# BashGuide

August 21, 2025



## **Contents**

1	Intr	roduction	7
	1.1	About This Guide	7
	1.2	A Definition	7
	1.3	Using Bash	8
	1.4	Contents	9
2	Con	nmands and Arguments	11
	2.1	Strings	13
	2.2	Types of commands	14
		2.2.1 Aliases	14
		2.2.2 Functions	14
		2.2.3 Builtins	15
		2.2.4 Keywords	15
		2.2.5 Executables	15
	2.3	Scripts	16
3	Spec	cial Characters	19
	3.1	Deprecated special characters (recognized, but not recommended)	21
4	Para	ameters	23
	4.1	Special Parameters and Variables	24
	4.2	Variable Types	27
	4.3	Parameter Expansion	28
5	Patt	terns	33
	5.1	Glob Patterns	33
	5.2	Extended Globs	35
	5.3	Regular Expressions	35
	5.4	Brace Expansion	36
6	Test	sAndConditionals	39
	6.1	Exit Status	39
	6.2	Control Operators (&& and   )	40
	6.3	Grouping Statements	41
	6.4	Conditional Blocks (if, test and [[])	42
	6.5	Conditional Loops (while, until and for)	47
	6.6		51

7	Arra	ays	55
	7.1	Creating Arrays	56
	7.2	Using Arrays	58
		7.2.1 Expanding Elements	58
		7.2.2 Expanding Indices	61
		7.2.3 Sparse Arrays	62
	7.3	Associative Arrays	62
	, ,,		-
8	Inpu	at and Output	<b>65</b>
	8.1	Command-line Arguments	65
	8.2	The Environment	66
	8.3	File Descriptors	67
	8.4	Redirection	69
		8.4.1 File Redirection	69
		8.4.2 File Descriptor Manipulation	72
		8.4.3 Heredocs And Herestrings	74
	8.5	Pipes	76
	8.6	Miscellaneous Operators	78
	0.0	8.6.1 Process Substitution	78
		0.0.1 Trocess substitution	70
9	Com	npound Commands	81
	9.1	Subshells	81
	9.2	Command grouping	81
	9.3	Arithmetic Evaluation	83
	9.4	Functions	84
	9.5	Aliases	86
	9.6	Destroying Constructs	87
		,	
10	Sour	rcing	89
11	Job	Control	91
	-	Theory	91
		Practice	91
		Job Specifications	92
		See Also	92
	11,7	dec ruso	72
12	Prac	ctices	93
	12.1	Choose Your Shell	93
		Quoting	94
		Readability	97
		Bash Tests	98

12.5	Don't I	Ever Do These	)1
12.6	Debugg	ing	)3
	12.6.1	Diagnose the Problem	3
	12.6.2	Minimize the Codebase	)3
	12.6.3	Activate Bash's Debug Mode	)4
	12.6.4	Step Your Code	)5
	12.6.5	The Bash Debugger	)5
	12.6.6	Reread the Manual	)6
	12.6.7	Read the FAQ / Pitfalls	)6
	12.6.8	Ask Us on IRC	Э6

## 1 Introduction

You are invited to make additions or modifications at http://mywiki.wooledge.org/BashGuide so long as you can keep them accurate. Please test any code samples you write.

All the information here is presented without any warranty or guarantee of accuracy. Use it at your own risk. When in doubt, please consult the man pages or the GNU info pages as the authoritative references.

#### 1.1 About This Guide

A new version of this guide<sup>1</sup> is currently being drafted. For now, this guide is still the most complete and best reviewed. Any contributions to the new guide are welcome via GitHub forks<sup>2</sup>.

This guide aims to aid people interested in learning to work with BASH<sup>3</sup>. It aspires to teach good practice techniques for using BASH, and writing simple scripts.

This guide is targeted at beginning users. It assumes no advanced knowledge -- just the ability to login to a Unix-like system and open a command-line (terminal) interface. It will help if you know how to use a text editor; we will not be covering editors, nor do we endorse any particular editor choice. Familiarity with the fundamental Unix tool set, or with other programming languages or programming concepts, is not required, but those who have such knowledge may understand some of the examples more quickly.

If something is unclear to you, you are invited to report this (use BashGuideFeedback<sup>4</sup>, or the #bash channel on irc.libera.chat<sup>5</sup>) so that it may be clarified in this document for future readers.

You are invited to contribute to the development of this document by extending it or correcting invalid or incomplete information.

The primary maintainer(s) of this document:

- -- Lhunath<sup>6</sup> (primary author)
- -- GreyCat<sup>7</sup>

### 1.2 A Definition

BASH is an acronym for Bourne Again Shell. It is based on the Bourne shell and is mostly compatible with its features.

Shells are command interpreters. They are applications that provide users with the ability to give commands to their operating system interactively, or to execute batches of commands quickly. In no way are they required for the execution of programs; they are merely a layer between system function calls and the user.

http://quide.bash.academy/

<sup>2</sup>https://github.com/lhunath/bash.academy

<sup>3</sup>http://mywiki.wooledge.org/BASH

<sup>4</sup>http://mywiki.wooledge.org/BashGuideFeedback

<sup>5</sup>https://libera.chat/

<sup>6</sup>http://mywiki.wooledge.org/Lhunath

<sup>7</sup>http://mywiki.wooledge.org/GreyCat

Think of a shell as a way for you to speak to your system. Your system doesn't need it for most of its work, but it is an excellent interface between you and what your system can offer. It allows you to perform basic math, run basic tests and execute applications. More importantly, it allows you to combine these operations and connect applications to each other to perform complex and automated tasks.

BASH is **not** your operating system. It is not your window manager. It is not your terminal (but it oftens runs *inside* your terminal). It does not control your mouse or keyboard. It does not configure your system, activate your screensaver, or open your files when you double-click them. It is generally not involved in launching applications from your window manager or desktop environment. It's important to understand that BASH is only an interface for you to execute statements (using BASH syntax), either at the interactive BASH prompt or via BASH scripts.

- In The Manual: Introduction<sup>1</sup>
- *Shell*: A (possibly interactive) command interpreter, acting as a layer between the user and the system. *Bash*: The Bourne Again Shell, a *Bourne* compatible shell.

## 1.3 Using Bash

Most users that think of BASH think of it as a prompt and a command line. That is BASH in *interactive mode*. BASH can also run in *non-interactive mode*, as when executing scripts. We can use scripts to automate certain logic. Scripts are basically lists of commands (just like the ones you can type on the command line), but stored in a file. When a script is executed, all these commands are (generally) executed sequentially, one after another.

We'll start with the basics in an *interactive shell*. Once you're familiar with those, you can put them together in scripts. Important! You should make yourself familiar with the man and apropos commands on the shell. They will be vital to your self-tutoring.

```
$ man man
$ man apropos
```

In this guide, the \$ at the beginning of a line represents your BASH prompt. Traditionally, a shell prompt either ends with \$, \$ or #. If it ends with \$, this indicates a shell that's compatible with the Bourne shell (such as a POSIX shell, or a Korn shell, or Bash). If it ends with \$, this indicates a *C shell* (csh or tcsh); this guide does not cover *C* shell. If it ends with #, this indicates that the shell is running as the system's superuser account (root), and that you should be extra careful.

Your actual BASH prompt will probably be much longer than \$. Prompts are often highly individualized.

The man command stands for "manual"; it opens documentation (so-called "man pages") on various topics. You use it by running the command man [topic] at the BASH prompt, where [topic] is the name of the "page" you wish to read. Note that many of these "pages" are considerably longer than one printed page; nevertheless, the name persists. Each command (application) on your system is likely to have a man page. There are pages for other things too, such as system calls or specific configuration files. In this guide, we will only be covering commands.

Note that if you're looking for information on BASH built-ins (commands provided by BASH, not by external applications) you should look in man bash instead. BASH's manual is extensive and detailed. It is an excellent reference, albeit more technical than this guide.

http://www.gnu.org/software/bash/manual/bashref.html#Introduction

Bash also offers a help command which contains brief summaries of its built-in commands (which we'll discuss in depth later on).

```
$ help
$ help read
```

- In the FAQ: Is there a list of which features were added to specific releases (versions) of Bash?<sup>1</sup>
- Interactive mode: A mode of operation where a prompt asks you for one command at a time.
- Script: A file that contains a sequence of commands to execute one after the other.

#### 1.4 Contents

The guide has been divided into sections, which are intended to be read roughly in the order presented. If you skip ahead to a specific section, you might find yourself missing some background information from previous sections. (Links to relevant sections are not always provided when a topic is mentioned.)

- Commands and Arguments Types of commands; argument splitting; writing scripts.
- Special Characters Characters special to bash
- Parameters Variables; special parameters; parameter types; parameter expansion.
- Patterns Globs; filename matching; extended globs; brace expansion; regular expressions.
- Tests and Conditionals Exit status; && and | |; if, test and [[; while, until and for; case and select.
- Arrays Arrays; associative arrays.
- Input and Output Redirection; here documents; here strings; pipes; process substitution.
- Compound Commands Subshells; command grouping; arithmetic evaluation; functions; aliases.
- Sourcing Reading commands from other files.
- Job Control Working with background jobs.
- Practices Choosing your shell; quoting; readability; debugging.

<sup>1</sup>http://mywiki.wooledge.org/BashFAQ/061

## 2 Commands and Arguments

BASH<sup>1</sup> reads commands from its input (which is usually either a terminal or a file). Each line of input that it reads is treated as a *command* — an instruction to be carried out. (There are a few advanced cases, such as commands that span multiple lines, that will be gotten to later.)

Bash divides each line into *words* that are demarcated by a whitespace character (spaces and tabs). The first word of the line is the name of the command to be executed. All the remaining words become *arguments* to that command (options, filenames, etc.).

Assume we're in an empty directory... (to try these commands, create an empty directory called test and enter that directory by running: mkdir test; cd test):

```
$ ls  # List files in the current directory (no output, no files).
$ touch a b c  # Create files 'a', 'b', and 'c'.
$ ls  # List files again, and this time outputs: 'a', 'b', and 'c'.
a b c
```

The command ls prints out the names of the files in the current directory. The first time we run the list command we get no output, because there are no files yet.

The # character at the start of a word indicates a *comment*. Any words following the comment are ignored by the shell, meant only for reading. If we run these examples in our own shell, we don't have to type the comments; but even if we do, the command will still work.

touch is an application that changes the *Last Modified* time of a file. If the filename that it is given does not exist yet, it creates a file of that name as a new and empty file. In this example, we passed three arguments. touch creates a file for each argument. 1s shows us that three files have been created.

rm is an application that removes all the files that it was given.  $\star$  is a glob<sup>2</sup>. It basically means *all* and in this case means all files in the current directory. We will talk more about globs later.

Now, did we notice that there are several spaces between a and b, and only one between b and c? Also, notice that the files that were created by touch are no different than the first time. The amount of whitespace between arguments does not matter! This is important to know. For example:

```
$ echo This is a test.
This is a test.
$ echo This is a test.
This is a test.
```

http://mywiki.wooledge.org/BASH

<sup>2</sup>http://mywiki.wooledge.org/glob

echo is a command that prints its arguments to *standard output* (which in our case is the terminal). In this example, we provide the echo command with four arguments: 'This', 'is', 'a', and 'test.'. echo takes these arguments, and prints them out one by one with a space in between. In the second case, the exact same thing happens. The extra spaces make no difference. If we *want* the extra whitespace, we need to pass the sentence as one single argument. We can do this by using quotes<sup>1</sup>:

```
$ echo "This is a test."
This is a test.
```

Quotes group everything inside them into a single argument. The argument is: 'This is a test.'... specifically spaced. Note that the quotes are not part of the argument — Bash removes them before handing the argument to echo. echo prints this single argument out just like it always does.

Be very careful to avoid the following:

We need to make sure we quote filenames properly. If we don't, we'll end up deleting the wrong things! rm takes filenames as arguments. If our filenames have spaces and we do not quote them, Bash thinks each word is a separate argument. Bash hands each argument to rm separately. Like individually wrapped slices of processed cheese, rm treats each argument as a separate file.

In the above example, rm tried to delete a file for each word in the filename of the song, rather than keeping the filename intact. That caused our file secret to be deleted, and our song to remain behind!

This is what we should have done:

```
$ rm "The secret voice in your head.mp3"
```

Arguments are separated from the command name and from each other by a whitespace. This is important to remember. For example, the following is **wrong**:

```
$ [-f file]
bash: [-f: command not found
```

This is intended to test for the existence of a file named "file". It's reasonable to assume that whitespace around [ and ] doesn't matter, but [ is actually a command, and it requires its last argument to be ]. (We will cover the [ command in more detail later.) Therefore, we must separate [ from -f and ] from file, otherwise Bash will think we are trying to execute a command named [-f with a single argument file]. The correct command separates all arguments with whitespace:

<sup>1</sup>http://mywiki.wooledge.org/Quotes

```
$ [ -f file ]
```

(We see a lot of people who are confused by this behavior; they think that they can omit the whitespace between it and its arguments, so we need to present this particular example early.)

And, if our *filename* contains whitespace or other special characters, it should also be quoted:

```
$ [ -f "my file" ]
```

Have a good look at Arguments<sup>1</sup>, Quotes<sup>2</sup>, and WordSplitting<sup>3</sup> if all this isn't very clear yet. It is important to have a good grasp of how the shell interprets the whitespace and special characters before continuing with this guide.

#### Review:

- Arguments: These are additional words specified after the command ('1s -1 foo' executes 1s with two arguments).
- Quotes: The two forms of quotes, single and double (' and "), are used to group words and can protect special characters. The difference between ' and " will be discussed later.

#### Additionally:

- Tip *Always* quote sentences or strings that belong together, even if it's not absolutely necessary. This developed practice will reduce the risk of human error in the scripts. (e.g. quoting a sentence of an echo command).
- FAQ I'm trying to put a command in a variable, but the complex cases always fail! BashFAQ/050<sup>4</sup>.
- FAQ How can I handle command line arguments (options) to my script easily? BashFAQ/035<sup>5</sup>.

## 2.1 Strings

The term *string* refers to a sequence of characters which is treated as a single unit. The term is used loosely throughout this guide, as well as in almost every other programming language.

In Bash programming, almost *everything* is a string. When we type a command, the command's name is a string and each argument is a string; variable names are strings, and the contents of variables are strings as well; a filename is a string, and most files *contain* strings. They're everywhere!

An entire command can also be considered a string. This is not normally a useful point of view, but it illustrates the fact that *parts* of strings can sometimes be considered strings in their own right. A string which is part of a larger string is called a *substring*.

