	Nov-2020

Version	Author	Description	Issue Date
3.x	tb	Aim to cover fossil 2.18 ... still much to do	Feb-2022