Strings do not have any intrinsic meaning. Their meaning is defined by how and where they are used.

Let's try another example. With the editor, write a shopping list and save it with the filename "list", and use cat to print it:

http://mywiki.wooledge.org/Arguments

<sup>2</sup>http://mywiki.wooledge.org/Quotes

<sup>&</sup>lt;sup>3</sup>http://mywiki.wooledge.org/WordSplitting

<sup>4</sup>http://mywiki.wooledge.org/BashFAQ/050

<sup>5</sup>http://mywiki.wooledge.org/BashFAQ/035

```
$ cat list
shampoo
tissues
milk (skim, not whole)
```

We typed a command: cat list. The shell reads this command as a string, and then divides it into the substrings cat and list. As far as the shell is concerned, list has no meaning, it's just a string with four characters in it. cat receives the argument list, which is a string of a filename. The string list has become meaningful because of *how it was used*.

The file happens to contain some text, which we see on our terminal. The entire file content — taken as a whole — is a string, but that string is not meaningful. However, if we divide the file into *lines* (and therefore treat each line as a separate string), then we see each individual *line* has meaning.

We can divide the final line into words, but these words are not meaningful by themselves. We can't buy "(skim" at the store, and we might get the wrong kind of "milk". Dividing the lines into words is not a useful thing to do in this example. But the shell doesn't know any of this — only we do!

So, when we are dealing with commands, data, and variables — all of which are just strings to the shell — we have *all* the responsibility. We need to be sure everything that needs to be separated is separated properly, and everything that needs to stay together stays together properly. We'll touch on these concepts repeatedly as we continue.

## 2.2 Types of commands

Bash understands several different types of commands: aliases, functions, builtins, keywords, and executables.

#### 2.2.1 Aliases

An alias is a way of shortening a command. (They are only used in **interactive** shells and *not* in scripts — this is one of the *very* few differences between a script and an interactive shell.) An alias is a *word* that is mapped to a certain *string*. Whenever that *word* is used as a command name, it is replaced by the *string* before executing the command. So, instead of executing:

```
$ nmap -Pn -A --osscan-limit 192.168.0.1
```

We could use an alias like this:

```
$ alias nmapp="nmap -Pn -A --osscan-limit"
$ nmapp 192.168.0.1
```

Aliases are limited in power; the replacement only happens in the first word. To have more flexibility, use a function. Aliases are only useful as simple textual shortcuts.

#### 2.2.2 Functions

Functions in Bash are somewhat like aliases, but more powerful. Unlike aliases, they can be used in **scripts**. A function contains shell commands, and acts very much like a small script; they can even take arguments and create local variables. When a function is called, the commands in it are executed. Functions will be covered in depth later in the guide<sup>1</sup>.

<sup>1</sup>http://mywiki.wooledge.org/BashGuide/CompoundCommands#Functions

#### 2.2.3 Builtins

Bash has some basic commands built into it, such as: cd (change directory), echo (write output), and so on. They can be thought of as functions that are already provided.

#### 2.2.4 Keywords

Keywords are like builtins, with the main difference being that keywords are actually Bash syntax and may be parsed using special rules. For example, [ is a Bash builtin, while [ [ is a Bash keyword; they are both used to test for a variety of conditions. Here we try to use them to compare the words "a" and "b" lexicographically:

```
$ [ a < b ]
-bash: b: No such file or directory
$ [[ a < b ]]</pre>
```

The first example returns an error because, as usual, Bash treats < as a file redirection operator<sup>1</sup> and attempts to redirect the nonexistent file b to the command [ a ]. The second example works because Bash parses words between [ [ and ] ] using different rules that don't use < for redirection.

#### 2.2.5 Executables

The last kind of command that can be executed by Bash is an *executable*. (Executables may also be called *external commands* or *applications*.) Executables are commonly invoked by typing only their name. This can be done because a pre-defined variable makes known to Bash a list of common, executable, file paths. This variable is called PATH. It is a set of directory names separated by colons (e.g./usr/bin:/bin). When a command is specified (e.g. myprogram, or ls) without a file path (and it isn't an alias, function, builtin or keyword), Bash searches through the directories in PATH. The search is done in order, from left to right, to see whether they contain an executable of the command typed.

If the executable is *outside* a known path... the executables file path will need to be defined. For an executable in the current directory, use ./myprogram; if it's in the directory /opt/somedirectory, use /opt/somedirectory/myprogram.

#### **Review:**

- *Alias* A word that is mapped to a string. Whenever that word is used as a command, it is replaced by the string it has mapped.
- Function A name that is mapped to a set of commands. Whenever the function is used as a command, it is called with the arguments following it. Functions are the basic method of making new commands.
- Builtin A command that has been built into Bash. Builtins are handled directly by the Bash executable and do not create new processes.
- Keyword A command that is actually part of Bash's syntax. Bash may parse keywords differently from regular commands.

<sup>1</sup>http://mywiki.wooledge.org/BashGuide/InputAndOutput#File\_Redirection

• Executable — A program that can be executed by referring to its file path (e.g. /bin/ls), or simply by its name if its location is in the PATH variable.

#### Additionally:

- Manual Simple Commands<sup>1</sup>
- FAQ What is the difference between test, [ and [ [? BashFAQ/031<sup>2</sup>.
- FAQ How can I make an alias that takes an argument? BashFAQ/080<sup>3</sup>.
- Tip The type command can be used to get a description of the command type:

```
$ type rm
rm is hashed (/bin/rm)
$ type cd
cd is a shell builtin
```

### 2.3 Scripts

A script is basically a sequence of commands in a file. Bash reads the file and processes the commands **in order**. It moves on to the next command only when the current one has **ended**. (The exception being if a command has been specified to run asynchronously, in the background. Don't worry too much about this case yet — we'll learn about how that works later on.)

Virtually any example that exists on the command line in this guide can also be used in a script.

Making a script is easy. Begin by making a new file, and put this on the first line:

```
#!/bin/bash
```

The header is called an *interpreter directive* (it is also called a *hashbang* or *shebang*). It specifies that /bin/bash is to be used as the interpreter when the file is used as the executable in a command. When the kernel executes a non-binary file, it reads the first line of the file. If the line begins with #!, the kernel uses the line to determine the interpreter to relay the file to. (There are other valid ways to do this as well, see below.) The #! must be at the very start of the file, with no spaces or blank lines before it. Our script's commands will appear on separate lines below this.

• Tip — Instead of #!/bin/bash, you could use

```
#!/usr/bin/env bash
```

env searches \$PATH for the executable named by its first argument (in this case, "bash"). For a more detailed explanation of this technique and how it differs from plain ol' #!/bin/bash, see this question in StackOverflow<sup>4</sup>.

http://www.gnu.org/software/bash/manual/bashref.html#Simple-Commands

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/031

 $<sup>^3</sup>$ http://mywiki.wooledge.org/BashFAQ/080

<sup>4</sup>https://stackoverflow.com/questions/16365130/what-is-the-difference-between-usr-bin-env-bash-and-usr-bin-bash/16365367#16365367

Please do not be fooled by scripts or examples on the Internet that use /bin/sh as the interpreter. **sh is not bash!** Bash itself is a "sh-compatible" shell (meaning that it can run most 'sh' scripts and carries much of the same syntax) however, the opposite is not true; some features of Bash will break or cause unexpected behavior in sh.

Also, please refrain from giving scripts a .sh extension. It serves no purpose, and it's completely misleading (since it's going to be a bash script, not an sh script).

It is perfectly fine to use **Windows** to write scripts. Avoid, however, using **Notepad**. "Microsoft Notepad" can only make files with DOS-style line-endings. DOS-style line-endings end with two characters: a Carriage Return (ASCII CR; 0xD) and a Line Feed (ASCII LF; 0xA) character. Bash understands line-endings with only Line Feed characters. As a result, the Carriage Return character will cause an unexpected surprise if one doesn't know it's there (very weird error messages). If at all possible, use a more capable editor like Vim<sup>1</sup>, Emacs<sup>2</sup>, kate, GEdit... If one doesn't, the carriage returns will need to be removed from the scripts before running them.

Once the script file has been created, it can be executed by doing:

```
$ bash myscript
```

In this example, we execute bash and tell it to read the script. When we do it like this, the #! line is just a comment, Bash does not do anything at all with it.

Alternatively, we can give our scripts executable permissions. With this method, instead of calling Bash manually, we can execute the script as an application:

```
$ chmod +x myscript # Mark the file as executable.
$ ./myscript # Now, myscript can be executed directly instead of having to pass it to bash.
```

When we run the command this way, the operating system (OS) uses the #! line to determine what interpreter to use.

To decide where to put the script, a couple alternatives exists. Generally, people like to keep their scripts in a personally-defined directory; this prevents your script from interfering with other users on the system. Some like to keep their scripts in a pre-existing directory in the PATH, because these people think that they will never make a mistake.

To use a personal directory:

```
$ mkdir -p "$HOME/bin"
$ echo 'PATH="$HOME/bin:$PATH"' >> "$HOME/.bashrc"
$ source "$HOME/.bashrc"
```

The first command will make a directory called bin inside your *home directory* (the directory that belongs to you, personally). It is traditional for directories that contain commands to be named bin, even when those commands are scripts and not compiled ("binary") programs. The second command adds a line containing a variable assignment to a file. The variable is PATH, and we are adding this line to the Bash configuration file (.bashrc). Every new *interactive* instance of Bash will now check for executable scripts in our new directory before checking any directories that were already in PATH. The third command tells Bash to re-read its configuration file.

Some people prefer to use a different directory to hold their personal scripts, such as \$HOME/.config/bin or \$HOME/.local/bin You can use whatever you prefer, as long as you are consistent.

http://www.vim.org/

<sup>2</sup>http://www.gnu.org/software/emacs/

Changes to the Bash configuration file will not have an immediate effect; we have to make the step to re-read the file. In the example above, we used source "\$HOME/.bashro". We could have also used exec bash, or we could close the existing terminal and open a new one. Bash would then initialize itself again by reading .bashro (and possibly other files).

In any case, we can now put our script in our bin directory and execute it as a normal command — we no longer need to prepend our script's name with the file path (which was the . / part in the previous example):

```
$ mv myscript "$HOME/bin"
$ myscript
```

#### **Additionally:**

- Tip If the system has multiple versions of bash installed, specifying the interpreter by absolute path may be desirable in order to ensure the right bash version is used. For example: #!/usr/bin/bash. Type "type -a bash" to print the paths to all Bash executables in PATH.
- Tip The interpreter can optionally be followed by one word of interpreter's options. For example, the following options will turn on verbose debugging: "#!/usr/bin/bash -x". To learn more, see Debugging<sup>1</sup>.
- Tip After defining the interpreter in the header, it is recommended to summarize the scripts purpose. If desired, copyright information can be listed...:

```
#!/usr/bin/env bash
# scriptname - a short explanation of the scripts purpose.
# Copyright (C) <date> <name>...
# # copyright (C) <date> in the script are script are script are script are scriptname [option] [argument] ...
```

 $<sup>^{1} \</sup>verb|http://mywiki.wooledge.org/BashGuide/Practices#Debugging|$ 

## 3 Special Characters

Some characters are evaluated by Bash<sup>1</sup> to have a *non-literal* meaning. Instead, these characters carry out a special instruction, or have an alternate meaning; they are called "special characters", or "meta-characters".

Here are some of the more common special characters uses:

Char.	Description
	Whitespace — this is a tab, newline, vertical tab, form
	feed, carriage return, or space. Bash uses whitespace to
пп	determine where words begin and end. The first word is
	the command name and additional words become
	arguments to that command.
	Expansion — introduces various types of expansion:
	parameter expansion (e.g. \$var or \${var}), command
\$	substitution <sup>2</sup> (e.g. \$ (command)), or arithmetic
	expansion (e.g. \$ ((expression))). More on
	expansions later.
	Single quotes — protect the text inside them so that it has
	a <i>literal</i> meaning. With them, generally any kind of
11	interpretation by Bash is ignored: special characters are
	passed over and multiple words are prevented from being
	split.
	Double quotes — protect the text inside them from being
" "	split into multiple words or arguments, yet allow
	substitutions to occur; the meaning of most other special
	characters is usually prevented.
	Escape — (backslash) prevents the next character from
	being interpreted as a special character. This works
\	outside of quoting, inside double quotes, and generally
	ignored in single quotes.
	Comment — the # character begins a commentary that
#	extends to the end of the line. Comments are notes of
	explanation and are not processed by the shell.
	Assignment assign a value to a variable (e.g.
=	logdir=/var/log/myprog). Whitespace is not
	allowed on either side of the = character.

<sup>1</sup>http://mywiki.wooledge.org/Bash

<sup>&</sup>lt;sup>2</sup>http://mywiki.wooledge.org/CommandSubstitution

	Test — an evaluation of a conditional expression to determine whether it is "true" or "false". Tests are used in
	Bash to compare strings, check the existence of a file, etc.  More of this will be covered later.
	Negate — used to negate or reverse a test or exit status.
!	For example: ! grep text file; exit \$?.
>, >>, <	Redirection — redirect a command's output or input to a
, ,	file. Redirections will be covered later.
	Pipe — send the output from one command to the input
	of another command. This is a method of chaining
	commands together. Example: echo "Hello
	beautiful."   grep -o beautiful.  Command separator — used to separate multiple
<b>;</b>	commands that are on the same line.
	Inline group — commands inside the curly braces are
	treated as if they were one command. It is convenient to
{ }	use these when Bash syntax requires only one command
	and a function doesn't feel warranted.
	Subshell group — similar to the above but where
	commands within are executed in a subshell <sup>1</sup> (a new
( )	process). Used much like a sandbox, if a command causes
	side effects (like changing variables), it will have no
	effect on the current shell.
	Arithmetic expression — with an arithmetic expression <sup>2</sup> ,
	characters such as +, -, *, and / are mathematical
(())	operators used for calculations. They can be used for
	variable assignments like (( a = 1 + 4 )) as well
	as tests like if (( a < b )). More on this later.
	Arithmetic expansion — Comparable to the above, but the
\$(())	expression is replaced with the result of its arithmetic
	evaluation. Example: echo "The average is \$((
	(a+b)/2 ))".
*,?	Globs <sup>3</sup> "wildcard" characters which match parts of
^,:	filenames (e.g. ls *.txt).

<sup>1</sup>http://mywiki.wooledge.org/SubShell
2http://mywiki.wooledge.org/ArithmeticExpression
3http://mywiki.wooledge.org/glob

~	Home directory — the tilde is a representation of a home directory. When alone or followed by a /, it means the current user's home directory; otherwise, a username must be specified (e.g. ls ~/Documents; cp ~john/.bashrc .).
&	Background when used at the end of a command, run the command in the background (do not wait for it to complete).

#### Examples:

```
$ echo "I am $LOGNAME"
I am lhunath
$ echo 'I am $LOGNAME'
I am $LOGNAME
$ # boo
$ echo An open \ \ space
An open space
$ echo "My computer is $(hostname)"
My computer is Lyndir
$ echo boo > file
$ echo $((5 + 5))
10
$ ((5 > 0)) && echo "Five is greater than zero."
Five is greater than zero.
```

## 3.1 Deprecated special characters (recognized, but not recommended)

This group of characters will also be evaluated by Bash<sup>1</sup> to have a *non-literal* meaning, but are generally included for backwards compatibility only. These are not recommended for use, but often appear in older or poorly written scripts.

Char.	Description
` `	Command substitution <sup>2</sup> - use $\$$ ( ) instead.
	Test - an alias for the old test command. Commonly
	used in POSIX shell scripts. Lacks many features of [ [
	]].
\$[]	Arithmetic expression <sup>3</sup> - use $\$$ ( ( ) ) instead.

## Additionally:

 $<sup>^{1} \</sup>verb|http://mywiki.wooledge.org/Bash|$ 

<sup>2</sup>http://mywiki.wooledge.org/CommandSubstitution

<sup>3</sup>http://mywiki.wooledge.org/ArithmeticExpression

	_			_			1
•	In	The	Manu	ıal —	Shell	<b>Syntax</b>	1

• Special Characters — Characters that have a special meaning to Bash. Usually their meaning is interpreted and then they are removed from the command before executing it.

<sup>1</sup> http://www.gnu.org/software/bash/manual/html\_node/Shell-Syntax.html#Shell-Syntax

## 4 Parameters

Parameters are a sort of named space in memory you can use to retrieve or store information. Generally speaking, they will store string data, but can also be used to store integers, indexed and associative arrays.

Parameters come in two flavors: *variables* and *special parameters*. Special parameters are read-only, pre-set by BASH, and used to communicate some type of internal status. Variables are parameters that you can create and update yourself. Variable names are bound by the following rule:

• *Name*: A word consisting only of letters, digits and underscores, and beginning with a letter or an underscore. Also referred to as an *identifier*.

To store data in a variable, we use the following assignment syntax:

```
$ varname=vardata
```

This command assigns the data vardata to the variable by name of varname.

Please note that you *cannot* use spaces around the = sign in an assignment. If you write this:

```
# This is wrong!
$ varname = vardata
```

BASH<sup>1</sup> will not know that you are attempting to assign something. The parser will see varname with no = and treat it as a command name, and then pass = and vardata to it as arguments.

To access the data stored in a variable, we use parameter expansion. Parameter expansion is the *substitution* of a parameter by its value, which is to say, the syntax tells bash that you want to use the contents of the variable. After that, Bash *may still perform additional manipulations on the result*. This is a very important concept to grasp correctly, because it is very much unlike the way variables are handled in other programming languages!

To illustrate what parameter expansion is, let's use this example:

```
$ foo=bar
$ echo "Foo is $foo"
```

When Bash is about to execute your code, it first changes the command by taking your parameter expansion (the \$foo), and replacing it by the contents of foo, which is bar. The command becomes:

```
$ echo "Foo is bar"
Foo is bar
```

Now, Bash is ready to execute the command. Executing it shows us the simple sentence on screen.

It is important to understand that parameter expansion causes the \$parameter to be **replaced** by its contents. Note the following case, which relies on an understanding of the previous chapter on argument splitting:

```
$ song="My song.mp3"
$ rm $song
rm: My: No such file or directory
rm: song.mp3: No such file or directory
```

<sup>1</sup>http://mywiki.wooledge.org/BASH

Why did this not work? Because Bash replaced your \$song by its contents, being My song.mp3; then it performed word splitting; and only THEN executed the command. It was as if you had typed this:

```
$ rm My song.mp3
```

And according to the rules of word splitting, Bash thought you meant for My and song.mp3 to mean two different files, because there is white space between them and it wasn't quoted. How do we fix this? We remember to **put double quotes around every parameter expansion!** 

• Parameters: Parameters store data that can be retrieved through a symbol or a name.

## 4.1 Special Parameters and Variables

Let's get our vocabulary straight before we get into the real deal. There are *Parameters* and *Variables*. Variables are actually just one kind of parameter: parameters that are denoted by a name. Parameters that aren't variables are called *Special Parameters*. I'm sure you'll understand things better with a few examples:

```
$ # Some parameters that aren't variables:
$ echo "My shell is $0, and has these options set: $-"
My shell is -bash, and has these options set: himB
$ # Some parameters that ARE variables:
$ echo "I am $LOGNAME, and I live at $HOME."
I am lhunath, and I live at /home/lhunath.
```

**Please note**: Unlike PHP/Perl/... parameters do NOT start with a \$-sign. The \$-sign you see in the examples merely causes the parameter that follows it to be *expanded*. Expansion basically means that the shell replaces the parameter by its content. As such, LOGNAME is the parameter (variable) that contains your username. \$LOGNAME is an expression that will be replaced with the content of that variable, which in my case is lhunath.

I think you've got the drift now. Here's a summary of most of the Special Parameters:

Parameter Name	Usage	Description
0	"\$0"	Contains the name, or the path, of the
V	~ \$0 ··	script. This is not always reliable.
		Positional Parameters contain the
<b>12</b> etc.	"\$1" etc.	arguments that were passed to the
		current script or function.
		Expands to all the words of all the
		positional parameters. Double
*	"\$*"	quoted, it expands to a single string
	J *	containing them all, separated by the
		first character of the <b>IFS</b> variable
		(discussed later).

@	<b>"</b> \$@"	Expands to all the words of all the positional parameters. Double quoted, it expands to a list of them all as individual words.
#	\$#	Expands to the number of positional parameters that are currently set.
?	\$?	Expands to the exit code of the most recently completed foreground command.
\$	\$\$	Expands to the PID <sup>1</sup> (process ID number) of the current shell.
!	\$!	Expands to the PID of the command most recently executed in the background.
-	"\$_"	Expands to the last argument of the last command that was executed.

And here are a few examples of Variables that the shell provides for you:

- BASH\_VERSION: Contains a string describing the version of Bash.
- **HOSTNAME**: Contains the hostname of your computer, I swear. Either short or long form, depending on how your computer is set up.
- **PPID**: Contains the PID of the parent process of this shell.
- PWD: Contains the current working directory.
- RANDOM: Each time you expand this variable, a (pseudo)random number between 0 and 32767 is generated.
- UID: The ID number of the current user. Not reliable for security/authentication purposes, alas.
- COLUMNS: The number of characters that fit on one line in your terminal. (The width of your terminal in characters.)
- LINES: The number of lines that fit in your terminal. (The height of your terminal in characters.)
- **HOME**: The current user's home directory.
- PATH: A colon-separated list of paths that will be searched to find a command, if it is not an alias, function, builtin command, or shell keyword, and no pathname is specified.
- PS1: Contains a string that describes the format of your shell prompt.
- TMPDIR: Contains the directory that is used to store temporary files (by the shell).

http://mywiki.wooledge.org/ProcessManagement

(There are many more -- see the manual for a comprehensive list.) Of course, you aren't restricted to only these variables. Feel free to define your own:

```
$ country=Canada
$ echo "I am $LOGNAME and I currently live in $country."
I am lhunath and I currently live in Canada.
```

Notice what we did to assign the value Canada to the variable country. Remember that you are **NOT** allowed to have any spaces before or after that equals sign!

```
$ language = PHP
-bash: language: command not found
$ language=PHP
$ echo "I'm far too used to $language."
I'm far too used to PHP.
```

Remember that Bash is not Perl or PHP. You need to be very well aware of how *expansion* works to avoid **big** trouble. If you don't, you'll end up creating very dangerous situations in your scripts, especially when making this mistake with rm:

```
$ ls
no secret secret
$ file='no secret'
$ rm $file
rm: cannot remove `no': No such file or directory
```

Imagine we have two files, no secret and secret. The first contains nothing useful, but the second contains the secret that will save the world from impending doom. Unthoughtful as you are, you forgot to **quote** your parameter expansion of file. Bash expands the parameter and the result is rm no secret. Bash splits the arguments up by their whitespace as it normally does, and rm is passed two arguments: 'no' and 'secret'. As a result, it fails to find the file no and it deletes the file secret. The secret is lost!

#### Good Practice:

You should always keep parameter expansions properly quoted. This prevents the whitespace or the possible globs inside of them from giving you gray hair or unexpectedly wiping stuff off your computer. The only good PE, is a quoted PE.

- In The Manual: Shell Parameters<sup>1</sup>, Shell Variables<sup>2</sup>
- In the FAQ:

How can I concatenate two variables? How do I append a string to a variable?<sup>3</sup>

How can I access positional parameters after \$9?4

http://www.gnu.org/software/bash/manual/bashref.html#Shell-Parameters

<sup>&</sup>lt;sup>2</sup>http://www.gnu.org/software/bash/manual/bashref.html#Shell-Variables

<sup>3</sup>http://mywiki.wooledge.org/BashFAQ/013

<sup>4</sup>http://mywiki.wooledge.org/BashFAQ/025

- *Variable*: A variable is a kind of parameter that you can create and modify directly. It is denoted by a name, which must begin with a letter or underscore (\_), and must consist only of letters, digits, and the underscore. Variable names are case-sensitive.
- *Expansion*: Expansion happens when a parameter is prefixed by a dollar sign. Bash takes the parameter's value and replaces the parameter's expansion by its value before executing the command. This is also called *substitution*.

## 4.2 Variable Types

Although Bash is not a typed language, it does have a few different types of variables. These types define the kind of content they are allowed to have. Type information is stored internally by Bash.

- Array: declare -a variable: The variable is an array of strings.
- Associative array: declare -A variable: The variable is an associative array of strings (bash 4.0 or higher).
- **Integer**: declare -i variable: The variable holds an integer. Assigning values to this variable automatically triggers *Arithmetic Evaluation*.
- **Read Only**: declare -r *variable*: The variable can no longer be modified or unset.
- Export: declare -x *variable*: The variable is marked for export which means it will be inherited by any child process.

Arrays are basically indexed lists of strings. They are very convenient for their ability to store multiple strings together without relying on a *delimiter* to split them apart (which is tedious when done correctly and error-prone when not).

Defining variables as integers has the advantage that you can leave out some syntax when trying to assign or modify them:

```
$ a=5; a+=2; echo "$a"; unset a
52
$ a=5; let a+=2; echo "$a"; unset a
7
$ declare -i a=5; a+=2; echo "$a"; unset a
7
$ a=5+2; echo "$a"; unset a
5+2
$ declare -i a=5+2; echo "$a"; unset a
7
```

However, in practice the use of declare —i is exceedingly rare. In large part, this is because it creates behavior that can be surprising to anyone trying to maintain the script, who misses the declare statement. Most experienced shell scripters prefer to use explicit arithmetic commands (with ((...)) or let) when they want to perform arithmetic.

It is also rare to see an explicit declaration of an array using declare -a. It is sufficient to write array=(...) and Bash will know that the variable is now an array. The exception to this is the associative array, which *must* be declared explicitly: declare -A myarray.

- String: A string is a sequence of characters.
- Array: An array is a list of strings indexed by numbers.
- *Integer*: An integer is a whole number (positive, negative or zero).
- Read Only: Parameters that are read-only cannot be modified or unset.
- Export: Variables that are marked for export will be inherited by any child process. Variables inherited in this way are called Environment Variables.
- In the FAQ:

How can I use array variables?<sup>1</sup>

## 4.3 Parameter Expansion

*Parameter Expansion* is the term that refers to any operation that causes a parameter to be expanded (replaced by content). In its most basic appearance, the expansion of a parameter is achieved by prefixing that parameter with a \$ sign. In certain situations, additional curly braces around the parameter's name are required:

```
$ echo "'$USER', '$USER's'"
'lhunath', '', 'lhunaths'
```

This example illustrates what basic parameter expansions (PE) look like. The second PE results in an empty string. That's because the parameter USERs is empty. We did not intend to have the s be part of the parameter name. Since there's no way Bash could know you want a literal s appended to the parameter's value, you need to use curly braces to mark the beginning and end of the parameter name. That's what we do in the third PE in our example above.

Parameter Expansion also gives us tricks to modify the string that will be expanded. These operations can be terribly convenient:

```
$ for file in *.JPG *.jpeg
do mv -- "$file" "${file%.*}.jpg"
done
```

The code above can be used to rename all JPEG files with a .JPG or a .jpeg extension to have a normal .jpg extension. The expression \$ {file%.\*} cuts off everything from the end starting with the last period (.). Then, in the same quotes, a new extension is appended to the expansion result.

Here's a summary of most of the PE tricks that are available:

<sup>1</sup>http://mywiki.wooledge.org/BashFAQ/005

Syntax	Description
	Use Default Value. If 'parameter' is unset or null,
<pre>\${parameter:-word}</pre>	'word' (which may be an expansion) is substituted.
	Otherwise, the value of 'parameter' is substituted.
	Assign Default Value. If 'parameter' is unset or null,
4.6	'word' (which may be an expansion) is assigned to
<pre>\${parameter:=word}</pre>	'parameter'. The value of 'parameter' is then
	substituted.
	Use Alternate Value. If 'parameter' is null or unset,
<pre>\${parameter:+word}</pre>	nothing is substituted, otherwise 'word' (which may be
	an expansion) is substituted.
	Substring Expansion. Expands to up to 'length'
	characters of 'parameter' starting at the character
	specified by 'offset' (0-indexed). If ':length' is
	omitted, go all the way to the end. If 'offset' is
	negative (use parentheses!), count backward from the end
<pre>\${parameter:offset:length}</pre>	of 'parameter' instead of forward from the beginning.
	If 'parameter' is @ or an indexed array name
	subscripted by @ or *, the result is 'length' positional
	parameters or members of the array, respectively, starting
	from 'offset'.
	The length in characters of the value of 'parameter' is
<pre>\${#parameter}</pre>	substituted. If 'parameter' is an array name
( "parameter)	subscripted by @ or *, return the number of elements.
	The 'pattern' is matched against the <b>beginning</b> of
	'parameter'. The result is the expanded value of
	'parameter' with the shortest match deleted.
<pre>\${parameter#pattern}</pre>	If 'parameter' is an array name subscripted by @ or *,
	this will be done on each element. Same for all following
	items.
\${parameter##pattern}	As above, but the <i>longest</i> match is deleted.
\{\parameter=\partarelarger}	The 'pattern' is matched against the <b>end</b> of
\${parameter%pattern}	'parameter'. The result is the expanded value of
3{parameterspactern;	'parameter' with the shortest match deleted.
¢(naramatar%%nattarn)	As above, but the <i>longest</i> match is deleted.
<pre>\${parameter%%pattern}</pre>	Results in the expanded value of 'parameter' with the
	<del>-</del>
<pre>\${parameter/pat/string}</pre>	first (unanchored) match of 'pat' replaced by
	'string'. Assume null string when the '/string' part
Classical Last Last Last Last Last Last Last Last	is absent.
<pre>\${parameter//pat/string}</pre>	As above, but <b>every</b> match of 'pat' is replaced.

\${parameter/#pat/string}	As above, but matched against the <b>beginning</b> . Useful for adding a common prefix with a null pattern:  "\${array[@]/#/prefix}".
\${parameter/%pat/string}	As above, but matched against the <b>end</b> . Useful for adding a common suffix with a null pattern.

You will learn them through experience. They come in handy far more often than you think they might. Here are a few examples to kickstart you:

```
$ file="$HOME/.secrets/007"; \
echo "File location: $file"; \
echo "Filename: ${file##*/}"; \
echo "Directory of file: ${file%/*}"; \
echo "Non-secret file: ${file/secrets/not_secret}"; \
echo; \
echo "Other file location: ${other:-There is no other file}"; \
echo "Using file if there is no other file: ${other:=$file}"; \
echo "Other filename: ${other##*/}"; \
echo "Other file location length: ${#other}"
File location: /home/lhunath/.secrets/007
Filename: 007
Directory of file: /home/lhunath/.secrets
Non-secret file: /home/lhunath/.not_secret/007
Other file location: There is no other file
Using file if there is no other file: /home/lhunath/.secrets/007
Other filename: 007
Other file location length: 26
```

Remember the difference between  $\{v \neq p\}$  and  $\{v \neq p\}$ . The doubling of the # character means patterns will become greedy. The same goes for %:

```
$ version=1.5.9; echo "MAJOR: ${version%*.*}, MINOR: ${version#*.}."
MAJOR: 1, MINOR: 5.9.
$ echo "Dash: ${version/./-}, Dashes: ${version//./-}."
Dash: 1-5.9, Dashes: 1-5-9.
```

Note: You **cannot** use multiple PEs together. If you need to execute multiple PEs on a parameter, you will need to use multiple statements:

```
$ file=$HOME/image.jpg; file=${file##*/}; echo "${file%.*}"
image
```

#### • Good Practice:

You may be tempted to use external applications such as sed, awk, cut, perl or others to modify your strings. Be aware that all of these require an extra process to be started, which in some cases can cause slowdowns. Parameter Expansions are the perfect alternative.

- In The Manual: Shell Parameter Expansion<sup>1</sup>
- In the FAQ:

How do I do string manipulations in bash?<sup>2</sup>

- How can I rename all my \*.foo files to \*.bar, or convert spaces to underscores, or convert upper-case file names to lower case?<sup>3</sup>
- How can I use parameter expansion? How can I get substrings? How can I get a file without its extension, or get just a file's extension?<sup>4</sup>
- How do I get the effects of those nifty Bash Parameter Expansions in older shells?<sup>5</sup>
- How do I determine whether a variable is already defined? Or a function?<sup>6</sup>
- *Parameter Expansion*: Any expansion (see earlier definition) of a parameter. Certain operations are possible during this expansion that are performed on the value that will be expanded.

 $<sup>^{1} \</sup>verb|http://www.gnu.org/software/bash/manual/bashref.html \#Shell-Parameter-Expansion$ 

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/100

<sup>3</sup>http://mywiki.wooledge.org/BashFAQ/030

<sup>4</sup>http://mywiki.wooledge.org/BashFAQ/073

<sup>5</sup>http://mywiki.wooledge.org/BashFAQ/074

<sup>6</sup>http://mywiki.wooledge.org/BashFAQ/083

## 5 Patterns

BASH<sup>1</sup> offers three different kinds of *pattern matching*. Pattern matching serves two roles in the shell: selecting filenames within a directory, or determining whether a string conforms to a desired format.

On the command line you will mostly use *globs*. These are a fairly straight-forward form of patterns that can easily be used to match a range of files, or to check variables against simple rules.

The second type of pattern matching involves *extended globs*, which allow more complicated expressions than regular globs.

Since version 3.0, Bash also supports *regular expression* patterns. These will be useful mainly in scripts to test user input or parse data. (You can't use a regular expression to select filenames; only globs and extended globs can do that.)

• Pattern: A pattern is a string with a special format designed to match filenames, or to check, classify or validate data strings.

#### 5.1 Glob Patterns

Globs<sup>2</sup> are a very important concept in Bash, if only for their incredible convenience. Properly understanding globs will benefit you in many ways. Globs are basically patterns that can be used to match filenames or other strings.

Globs are composed of normal characters and metacharacters. Metacharacters are characters that have a special meaning. These are the metacharacters that can be used in globs:

- \*: Matches any string, including the null string.
- ?: Matches any single character.
- [...]: Matches any one of the enclosed characters.

Globs are implicitly *anchored* at both ends. What this means is that a glob must match a *whole* string (filename or data string). A glob of a\* will not match the string cat, because it only matches the at, not the whole string. A glob of ca\*, however, would match cat.

Here's an example of how we can use glob patterns to expand to filenames:

```
$ 1s
a abc b c
$ echo *
a abc b c
$ echo a*
a abc
```

Bash sees the glob, for example a\*. It *expands* this glob, by looking in the current directory and matching it against all files there. Any filenames that match the glob are gathered up and sorted, and then the list of filenames is used in place of the glob. As a result, the statement echo a\* is replaced by the statement echo a abc, which is then executed.

<sup>1</sup>http://mywiki.wooledge.org/BASH

<sup>2</sup>http://mywiki.wooledge.org/glob

When a glob is used to match *filenames*, the  $\star$  and ? characters cannot match a slash (/) character. So, for instance, the glob  $\star$ /bin might match foo/bin but it cannot match /usr/local/bin. When globs match *patterns*, the / restriction is removed.

Bash performs filename expansions *after* word splitting has already been done. Therefore, filenames generated by a glob will not be split; they will always be handled correctly. For example:

```
$ touch "a b.txt"
$ ls
a b.txt
$ rm *
$ ls
```

Here, \* is expanded into the single filename "a b.txt". This filename will be passed as a single argument to rm. Using globs to enumerate files is **always** a better idea than using `ls` for that purpose. Here's an example with some more complex syntax which we will cover later on, but it will illustrate the reason very well:

```
$ ls
a b.txt
$ for file in `ls`; do rm "$file"; done
rm: cannot remove `a': No such file or directory
rm: cannot remove `b.txt': No such file or directory
$ for file in *; do rm "$file"; done
$ ls
```

Here we use the for command to go through the output of the ls command. The ls command prints the string a b.txt. The for command splits that string into words over which it iterates. As a result, for iterates over first a, and then b.txt. Naturally, this is **not** what we want. The glob, however, expands in the proper form. It results in the string "a b.txt", which for takes as a single argument.

In addition to filename expansion, globs may also be used to check whether data matches a specific format. For example, we might be given a filename, and need to take different actions depending on its extension:

```
$ filename="somefile.jpg"
$ if [[ $filename = *.jpg ]]; then
> echo "$filename is a jpeg"
> fi
somefile.jpg is a jpeg
```

The [ [ keyword and the case keyword (which we will discuss in more detail later) both offer the opportunity to check a string against a glob -- either regular globs, or extended globs, if the latter have been enabled.

- Good Practice: You should always use globs instead of ls (or similar) to enumerate files. Globs will always expand safely and minimize the risk for bugs. You can sometimes end up with some very weird filenames. Most scripts aren't tested against all the odd cases that they may end up being used with. Don't let your script be one of those!
- In The Manual: Pattern Matching<sup>1</sup>

Ihttp://www.gnu.org/software/bash/manual/bashref.html#Pattern-Matching

- In the FAQ: How can I use a logical AND/OR/NOT in a shell pattern (glob)?<sup>1</sup>
- Glob: A glob is a string that can match certain strings or filenames.

#### 5.2 Extended Globs

Bash also supports a feature called *Extended Globs*. These globs are more powerful in nature; technically, they are equivalent to regular expressions, although the syntax looks different than most people are used to. This feature is turned off by default, but can be turned on with the shopt command, which is used to toggle **she**ll **opt**ions:

```
$ shopt -s extglob
```

- ?(list): Matches zero or one occurrence of the given patterns.
- \*(list): Matches zero or more occurrences of the given patterns.
- +(list): Matches one or more occurrences of the given patterns.
- @(list): Matches one of the given patterns.
- !(list): Matches anything but the given patterns.

The list inside the parentheses is a list of globs or extended globs separated by the | character. Here's an example:

```
$ ls
names.txt tokyo.jpg california.bmp
$ echo !(*jpg|*bmp)
names.txt
```

Our extended glob expands to anything that does not match the \*jpg or the \*bmp pattern. Only the text file passes for that, so it is expanded.

### **5.3** Regular Expressions

Regular expressions (regex) are similar to *Glob Patterns*, but they can only be used for pattern matching, not for filename matching. Since 3.0, Bash supports the  $=\sim$  operator to the [[keyword. This operator matches the string that comes before it against the regex pattern that follows it. When the string matches the pattern, [[returns with an exit code of 0 ("true"). If the string does not match the pattern, an exit code of 1 ("false") is returned. In case the pattern's syntax is invalid, [[will abort the operation and return an exit code of 2.

Bash uses the *Extended Regular Expression* (ERE) dialect. We will not cover regexes in depth in this guide, but if you are interested in this concept, please read up on RegularExpression<sup>2</sup>, or Extended Regular Expressions<sup>3</sup>.

```
http://mywiki.wooledge.org/BashFAQ/016
```

<sup>2</sup>http://mywiki.wooledge.org/RegularExpression

 $<sup>^3 \</sup>verb|http://pubs.opengroup.org/onlinepubs/9699919799/basedefs/V1\_chap09.html#tag\_09\_04|$ 

Regular Expression patterns that use capturing groups (parentheses) will have their captured strings assigned to the BASH REMATCH variable for later retrieval.

Let's illustrate how regex can be used in Bash:

```
$ langRegex='(..)_(..)'
$ if [[ $LANG =~ $langRegex ]]
> then
> echo "Your country code (ISO 3166-1-alpha-2) is ${BASH_REMATCH[2]}."
> echo "Your language code (ISO 639-1) is ${BASH_REMATCH[1]}."
> else
> echo "Your locale was not recognised"
> fi
```

Be aware that regex parsing in Bash has changed between releases 3.1 and 3.2. Before 3.2 it was safe to wrap your regex pattern in quotes but this has changed in 3.2. Since then, regex should always be unquoted. You should protect any special characters by escaping it using a backslash. The best way to always be compatible is to put your regex in a variable and expand that variable in [ [ without quotes, as we showed above.

- Good Practice: Since the way regex is used in 3.2 is also valid in 3.1 we *highly* recommend you just never quote your regex. Remember to keep special characters properly escaped!
- For cross-compatibility (to avoid having to escape parentheses, pipes and so on) use a variable to store your regex, e.g. re='^\\*( >| \*Applying |.\*\.diff|.\*\.patch)'; [[ \$var =~ \$re ]] This is much easier to maintain since you only write ERE syntax and avoid the need for shell-escaping, as well as being compatible with all 3.x BASH versions.
- See also Chet Ramey's Bash FAQ<sup>1</sup>, section E14.
- In The Manual: Regex(3)<sup>2</sup>
- In the FAQ: I want to check if [[ \$var == foo or \$var == bar or \$var == more ... without repeating \$var n times.<sup>3</sup>
- Regular Expression: A regular expression is a more complex pattern that can be used to match specific strings (but unlike globs cannot expand to filenames).

#### 5.4 Brace Expansion

Then, there is *Brace Expansion*. Brace Expansion technically does not fit in the category of patterns, but it is similar. Globs only expand to actual filenames, but brace expansions will expand to any possible permutation of their contents. Here's how they work:

http://tiswww.case.edu/php/chet/bash/FAQ

<sup>2</sup>http://www.daemon-systems.org/man/regex.3.html

 $<sup>^{3}</sup>$ http://mywiki.wooledge.org/BashFAQ/066

```
$ echo th{e,a}n
then than
$ echo {/home/*,/root}/.*profile
/home/axxo/.bash_profile /home/lhunath/.profile /root/.bash_profile /root/.profile
$ echo {1..9}
1 2 3 4 5 6 7 8 9
$ echo {0,1}{0..9}
00 01 02 03 04 05 06 07 08 09 10 11 12 13 14 15 16 17 18 19
```

The brace expansion is replaced by a list of words, just like a glob is. However, these words aren't necessarily filenames, and they are not sorted (than would have come before then if they were).

Brace expansion happens *before* filename expansion. In the second echo command above, we used a combination of brace expansion and globs. The brace expansion goes first, and we get:

```
$ echo /home/*/.*profile /root/.*profile
```

After the brace expansion, the globs are expanded, and we get the filenames as the final result.

Brace expansions can only be used to generate lists of words. They cannot be used for pattern matching.

There are a few interesting and not very intuitive differences between ranges in character classes like [a-z] and brace expansion. For example, brace expansion allows counting backward, as can be seen with  $\{5..1\}$  or even  $\{b..Y\}$ , whereas [5-1] isn't expanded by the shell. [b-Y] may or may not be expanded, depending on your locale. Also, character ranges in brace expansions ignore locale variables like LANG and LC\_COLLATE and always use ASCII ordering. Globbing on the other hand is affected by language settings. The following fragment is an example for counting down and for displaying characters in the order of their ASCII codes:

```
$ echo {b..Y}
b a ` _ ^ ] [ Z Y
```

<sup>1</sup>http://mywiki.wooledge.org/locale

## 6 TestsAndConditionals

Sequential execution of commands is one thing, but to achieve any advanced logic in your scripts or your command line one-liners, you'll need tests and conditionals. Tests determine whether something is true or false. Conditionals are used to make decisions which determine the execution flow of a script.

#### **6.1** Exit Status

Every command results in an exit code whenever it terminates. This exit code is used by whatever application started it to evaluate whether everything went OK. This exit code is like a return value from functions. It's an integer between 0 and 255 (inclusive). Convention dictates that we use 0 to denote success, and any other number to denote failure of some sort. The specific number is entirely application-specific, and is used to hint as to what exactly went wrong.

For example, the ping command sends ICMP packets over the network to a certain host. That host normally responds to this packet by sending the exact same one right back. This way, we can check whether we can communicate with a remote host. ping has a range of exit codes which can tell us what went wrong, if anything did:

### From the Linux ping manual:

• If ping does not receive any reply packets at all it will exit with code 1. If a packet count and deadline are both specified, and fewer than count packets are received by the time the deadline has arrived, it will also exit with code 1. On other error it exits with code 2. Otherwise it exits with code 0. This makes it possible to use the exit code to see if a host is alive or not.

The special parameter? shows us the exit code of the last foreground process that terminated. Let's play around a little with ping to see its exit codes:

```
$ ping Good
ping: unknown host Good
$ echo $?
2
$ ping -c 1 -W 1 1.1.1.1
PING 1.1.1.1 (1.1.1.1) 56(84) bytes of data.
--- 1.1.1.1 ping statistics ---
1 packets transmitted, 0 received, 100% packet loss, time 0ms
$ echo $?
1
```

• Good Practice: You should make sure that your scripts always return a non-zero exit code if something unexpected happened in their execution. You can do this with the exit builtin:

```
rm file || { echo 'Could not delete file!' >&2; exit 1; }
```

• In The Manual: Exit Status<sup>1</sup>

http://www.gnu.org/software/bash/manual/bashref.html#Exit-Status

• Exit Code / Exit Status: Whenever a command ends it notifies its parent (which in our case will always be the shell that started it) of its exit status. This is represented by a number ranging from 0 to 255. This code is a hint as to the success of the command's execution.

## 6.2 Control Operators (&& and ||)

Now that we know what exit codes are, and that an exit code of '0' means the command's execution was successful, we'll learn to use this information. The easiest way of performing a certain action depending on the success of a previous command is through the use of *control operators*. These operators are && and ||, which respectively represent a logical AND and a logical OR. These operators are used between two commands, and they are used to control whether the second command should be executed depending on the success of the first. This concept is called *conditional execution*.

Let's put that theory in practice:

```
$ mkdir d && cd d
```

This simple example has two commands, mkdir d and cd d. You could use a semicolon there to separate the commands and execute them sequentially; but we want something more. In the above example, BASH<sup>1</sup> will execute mkdir d, then && will check the result of the mkdir application after it finishes. If the mkdir application was successful (exit code 0), then Bash will execute the next command, cd d. If mkdir d failed, and returned a non-0 exit code, Bash will skip the next command, and we will stay in the current directory.

Another example:

```
$ rm /etc/some_file.conf || echo "I couldn't remove the file"
rm: cannot remove `/etc/some_file.conf': No such file or directory
I couldn't remove the file
```

 $|\cdot|$  is much like &&, but it does the exact opposite. It only executes the next command if the first **failed**. As such, the message is only echoed if the rm command was unsuccessful.

In general, it's *not* a good idea to string together multiple different control operators in one command (we will explore this in the next section). & and  $| \cdot |$  are quite useful in simple cases, but not in complex ones. In the next few sections we'll show some other tools you can use for decision-making.

- Good Practice: It's best not to get overzealous when dealing with conditional operators. They can make your script hard to understand, especially for a person that's assigned to maintain it and didn't write it themselves.
- In The Manual: Lists of Commands<sup>2</sup>
- *Control Operators*: These operators are used to link commands together. They check the exit code of the previous command to determine whether or not to execute the next command in the sequence.

<sup>1</sup>http://mywiki.wooledge.org/BASH

 $<sup>^2</sup>$ http://www.gnu.org/software/bash/manual/bashref.html#Lists

## **6.3** Grouping Statements

Using conditional operators is easy and terse if we want to do simple error checking. Things get a bit more dangerous, though, when we want to run multiple statements if a condition holds true, or if we need to evaluate multiple conditions.

Suppose you want to delete a file if it contains a certain "good" word but also doesn't contain another "bad" word. Using grep (a command that checks its input for patterns), we translate these conditions to:

```
grep -q goodword "$file"  # exit status 0 (success) if "$file" contains 'goodword'
! grep -q "badword" "$file"  # exit status 0 (success) if "$file" does not contain 'badword'
```

We use -q (quiet) on grep because we don't want it to output the lines that match; we just want the exit code to be set.

The ! in front of a command causes Bash to *negate* the command's exit status. If the command returns 0 (success), the ! turns it into a failure. Likewise, if the command returns non-zero (failure), the ! turns it into a success.

Now, to put these conditions together and delete the file as a result of both holding true, we could use **Conditional Operators**:

```
$ grep -q goodword "$file" && ! grep -q badword "$file" && rm "$file"
```

This works great. (In fact, we can string together as many & & as we want, without any problems.) Now, imagine we want to show an error message in case the deletion of the file failed:

This looks OK, at first sight. If rm's exit code is not 0 (success), then the | | operator will trigger the next command and echo the error message (>&2: to standard error).

But there's a problem. When we have a sequence of commands separated by *Conditional Operators*, Bash looks at every one of them, in order from left to right. The exit status is carried through from whichever command was most recently executed, and skipping a command doesn't change it.

So, imagine the first grep fails (sets the exit status to 1). Bash sees a && next, so it skips the second grep altogether. Then it sees another &&, so it also skips the rm which follows that one. Finally, it sees a || operator. Aha! The exit status is "failure", and we have a ||, so Bash executes the echo command, and tells us that it couldn't delete a file -- even though it never actually *tried* to! That's not what we want.

This doesn't sound too bad when it's just a wrong error message you receive, but if you're not careful, this **will** eventually happen on more dangerous code. You wouldn't want to accidentally delete files or overwrite files as a result of a failure in your logic!

The failure in our logic is in the fact that we want the rm and the echo statements to belong together. The echo is related to the rm, not to the greps. So what we need is to *group* them. Grouping is done using curly braces:

```
$ grep -q goodword "$file" && ! grep -q badword "$file" && { rm "$file" || echo "Couldn't delete: \leftrightarrow $file" >&2; }
```

(Note: don't forget that you need a semicolon or newline before the closing curly brace!)

Now we've grouped the rm and echo command together. That effectively means the group is considered **one statement** instead of several. Going back to our situation of the first grep failing, instead of Bash trying the && rm "\$file"

statement, it will now try the && { . . . } statement. Since it is preceded by a && and the last command it ran failed (the failed grep), it will skip this group and move on.

Command grouping can be used for more things than just *Conditional Operators*. We may also want to group them so that we can redirect input to a group of statements instead of just one:

```
{
    read firstLine
    read secondLine
    while read otherLine; do
        something
    done
} < file</pre>
```

Here we're redirecting<sup>1</sup> file to a group of commands that read input. The file will be opened when the command group starts, stay open for the duration of it, and be closed when the command group finishes. This way, we can keep sequentially reading lines from it with multiple commands.

Another common use of grouping is in simple error handling:

```
# Check if we can go into appdir. If not, output an error and exit the script. cd "$appdir" || { echo "Please create the appdir and try again" >&2; exit 1; }
```

## 6.4 Conditional Blocks (if, test and [])

if is a shell keyword that executes a command (or a set of commands), and checks that command's exit code to see whether it was successful. Depending on that exit code, if executes a specific, different, block of commands.

```
$ if true
> then echo "It was true."
> else echo "It was false."
> fi
It was true.
```

Here you see the basic outline of an *if-statement*. We start by calling if with the command true. true is a builtin command that always ends successfully. if runs that command, and once the command is done, if checks the exit code. Since true always exits successfully, if continues to the then-block, and executes that code. Should the true command have failed somehow, and returned an unsuccessful exit code, the if statement would have skipped the then code, and executed the else code block instead.

Different people have different preferred styles for writing if statements. Here are some of the common styles:

```
if COMMANDS
then OTHER COMMANDS
fi

if COMMANDS
then
```

 $<sup>^{1} \</sup>verb|http://mywiki.wooledge.org/BashGuide/InputAndOutput\#Redirection|$ 

```
OTHER COMMANDS

fi

if COMMANDS; then
OTHER COMMANDS

fi
```

There are some commands designed specifically to *test* things and return an exit status based on what they find. The first such command is test (also known as [). A more advanced version is called [[. [ or test is a normal command that reads its arguments and does some checks with them. [[ is much like [, but it's special (a shell keyword), and it offers far more versatility. Let's get practical:

```
$ if [ a = b ]
> then echo "a is the same as b."
> else echo "a is not the same as b."
> fi
a is not the same as b.
```

if executes the command [ (remember, you don't **need** an if to run the [ command!) with the arguments a, =, b and ]. [ uses these arguments to determine what must be checked. In this case, it checks whether the string a (the first argument) is equal (the second argument) to the string b (the third argument), and if this is the case, it will exit successfully. However, since the string "a" is not equal to the string "b", [ will not exit successfully (its exit code will be 1). if sees that [ terminated unsuccessfully and executes the code in the else block.

The last argument, "]", means nothing to [, but it is required. See what happens when you omit it. Here's an example of a common pitfall when [ is used:

```
$ myname='Greg Wooledge' yourname='Someone Else'
$ [ $myname = $yourname ]
-bash: [: too many arguments
```

Can you guess what caused the problem?

[ was executed with the arguments Greg, Wooledge, =, Someone, Else and ]. That is 6 arguments, not 4! [ doesn't understand what test it's supposed to execute, because it expects either the first or second argument to be an operator. In our case, the operator is the third argument. Yet another reason why quotes are so terribly important. Whenever we type whitespace in Bash that belongs together with the words before or after it, we need to quote it, and the same thing goes for parameter expansions:

```
$ [ "$myname" = "$yourname" ]
```

This time, [ sees an operator (=) in the second argument and it can continue with its work.

To help us out a little, the Korn shell introduced (and Bash adopted) a new style of conditional test. Original as the Korn shell authors are, they called it [[. [[ is loaded with several very interesting features that [ lacks.

One of the features of [ [ is pattern matching:

```
$ [[ $filename = *.png ]] && echo "$filename looks like a PNG file"
```

<sup>1</sup>http://mywiki.wooledge.org/Quotes

Another feature of [ [ helps us in dealing with parameter expansions:

```
$ [[ $me = $you ]]  # Fine.
$ [[ I am $me = I am $you ]] # Not fine!
-bash: conditional binary operator expected
-bash: syntax error near `am'
```

This time, \$me and \$you did not need quotes. Since [ [ isn't a normal command (like [ is), but a *shell keyword*, it has special magical powers. It parses its arguments before they are expanded by Bash and does the expansion itself, taking the result as a single argument, even if that result contains whitespace. (In other words, [ [ does not allow word-splitting of its arguments.) *However*, be aware that simple strings still have to be quoted properly. [ [ treats a space outside of quotes as an argument separator, just like Bash normally would. Let's fix our last example:

```
$ [[ "I am $me" = "I am $you" ]]
```

Also; there is a subtle difference between quoting and not quoting the **right-hand side** of the comparison in [ [. The = operator does pattern matching by default, whenever the *right-hand side* is **not** quoted:

```
$ foo=[a-z]* name=lhunath $ [[ $name = $foo ]] && echo "Name $name matches pattern $foo" Name lhunath matches pattern <math>[a-z]* $ [[ $name = "$foo" ]] || echo "Name $name is not equal to the string $foo" Name lhunath is not equal to the string <math>[a-z]*
```

The first test checks whether \$name matches the *pattern* in \$foo. The second test checks whether \$name is equal to the *string* in \$foo. The quotes really do make that much difference -- a subtlety worth noting.

**Remember:** Quoting is usually going to give you the behavior that you want, so make it a habit; omit only when the specific situation requires unquoted behavior. Unfortunately, bugs caused by incorrect quoting are often hard to find, because code is often valid with or without quotes, but may have different meanings. In such cases, bash cannot tell that you did something wrong; it just does what you tell it, even if that's not what you intended.

You could also combine several if statements into one using elif instead of else, where each test indicates another possibility:

```
$ name=lhunath
$ if [[ $name = "George" ]]
> then echo "Bonjour, $name"
> elif [[ $name = "Hans" ]]
> then echo "Goeie dag, $name"
> elif [[ $name = "Jack" ]]
> then echo "Good day, $name"
> else
> echo "You're not George, Hans or Jack. Who the hell are you, $name?"
> fi
```

Note that "<" and ">" have special significance in bash. Pop quiz: Predict what happens when you do [ apple < banana ]. Test your hypothesis (don't cheat by trying without first forming a hypothesis!). Cue Jeopardy music... Answer: bash looks for a file named "banana" in the current directory so that its contents can be sent to [ apple (via standard input). Assuming you don't have a file named "banana" in your current directory, this will result in an error. Pop quiz: Assuming the original

intention of that command is determine whether "apple" comes before "banana", how would you change the command to get the desired effect? Cue Jeopardy music... Answer: escape the "<" with a backslash like that: [ apple \< banana ] or use [ [ instead of [.

Note that the comparison operators =, !=, >, and < treat their arguments as strings. In order for the operands to be treated as numbers, you need to use one of a different set of operators: -eq, -ne (not equal), -lt (less than), -gt, -le (less than or equal to), or -ge. Pop quiz: Come up with an example that shows the difference between < and -lt. Cue Jeopardy music... Since "314" comes before "9" lexicographically (i.e. the order that the dictionary would put them in), [ considers the former to be < than the later; whereas, [ considers "314" NOT to be -lt "9", because three hundred fourteen is NOT less than nine.

Now that you've got a decent understanding of quoting issues that may arise, let's have a look at some of the other features that [ and [ [ were blessed with:

- Tests supported by [ (also known as test) and [[:
  - **-e FILE**: True if file exists.
  - - f FILE: True if file is a regular file.
  - **-d FILE**: True if file is a directory.
  - **-h FILE**: True if file is a symbolic link.
  - -p PIPE: True if pipe exists.
  - **-r FILE**: True if file is readable by you.
  - -s FILE: True if file exists and is not empty.
  - -t FD: True if FD is opened on a terminal.
  - -w FILE: True if the file is writable by you.
  - -x FILE: True if the file is executable by you.
  - - O FILE: True if the file is effectively owned by you.
  - -G FILE: True if the file is effectively owned by your group.
  - FILE -nt FILE: True if the first file is newer than the second.
  - FILE -ot FILE: True if the first file is older than the second.
  - -z STRING: True if the string is empty (its length is zero).
  - - n STRING: True if the string is not empty (its length is not zero).
  - String operators:
    - \* **STRING = STRING**: True if the first string is identical to the second.
    - \* **STRING** != **STRING**: True if the first string is not identical to the second.
    - \* **STRING < STRING**: True if the first string sorts before the second.
    - \* **STRING > STRING**: True if the first string sorts after the second.
  - ! EXPR: Inverts the result of the expression (logical NOT).

- Numeric operators:
  - \* **INT -eq INT**: True if both integers are identical.
  - \* INT -ne INT: True if the integers are not identical.
  - \* **INT -lt INT**: True if the first integer is less than the second.
  - \* INT -gt INT: True if the first integer is greater than the second.
  - \* **INT -le INT**: True if the first integer is less than or equal to the second.
  - \* **INT -ge INT**: True if the first integer is greater than or equal to the second.
- Additional tests supported only by [ [:
  - **STRING** = (**or** ==) **PATTERN**: Not string comparison like with [ (or test), but *pattern matching* is performed. True if the string matches the glob pattern.
  - **STRING** != **PATTERN**: Not string comparison like with [ (or test), but *pattern matching* is performed. True if the string does not match the glob pattern.
  - STRING =~ REGEX: True if the string matches the regex pattern.
  - (EXPR): Parentheses can be used to change the evaluation precedence.
  - **EXPR && EXPR**: Much like the '-a' operator of test, but does not evaluate the second expression if the first already turns out to be false.
  - EXPR || EXPR: Much like the '-o' operator of test, but does not evaluate the second expression if the first already turns out to be true.
- Tests exclusive to [ (and test):
  - EXPR -a EXPR: True if both expressions are true (logical AND).
  - **EXPR** o **EXPR**: True if either expression is true (logical OR).

### Some examples? Sure:

```
$ test -e /etc/X11/xorg.conf && echo 'Your Xorg is configured!'
Your Xorg is configured!
$ test -n "$HOME" && echo 'Your homedir is set!'
Your homedir is set!
$ [[ boar != bear ]] && echo "Boars aren't bears."
Boars aren't bears!
$ [[ boar != b?ar ]] && echo "Boars don't look like bears."
$ [[ $DISPLAY ]] && echo "Your DISPLAY variable is not empty, you probably have Xorg running."
Your DISPLAY variable is not empty, you probably have Xorg running.
$ [[ ! $DISPLAY ]] && echo "Your DISPLAY variable is not not empty, you probably don't have Xorg running."
```

• Good Practice: Whenever you're making a Bash script, you should always use [[ rather than [. Whenever you're making a Shell script, which may end up being used in an environment where Bash is not available, you should use [, because it is far more portable. (While being built in to Bash and some other shells, [ should be available as an external application as well; meaning it will work as argument to, for example, find's -exec and xargs.) Don't ever use the -a or -o tests of the [ command. Use multiple [ commands instead (or use [[ if you can). POSIX doesn't define the behavior of [ with complex sets of tests, so you never know what you'll get.

```
if [ "$food" = apple ] && [ "$drink" = tea ]; then
  echo "The meal is acceptable."
fi
```

- In The Manual: Conditional Constructs<sup>1</sup>
- In the FAO:
- How can I group expressions, e.g. (A AND B) OR C?<sup>2</sup>
- What is the difference between the old and new test commands ([ and [[)?<sup>3</sup>
- How do I determine whether a variable contains a substring?<sup>4</sup>
- How can I tell whether a variable contains a valid number?<sup>5</sup>
- *if (keyword)*: Execute a list of commands and then, depending on their exit code, execute the code in the following then (or optionally else) block.

## 6.5 Conditional Loops (while, until and for)

Now you've learned how to make some basic decisions in your scripts. However, that's not enough for every kind of task we might want to script. Sometimes we need to repeat things. For that, we need to use a *loop*. There are two basic kinds of loops (plus a couple of variants), and using the correct kind of loop will help you keep your scripts readable and maintainable.

The two basic kinds of loops are called while and for. The while loop has a variant called until which simply reverses its check; and the for loop can appear in two different forms. Here's a summary:

- while *command*: Repeat so long as command is executed successfully (exit code is 0).
- until command: Repeat so long as command is executed unsuccessfully (exit code is not 0).
- for variable in words: Repeat the loop for each word, setting variable to each word in turn.

```
Ihttp://www.gnu.org/software/bash/manual/bashref.html#Conditional-Constructs
```

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/017

<sup>&</sup>lt;sup>3</sup>http://mywiki.wooledge.org/BashFAQ/031

<sup>4</sup>http://mywiki.wooledge.org/BashFAQ/041

<sup>&</sup>lt;sup>5</sup>http://mywiki.wooledge.org/BashFAQ/054

• for ((expression; expression; expression)): Starts by evaluating the first arithmetic expression; repeats the loop so long as the second arithmetic expression is successful; and at the end of each loop evaluates the third arithmetic expression.

Each loop form is followed by the key word do, then one or more commands in the *body*, then the key word done. The do and done are similar to the then and fi (and possible elif and/or else) from the if statement we saw earlier. Their job is to tell us where the body of the loop begins and ends.

In practice, the loops are used for different kinds of tasks. The for loop (first form) is appropriate when we have a list of things, and we want to run through that list sequentially. The while loop is appropriate when we don't know exactly how many times we need to repeat something; we simply want it to keep going until we find what we're looking for.

Here are some examples to illustrate the differences and also the similarities between the loops. (Remember: on most operating systems, you press **Ctrl-C** to kill a program that's running on your terminal.)

```
$ while true
> do echo "Infinite loop"
> done
```

```
$ while ! ping -c 1 -W 1 1.1.1.1; do
> echo "still waiting for 1.1.1.1"
> sleep 1
> done
```

```
$ (( i=10 )); while (( i > 0 ))
> do echo "$i empty cans of beer."
> (( i-- ))
> done
$ for (( i=10; i > 0; i-- ))
> do echo "$i empty cans of beer."
> done
$ for i in {10..1}
> do echo "$i empty cans of beer."
> done
```

The last three loops achieve exactly the same result, using different syntax. You'll encounter this many times in your shell scripting experience. There will nearly always be multiple approaches to solving a problem. The test of your skill soon won't be about solving a problem as much as about how *best* to solve it. You must learn to pick the best angle of approach for the job. Usually, the main factors to take into account will be the simplicity and flexibility of the resulting code. My personal favorite is the last of the examples. In that example I used *Brace Expansion* to generate the words; but there are other ways, too.

Let's take a closer look at that last example, because although it looks the easier of the two fors, it can often be the trickier, if you don't know exactly how it works.

As I mentioned before: for runs through a list of words and puts each one in the loop index variable, one at a time, and then loops through the body with it. The tricky part is how Bash decides what the words are. Let me explain myself by expanding the braces from that previous example:

```
$ for i in 10 9 8 7 6 5 4 3 2 1
> do echo "$i empty cans of beer."
> done
```

Bash takes the characters between in and the end of the line, and splits them up into words. This splitting is done on spaces and tabs, just like argument splitting. However, if there are any unquoted substitutions in there, they will be word-split as well (using IFS<sup>1</sup>). All these split-up words become the iteration elements.

As a result, be VERY careful not to make the following mistake:

```
$ ls
The best song in the world.mp3
$ for file in $(ls *.mp3)
> do rm "$file"
> done
rm: cannot remove `The': No such file or directory
rm: cannot remove `best': No such file or directory
rm: cannot remove `song': No such file or directory
rm: cannot remove `in': No such file or directory
rm: cannot remove `in': No such file or directory
rm: cannot remove `the': No such file or directory
rm: cannot remove `the': No such file or directory
```

You should already know to quote the <code>\$file</code> in the rm statement; but what's going wrong here? Bash expands the command substitution (<code>\$(ls \*.mp3))</code>, replaces it by its output, and then performs word splitting on it (because it was unquoted). Essentially, Bash executes for file in The best song in the world.mp3. Boom, you are dead.

You want to quote it, you say? Let's add another song:

Quotes will indeed protect the whitespace in your filenames; but they will do more than that. The quotes will protect **all the whitespace** from the output of ls. There is no way Bash can know which parts of the output of ls represent filenames; it's not psychic. The output of ls is a simple string, and Bash treats it as such. The for puts the whole quoted output in i and runs the rm command with it. *Damn, dead again*.

So what do we do? As suggested earlier, globs are your best friend:

```
$ for file in *.mp3
> do rm "$file"
> done
```

<sup>1</sup>http://mywiki.wooledge.org/IFS

This time, Bash **does** know that it's dealing with filenames, and it **does** know what the filenames are, and as such it can split them up nicely. The result of expanding the glob is this: for file in "The best song in the world.mp3" "The worst song in the world.mp3". Problem solved!

Now let's look at the while loop. The while loop is very interesting for its capacity to execute commands until something interesting happens. Here are a few examples of how while loops are very often used:

```
$ # The sweet machine; hand out sweets for a cute price.
$ while read -p $'The sweet machine.\nInsert 20c and enter your name: ' name
> do echo "The machine spits out three lollipops at $name."
> done

$ # Check your email every five minutes.
$ while sleep 300
> do kmail --check
> done

$ # Wait for a host to come back online.
$ while ! ping -c 1 -W 1 "$host"
> do echo "$host is still unavailable."
> done; echo -e "$host is available again.\a"
```

The until loop is barely ever used, if only because it is pretty much exactly the same as while !. We could rewrite our last example using an until loop:

```
$ # Wait for a host to come back online.
$ until ping -c 1 -W 1 "$host"
> do echo "$host is still unavailable."
> done; echo -e "$host is available again.\a"
```

In practice, most people simply use while! instead.

Lastly, you can use the continue builtin to skip ahead to the next iteration of a loop without executing the rest of the body, and the break builtin to jump out of the loop and continue with the script after it. This works in both for and while loops.

- In The Manual: Looping Constructs<sup>1</sup>
- In the FAQ:
- How can I run a command on all files with the extension .gz?<sup>2</sup>
- How can I use numbers with leading zeros in a loop, e.g. 01, 02?<sup>3</sup>
- How can I find and deal with file names containing newlines, spaces or both?<sup>4</sup>

```
1http://www.gnu.org/software/bash/manual/bashref.html#Looping-Constructs
2http://mywiki.wooledge.org/BashFAQ/015
3http://mywiki.wooledge.org/BashFAQ/018
```

<sup>4</sup>http://mywiki.wooledge.org/BashFAQ/020

- How can I rename all my \*.foo files to \*.bar, or convert spaces to underscores, or convert upper-case file names to lower case?<sup>1</sup>
- Can I do a spinner in Bash?<sup>2</sup>
- I want to check to see whether a word is in a list (or an element is a member of a set).<sup>3</sup>
- *Loop*: A loop is a structure that is designed to repeat the code within until a certain condition has been fulfilled. At that point, the loop stops and the code beyond it is executed.
- for (keyword): A for-loop is a type of loop that sets a variable to each of a list of values in turn, and repeats until the list is exhausted.
- while (keyword): A while-loop is a type of loop that continues to run its code so long as a certain command (run before each iteration) executes successfully.
- *until* (*keyword*): An until-loop is a type of loop that continues to run its code so long as a certain command (run before each iteration) executes unsuccessfully.

## 6.6 Choices (case and select)

Sometimes you want to build application logic depending on the content of a variable. This could be implemented by taking a different branch of an if statement depending on the results of testing against a glob:

```
shopt -s extglob
   if [[ \$LANG = en* ]]; then
       echo 'Hello!'
   elif [[ $LANG = fr* ]]; then
5
       echo 'Salut!'
   elif [[ $LANG = de* ]]; then
       echo 'Guten Tag!'
   elif [[ \$LANG = nl* ]]; then
9
       echo 'Hallo!'
10
   elif [[ $LANG = it* ]]; then
11
      echo 'Ciao!'
12
   elif [[ $LANG = es* ]]; then
       echo 'Hola!'
14
   elif [[ $LANG = @(C|POSIX) ]]; then
15
       echo 'hello world'
16
17
       echo 'I do not speak your language.'
18
19
```

http://mywiki.wooledge.org/BashFAQ/030

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/034

<sup>3</sup>http://mywiki.wooledge.org/BashFAQ/046

But all these comparisons are a bit redundant. Bash provides a keyword called case exactly for this kind of situation. A case statement basically enumerates several possible *Glob Patterns* and checks the content of your parameter against these:

```
case $LANG in
en*) echo 'Hello!';;
fr*) echo 'Salut!';;
de*) echo 'Guten Tag!';;
nl*) echo 'Hallo!';;
it*) echo 'Ciao!';;
es*) echo 'Hola!';;
C|POSIX) echo 'hello world';;

* esac
esac
```

Each choice in a case statement consists of a pattern (or a list of patterns with | between them), a right parenthesis, a block of code that is to be executed if the string matches one of those patterns, and two semi-colons to denote the end of the block of code (since you might need to write it on several lines). A left parenthesis can be added to the left of the pattern. Using ; & instead of ;; will grant you the ability to fall-through the case matching in bash, zsh and ksh. case stops matching patterns as soon as one is successful. Therefore, we can use the \* pattern in the end to match any case that has not been caught by the other choices.

When a matching pattern is found and the ; & operator is used after the block of code, then the block of code for the next choice (if any) will be executed, too, no matter if the pattern for that choice matches or not. To "fall through" several consecutive patterns, use the ; & operator for all but the last.

When a matching pattern is found and the ;; & operator is used after the block of code, instead of the block of code for the *next* choice, the block of code for the *next matching* pattern (if any) will be executed.

Another construct of choice is the select construct. This statement smells like a loop and is a convenience statement for generating a menu of choices that the user can choose from.

The user is presented by choices and asked to enter a number reflecting his choice. The code in the select block is then executed with a variable set to the choice the user made. If the user's choice was invalid, the variable is made empty:

```
$ echo "Which of these does not belong in the group?"; \
> select choice in Apples Pears Crisps Lemons Kiwis; do
> if [[ $choice = Crisps ]]
> then echo "Correct! Crisps are not fruit."; break; fi
> echo "Errr... no. Try again."
> done
```

The menu reappears so long as the break statement is not executed. In the example the break statement is only executed when the user makes the correct choice.

We can also use the PS3 variable to define the prompt the user replies on. Instead of showing the question before executing the select statement, we could choose to set the question as our prompt:

```
$ PS3="Which of these does not belong in the group (#)? "; \
> select choice in Apples Pears Crisps Lemons Kiwis; do
> if [[ $choice = Crisps ]]
> then echo "Correct! Crisps are not fruit."; break; fi
> echo "Errr... no. Try again."
```

All of these conditional constructs (if, for, while, and case) can be *nested*. This means you could have a for loop with a while loop inside it, or any other combination, as deeply as you need to solve your problem.

```
1
  # A simple menu:
   while true; do
      echo "Welcome to the Menu"
       echo " 1. Say hello"
      echo " 2. Say good-bye"
6
      read -p "-> " response
       case $response in
          1) echo 'Hello there!' ;;
           2) echo 'See you later!'; break ;;
10
           *) echo 'What was that?' ;;
11
12
       esac
  done
13
14
   # Alternative: use a variable to terminate the loop instead of an
15
   # explicit break command.
17
18
  while test -z "$quit"; do
19
    echo "...."
20
       read -p "-> " response
21
       case $response in
22
23
           2) echo 'See you later!'; quit=y ;;
24
25
           . . .
26
       esac
  done
27
```

- Good Practice: A select statement makes a simple menu simple, but it doesn't offer much flexibility. If you want something more elaborate, you might prefer to write your own menu using a while loop, some echo or printf commands, and a read command.
- In The Manual: Conditional Constructs<sup>1</sup>
- In the FAQ:
- I want to check if [[ \$var == foo or \$var == bar or \$var = more ... without repeating \$var n times.<sup>2</sup>
- How can I handle command-line arguments (options) to my script easily?<sup>3</sup>

<sup>1</sup> http://www.gnu.org/software/bash/manual/bashref.html#Conditional-Constructs

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/066

 $<sup>^3</sup>$ http://mywiki.wooledge.org/BashFAQ/035

- case (keyword): The case statement evaluates a parameter's value against several given patterns (choices).
- *select (keyword)*: The select statement offers the user the choice of several options and executes a block of code with the user's choice in a parameter. The menu repeats until a break command is executed.

## 7 Arrays

As mentioned earlier<sup>1</sup>, Bash<sup>2</sup> has multiple parameter types. Of importance here, variables can either contain a single value (scalars) or multiple values (arrays).

Strings are without a doubt the most used parameter type. But they are also the most misused parameter type. It is important to remember that a string holds just **one** element. Capturing the output of a command, for instance, and putting it in a string parameter means that parameter holds just **one** string of characters, regardless of whether that string represents twenty filenames, twenty numbers or twenty names of people.

And as is always the case when you put multiple items in a single string, these multiple items must be somehow delimited from each other. We, as humans, can usually decipher what the different filenames are when looking at a string. We assume that, perhaps, each line in the string represents a filename, or each word represents a filename. While this assumption is understandable, it is also inherently flawed. Each single filename can contain every character you might want to use to separate the filenames from each other in a string. That means there's technically no telling where the first filename in the string ends, because there's no character that can say: "I denote the end of this filename" because that character itself could be part of the filename.

Often, people make this mistake:

```
# This does NOT work in the general case
$ files=$(ls ~/*.jpg); cp $files /backups/
```

When this would probably be a better idea (using array notation, which is explained later, in the next section):

```
# This DOES work in the general case
$ files=(~/*.jpg); cp "${files[@]}" /backups/
```

The first attempt at backing up our files in the current directory is flawed. We put the output of ls in a string called files and then use the **unquoted** \$files parameter expansion to cut that string into arguments (relying on *Word Splitting*). As mentioned before, argument and word splitting cuts a string into pieces wherever there is whitespace. Relying on it means we assume that none of our filenames will contain any whitespace. If they do, the filename will be cut in half or more. Conclusion: **bad** 

The only safe way to represent **multiple** string elements in Bash is through the use of arrays. An array is a type of variable that **maps integers to strings**. That basically means that it holds a numbered list of strings. Since each of these strings is a separate entity (element), it can safely contain any character, even whitespace.

For the best results and the least headaches, remember that if you have a list of things, you should always put it in an array. Unlike some other programming languages, Bash does not offer lists, tuples, etc. Just arrays, and associative arrays (which are new in Bash 4).

• Array: An array is a numbered list of strings: It maps integers to strings.

http://mywiki.wooledge.org/BashGuide/Parameters#Variable\_Types

<sup>2</sup>http://mywiki.wooledge.org/Bash

## 7.1 Creating Arrays

There are several ways you can create or fill your array with data. There is no one single true way: the method you'll need depends on where your data comes from and what it is.

The easiest way to create a simple array with data is by using the = () syntax:

```
$ names=("Bob" "Peter" "$USER" "Big Bad John")
```

This syntax is great for creating arrays with static data or a known set of string parameters, but it gives us very little flexibility for adding lots of array elements. If you need more flexibility, you can also specify explicit indexes:

```
$ names=([0]="Bob" [1]="Peter" [20]="$USER" [21]="Big Bad John")
# or...
$ names[0]="Bob"
```

Notice that there is a gap between indices 1 and 20 in this example. An array with holes in it is called a *sparse array*. Bash allows this, and it can often be quite useful.

If you want to fill an array with filenames, then you'll probably want to use Globs in there:

```
$ photos=(~/"My Photos"/*.jpg)
```

Notice here that we quoted the My Photos part because it contains a space. If we hadn't quoted it, Bash would have split it up into photos=('~/My' 'Photos/'\*.jpg') which is obviously **not** what we want. Also notice that we quoted **only** the part that contained the space. That's because we cannot quote the ~ or the \*; if we do, they'll become literal and Bash won't treat them as special characters anymore.

Unfortunately, its really easy to equivocally create arrays with a bunch of filenames in the following way:

```
$ files=$(ls)  # BAD, BAD!
$ files=($(ls))  # STILL BAD!
```

Remember to **always avoid** using ls. The first would create a **string** with the output of ls. That string cannot possibly be used safely for reasons mentioned in the Arrays introduction. The second is closer, but it still splits up filenames with whitespace.

This is the right way to do it:

```
$ files=(*) # Good!
```

This statement gives us an array where each filename is a separate element. The \* is a glob pattern for any string which pathname-expands into all the filenames in the current directory (just like it would in eg. rm \*). After the pathname expansion, the command will look like files=([each file in the current directory that matches \*]) which assigns all of the files to the array files. Perfect!

This section that we're about to introduce contains some advanced concepts. If you get lost, you may want to return here after you've read the whole guide. You can skip ahead to Using Arrays if you want to keep things simple.

Now, sometimes we want to build an array from a string or the output of a command. Commands (generally) just output strings: for instance, running a find command will enumerate filenames, and separate these filenames with newlines (putting each filename on a separate line). So to parse that one big string into an array we need to tell Bash where each element ends.

(Note, this is a bad example, because filenames can **contain** a newline, so it is not safe to delimit them with newlines! But see below.)

Breaking up a string is what IFS is used for:

```
$ IFS=. read -ra ip_elements <<< "127.0.0.1"
```

Here we use IFS with the value. to cut the given IP address into array elements wherever there's a ., resulting in an array with the elements 127, 0, 0 and 1. (The builtin command read and the <<< operator will be covered in more depth in the Input and Output chapter.)

We could do the same thing with a find command, by setting IFS to a newline. But then our script would fail when someone creates a filename with a newline in it (either accidentally or maliciously).

So, is there any way to get a list of elements from an external program (like find) into a Bash array? In general, the answer is yes, provided there is a reliable way to delimit the elements.

In the specific case of filenames, the answer to this problem is NUL bytes. A NUL byte is a byte which is just all zeros: 00000000. Bash strings can't contain NUL bytes, because of an artifact of the "C" programming language: NUL bytes are used in C to mark the end of a string. Since Bash is written in C and uses C's native strings, it inherits that behavior.

A data stream (like the output of a command, or a file) can contain NUL bytes. Streams are like strings with three big differences: they are read sequentially (you usually can't jump around); they're *unidirectional* (you can read from them, or write to them, but typically not both); and they can contain NUL bytes.

File *names* cannot contain NUL bytes (since they're implemented as C strings by Unix), and neither can the vast majority of human-readable things we would want to store in a script (people's names, IP addresses, etc.). That makes NUL a great candidate for separating elements in a stream. Quite often, the command whose output you want to read will have an option that makes it output its data separated by NUL bytes rather than newlines or something else. find (on GNU and BSD, anyway) has the option <code>-print0</code>, which we'll use in this example:

```
$ files=()
$ while read -r -d ''; do
> files+=("$REPLY")
> done < <(find /foo -print0)</pre>
```

This is a safe way of parsing a command's output into strings. Understandably, it looks a little confusing and convoluted at first. So let's take it apart:

- The first line files=() creates an empty array named files.
- We're using a while loop<sup>1</sup> that runs a read command each time. The read command uses the -d '' option to specify the delimiter and it interprets the empty string as a NUL byte (\0) (as Bash arguments can not contain NULs). This means that instead of reading a line at a time (up to a newline), we're reading up to a NUL byte. It also uses -r to prevent it from treating backslashes specially.
- Once read has read some data and encountered a NUL byte, the while loop's body is executed. We put what we read (which is in the parameter REPLY) into our array.

<sup>1</sup>http://mywiki.wooledge.org/BashGuide/TestsAndConditionals#Conditional\_Loops

- To do this, we use the += () syntax. This syntax adds one or more element(s) to the end of our array.
- Finally, the < <(...) syntax is a combination of *File Redirection* (<) and *Process Substitution* (<(...)). Omitting the technical details for now, we'll simply say that this is how we send the output of the find command into our while loop.

The find command itself uses the -print0 option as mentioned before to tell it to separate the filenames it finds with a NUL byte.

As an aside, check out globstar<sup>1</sup> if you are using bash >= 4.0 and just want to recursively walk directories.

- Good Practice: Arrays are a safe list of strings. They are perfect for storing multiple filenames. If you have to parse a stream of data into component elements, there must be a way to tell where each element starts and ends. The NUL byte is very often the best choice for this job. If you have a list of things, keep it in list form as long as possible. Don't smash it into a string or a file until you absolutely have to. If you do have to write it out to a file and read it back in later, keep in mind the delimiter problem we mentioned above.
- In The Manual: Arrays<sup>2</sup>
- In the FAQ: How can I use array variables?<sup>3</sup> How can I use variable variables (indirect variables, pointers, references) or associative arrays?<sup>4</sup> How can I find and deal with file names containing newlines, spaces or both?<sup>5</sup> I set variables in a loop. Why do they suddenly disappear after the loop terminates? Or, why can't I pipe data to read?<sup>6</sup>

### 7.2 Using Arrays

Once we have an array, there are several things we can do with it.

## 7.2.1 Expanding Elements

First of all, we can print the contents to see what's in it:

```
$ declare -p myfiles
declare -a myfiles='([0]="/home/wooledg/.bashrc" [1]="billing codes.xlsx" [2]="hello.c")'
```

The declare -p command prints the contents of one or more variables. In addition, it shows you what Bash thinks the *type* of the variable is, and it does all of this using code that you could copy and paste into a script. In our case, the -a means this is an array. There are three elements, with indices 0, 1 and 2, and we can see what each one contains.

If we want something a bit less technical, we can also print the array using printf:

```
Inttp://mywiki.wooledge.org/glob#globstar_.28since_bash_4.0-alpha.29
Inttp://www.gnu.org/software/bash/manual/bashref.html#Arrays
Inttp://mywiki.wooledge.org/BashFAQ/005
Inttp://mywiki.wooledge.org/BashFAQ/006
Inttp://mywiki.wooledge.org/BashFAQ/020
Inttp://mywiki.wooledge.org/BashFAQ/024
```

```
$ printf '%s\n' "${myfiles[@]}"
/home/wooledg/.bashrc
billing codes.xlsx
hello.c
```

This prints each array element, in order by index, with a newline after each one. Note that if one of the array elements happens to *contain* a newline character, we won't be able to tell where each element starts and ends, or even how many there are. That's why it's important to keep our data safely contained in the array as long as possible. Once we print it out, there's no way to reverse that.

The syntax "\${myfiles[@]}" is extremely important. It works just like "\$@" does for the positional parameters: it expands to a list of words, with each array element as *one* word, no matter what it contains. Even if there are spaces, tabs, newlines, quotation marks, or any other kind of characters in one of the array elements, it'll still be passed along as one word to whichever command we're running.

The printf command implicitly loops over all of its arguments. But what if we wanted to do our own loop? In that case, we can use a for loop<sup>1</sup> to iterate over the elements:

```
$ for file in "${myfiles[@]}"; do
> cp "$file" /backups/
> done
```

We use the **quoted** form again here: " $\{myfiles[@]\}$ ". Bash replaces this syntax with each element in the array properly quoted – similar to how positional parameters (arguments that were passed to the current script or function) are expanded.

The following two examples have the same effect:

```
$ names=("Bob" "Peter" "$USER" "Big Bad John")
$ for name in "${names[@]}"; do echo "$name"; done
$ for name in "Bob" "Peter" "$USER" "Big Bad John"; do echo "$name"; done
```

The first example creates an array named names which is filled up with a few elements. Then the array is expanded into these elements, which are then used by the for loop. In the second example, we skipped the array and just passed the list of elements directly to for.

Remember to **quote** the \${arrayname[@]} expansion properly. If you don't, you'll lose all benefit of having used an array at all: leaving arguments unquoted means you're telling Bash it's OK to wordsplit them into pieces and break everything again.

The above example expanded the array in a for-loop statement. But you can expand the array anywhere you want to put its elements as arguments; for instance in a cp command:

```
$ myfiles=(db.sql home.tbz2 etc.tbz2)
$ cp "${myfiles[@]}" /backups/
```

This runs the cp command, replacing the "\${myfiles[@]}" part with every filename in the myfiles array, properly quoted. After expansion, Bash will effectively run this:

 $<sup>^{1} \</sup>verb|http://mywiki.wooledge.org/BashGuide/TestsAndConditionals\#Conditional\_Loops|$ 

```
$ cp "db.sql" "home.tbz2" "etc.tbz2" /backups/
```

cp will then copy the files to your /backups/ directory.

Of course, a for loop offers the ultimate flexibility, but printf and its implicit looping over arguments can cover many of the simpler cases. It can even produce NUL-delimited streams, perfect for later retrieval:

```
$ printf "%s\0" "${myarray[@]}" > myfile
```

You can also expand single array elements by referencing their element number (called **index**). Remember that by default, arrays are *zero-based*, which means that their **first element** has the index **zero**:

```
$ echo "The first name is: ${names[0]}"
$ echo "The second name is: ${names[1]}"
```

(You could create an array with no element 0. Remember what we said about *sparse arrays* earlier -- you can have "holes" in the sequence of indices, and this applies to the beginning of the array as well as the middle. It's your responsibility as the programmer to know which of your arrays are potentially sparse, and which ones are not.)

There is also a second form of expanding all array elements, which is " $\{arrayname[*]\}$ ". This form is **ONLY** useful for converting arrays into a single string with all the elements joined together. The main purpose for this is outputting the array to humans:

```
$ names=("Bob" "Peter" "$USER" "Big Bad John")
$ echo "Today's contestants are: ${names[*]}"
Today's contestants are: Bob Peter lhunath Big Bad John
```

Notice that in the resulting string, there's no way to tell where the names begin and end! This is why we keep everything separate as long as possible.

Remember to still keep everything nicely **quoted**! If you don't keep \${arrayname[\*]} quoted, once again Bash's WordSplitting<sup>1</sup> will cut it into bits.

You can combine IFS with "\${arrayname[\*]}" to indicate the character to use to delimit your array elements as you merge them into a single string. This is handy, for example, when you want to comma delimit names:

```
$ names=("Bob" "Peter" "$USER" "Big Bad John")
$ ( IFS=,; echo "Today's contestants are: ${names[*]}" )
Today's contestants are: Bob,Peter,lhunath,Big Bad John
```

Notice how in this example we put the IFS=,; echo ... statement in a Subshell<sup>2</sup> by wrapping (and) around it. We do this because we don't want to change the default value of IFS in the main shell. When the subshell exits, IFS still has its default value and no longer just a comma. This is important because IFS is used for a lot of things, and changing its value to something non-default will result in very odd behavior if you don't expect it!

Alas, the "\${array[\*]}" expansion only uses the *first* character of IFS to join the elements together. If we wanted to separate the names in the previous example with a comma and a space, we would have to use some other technique (for example, a for loop).

One final tip: you can get the number of elements of an array by using \${ #array [@] }

http://mywiki.wooledge.org/WordSplitting

 $<sup>^{2} \</sup>verb|http://mywiki.wooledge.org/BashGuide/CompoundCommands\#Subshells|$ 

```
$ array=(a b c)
$ echo ${#array[@]}
3
```

#### 7.2.2 Expanding Indices

Sometimes a problem requires more than just expanding the values of an array in order. You may need to refer to multiple elements at the same time, or refer to the same index in multiple arrays at the same time. In these cases, it's better to expand the array indices, instead of the array values.

Let's say we have two arrays: first and last. These will hold the first names, and the last names, of a list of people. Obviously we need to make sure that the first and last names of each person are properly matched up. We do this by keeping careful control of the indices.

```
$ first=(Jessica Sue Peter)
$ last=(Jones Storm Parker)
```

Now, to print the full name of the person with index 1:

```
$ echo "${first[1]} ${last[1]}"
Sue Storm
```

If we want to loop over all of the names, we can't just loop over "\${first[@]}" or "\${last[@]}". If we do that, we won't have an index into the *other* array, so we won't know how to match them up. Instead, we'll loop over the indices of one of the arrays (arbitrarily chosen), and then use that same index in both arrays together:

```
$ for i in "${!first[@]}"; do
> echo "${first[i]} ${last[i]}"
> done
Jessica Jones
Sue Storm
Peter Parker
```

So, we have a new piece of syntax: "\${!arrayname[@]}" expands to a list of the *indices* of an array, in sequential order.

Another feature worth mentioning is that the [...] around the index of an array actually creates an arithmetic context.

You can do math there, without wrapping it in \$((...)). Let's suppose we want to process an array two elements at a time, and let's also suppose we know this array can't be *sparse*. Then:

```
$ a=(a b c q w x y z)
$ for ((i=0; i<${#a[@]}; i+=2)); do
> echo "${a[i]} and ${a[i+1]}"
> done
```

We use the arithmetic expression i+1 as an array index. Bash will evaluate the i parameter first, and keep evaluating the value it receives as long as it is a valid *Name*, until it gets to an integer. Then it will add 1, and use that as the real index.

### 7.2.3 Sparse Arrays

We've mentioned sparse arrays already, so this will be brief. Most arrays simply have indices 0, 1, 2, 3, etc. But an array can also have "holes" in the sequence. This can be done either by assigning directly to an index that's way out past the current end of the array, or by *removing* an element from an existing array.

```
$ nums=(zero one two three four)
$ nums[70]="seventy"
$ unset 'nums[3]'
$ declare -p nums
declare -a nums='([0]="zero" [1]="one" [2]="two" [4]="four" [70]="seventy")'
```

Note that we quoted 'nums[3]' in the unset command. This is because an *unquoted* nums[3] could be interpreted by Bash as a filename glob. If it happens to match a file in the current directory, then nums[3] becomes nums3 and we will unset the wrong variable. That would be bad!

If you follow the practices we've outlined so far, sparse arrays shouldn't cause you any special concerns.

- Don't assume that your indices are sequential.
- If the index values matter, always iterate over the indices instead of making assumptions about them.
- If you loop over the values instead, don't assume anything about which index you might be on currently.
- In particular, don't assume that just because you're currently in the first iteration of your loop, that you must be on index 0!

When you expand the values of a sparse array using "\${arrayname[@]}" you will get a list with no gaps in it. There is no way to tell what kind of array these values came from. This can be useful if you want to *re-index* an array to remove all of the gaps:

```
$ array=("${array[@]}") # This re-creates the indices.
```

• Good Practice: Always quote your array expansions properly, just like you would your normal parameter expansions. Use "\${myarray[@]}" to expand all your array elements and ONLY use "\${myarray[\*]}" when you want to merge all your array elements into a single string.

## 7.3 Associative Arrays

Until recently, Bash could only use numbers (more specifically, non-negative integers) as keys of arrays. This means you could not "map" or "translate" one string to another. This is something a lot of people missed. People began to (ab)use variable indirection as a means to address the issue.

Since Bash 4 was released, there is no longer any excuse to use indirection (or **worse**, eval) for this purpose. You can now use full-featured associative arrays.

<sup>1</sup>http://mywiki.wooledge.org/BashFAQ/006

To create an associative array, you need to declare it as such (using declare -A). This is necessary, because otherwise bash doesn't know what kind of array you're trying to make. Here's how you make an associative array:

```
$ declare -A fullNames
$ fullNames=( ["lhunath"]="Maarten Billemont" ["greycat"]="Greg Wooledge" )
$ echo "Current user is: $USER. Full name: ${fullNames[$USER]}."
Current user is: lhunath. Full name: Maarten Billemont.
```

We can print the contents of an associative array very much like we did with regular arrays:

```
$ declare -A dict
$ dict[astro]="Foo Bar"
$ declare -p dict
declare -A dict='([astro]="Foo Bar")'
```

With the same syntax as for indexed arrays, you can iterate over the keys (indices) of associative arrays:

```
$ for user in "${!fullNames[@]}"
> do echo "User: $user, full name: ${fullNames[$user]}."; done
User: lhunath, full name: Maarten Billemont.
User: greycat, full name: Greg Wooledge.
```

Two things to remember, here: First, the order of the keys you get back from an associative array using the "\${ !array[@]}" syntax is unpredictable; it won't necessarily be the order in which you assigned elements, or any kind of sorted order. Likewise, if you expand the elements using "\${array[@]}" you will get them in an unpredictable order. Associative arrays are not well suited to storing lists that need to be processed in a specific order.

Second, you cannot omit the \$ if you're using a parameter as the key of an associative array. With standard indexed arrays, the [...] part is an arithmetic context. In an arithmetic context, a *Name* can't possibly be a valid number, and so BASH assumes it's a parameter and that you want to use its content. This doesn't work with associative arrays, since a *Name* could just as well be a valid associative array key.

Let's demonstrate with examples:

```
$ indexedArray=( "one" "two" )
$ declare -A associativeArray=( ["foo"]="bar" ["alpha"]="omega" )
$ index=0 key="foo"
$ echo "${indexedArray[$index]}"
one
$ echo "${indexedArray[index]}"
one
$ echo "${indexedArray[index + 1]}"
two
$ echo "${associativeArray[$key]}"
bar
$ echo "${associativeArray[key]}"
$ echo "${associativeArray[key]}"
```

# 8 Input and Output

Input and output in Bash scripts is a complex topic, because there is a great deal of flexibility in how it's done. This chapter will only scratch the surface of what is possible.

*Input* refers to any information that your program receives (or reads). Input to a Bash script can come from several different places:

- Command-line arguments (which are placed in the positional parameters)
- Environment variables, inherited from whatever process started the script
- Files
- Anything else a File Descriptor can point to (pipes, terminals, sockets, etc.). This will be discussed below.

*Output* refers to any information that your program produces (or writes). Output from a Bash script can also go to lots of different places:

- Files
- Anything else a File Descriptor can point to
- Command-line arguments to some other program
- Environment variables passed to some other program

Input and output are important in shell script programming. Figuring out where your input comes from, what it looks like, and what you must do to it in order to produce your desired output are core requirements for almost all scripts.

## 8.1 Command-line Arguments

For many scripts, the first (or the only) input we will care about are the arguments received by the script on the command line. As we saw in the Parameters chapter, there are some *Special Parameters* available to every script which contain these arguments. These are called the *Positional Parameters*. They are a very simple numerically indexed array of strings (in fact, in the POSIX shell, they are the *only* array the shell has). The first positional parameter is referred to with \$1; the second, with \$2; and so on. After the 9th one, you must use curly braces to refer to them:  $$\{10\}, $\{11\},$  etc. But in practice, it's exceedingly rare that you would ever need to do that, because there are better ways to deal with them as a group.

In addition to referring to them one at a time, you may also refer to the entire set of positional parameters with the "\$@" substitution. The double quotes here are **extremely important**. If you don't use the double quotes, each one of the positional parameters will undergo word splitting and globbing. You don't want that. By using the quotes, you tell Bash that you want to preserve each parameter as a separate word.

Another way to deal with the positional parameters is to eliminate each one as it is used. There is a special builtin command named shift which is used for this purpose. When you issue the shift command, the first positional parameter (\$1) goes away. The second one becomes \$1, the third one becomes \$2, and so on down the line. So, if you wish, you can write a loop that keeps using \$1 over and over.

In real scripts, a combination of these techniques is used. A loop to process \$1 as long as it begins with a – takes care of the options. Then, when all the options have been processed and shifted away, everything that's left (in "\$@") is presumably a filename that we want to process.

For brevity, we will not include examples of argument processing here. Instead, we will refer to the FAQ where those examples have already been written.

- Good Practice: Identify where your input comes from before you start writing. If you get to design the data flow into your script, then choose a way that makes sense for the kind of data you're dealing with. If you need to pass filenames, passing them as arguments is an excellent approach, because each one is encapsulated as a word, ready to go.
- In the FAQ: How can I handle command-line arguments (options) to my script easily?<sup>1</sup>

#### 8.2 The Environment

Every program inherits certain information, resources, privileges and restrictions from its parent process. (For a more advanced discussion of this topic, see process management<sup>2</sup>.) One of those resources is a set of variables called *Environment Variables*.

In Bash, environment variables work very much like the regular shell variables we're used to. The only real difference is that they're already set when the script starts up; we don't have to set them ourselves.

Traditionally, environment variables have names that are all capital letters, such as PATH or HOME. This helps you avoid creating any variables that would conflict with them; as long as your variables all contain at least one lower-case letter, you should never have to worry about accidentally colliding with the environment. (Bash's special variables are also capitalized, such as PIPESTATUS. This is done for the exact same reason -- so you can avoid having your variables trampled by Bash.)

Passing information to a program through the environment is useful in many situations. One of those is user preference. Not every user on a Unix-like system has the same likes and dislikes in applications, and in some cases, they may not all speak the same language. So, it's useful for users to be able to tell *every* application they run what their favorite editor is (the EDITOR environment variable), or what language they speak (the various environment variables that compose the user's locale<sup>3</sup>). Environment variables can be set in each user's DotFiles<sup>4</sup>, and then they will be passed automatically to every program the user runs from the login session.

Environment variables can also be tweaked on the fly extremely easily (more easily than if the same information were stored in a file). When you run a command in Bash, you have the option of specifying a temporary environment change which only takes effect for the duration of that command. This is done by putting VAR=value in front of the command. Here is an example:

```
$ ls /tpm
ls: no se puede acceder a /tpm: No existe el fichero o el directorio
$ LANG=C ls /tpm
ls: cannot access /tpm: No such file or directory
```

http://mywiki.wooledge.org/BashFAQ/035

<sup>&</sup>lt;sup>2</sup>http://mywiki.wooledge.org/ProcessManagement#theory

<sup>3</sup>http://mywiki.wooledge.org/locale

<sup>4</sup>http://mywiki.wooledge.org/DotFiles

The LANG=C temporary environment will not cause the user's locale to change for anything other than the one command where it was typed.

In a script, if you know that some information is in an environment variable, you can just use it like any other variable:

```
if [[ $DISPLAY ]]; then
   xterm -e top
else
   top
fi
```

This runs xterm -e top if the environment variable DISPLAY is set (and not empty); otherwise, it runs top.

If you want to put information into the environment for your child processes to inherit, you use the export command:

```
export MYVAR=something
```

The tricky part here is that your environment changes are only inherited by your **descendants**. You can't change the environment of a program that is already running, or of a program that you don't run.

Changing the environment and then running some other program is extremely common. A script that does this as its primary task is called a WrapperScript<sup>1</sup>.

- Good Practice: Don't use all-capital variable names in your scripts, unless they are environment variables. Use lower-case or mixed-case variable names, to avoid accidents.
- In the FAQ: I'm trying to write a script that will change directory (or set a variable), but after the script finishes, I'm back where I started (or my variable isn't set)!<sup>2</sup>

## **8.3** File Descriptors

*File Descriptors* (in short: FDs) are the way programs refer to files, or to other resources that work like files (such as pipes, devices, sockets, or terminals). FDs are kind of like pointers to sources of data, or places data can be written. When something reads from or writes to that FD, the data is read from or written to that FD's resource.

By default, every new process starts with three open FDs:

- Standard Input (stdin): File Descriptor 0
- Standard Output (stdout): File Descriptor 1
- Standard Error (stderr): File Descriptor 2

In an interactive shell, or in a script running on a terminal, the *Standard Input* is how bash sees the characters you type on your keyboard. The *Standard Output* is where the program sends most of its normal information so that the user can see it, and the *Standard Error* is where the program sends its error messages.

<sup>!</sup>http://mywiki.wooledge.org/WrapperScript

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/060

GUI applications also have these FDs, but they don't normally work with them. Usually, they do all their user interaction via the GUI, making it hard for BASH<sup>1</sup> to control them. As a result, we'll stick to simple terminal applications. With those, we can easily feed data to them on their Standard Input, and read data from them on their Standard Output and Standard Error.

Let's make these definitions a little more concrete. Here's a demonstration of how Standard Input and Standard Output work:

```
$ read -p "What is your name? " name; echo "Good day, $name. Would you like some tea?"
What is your name? lhunath
Good day, lhunath. Would you like some tea?
```

read is a command that reads information from stdin and stores it in a variable. We specified name to be that variable. Once read has read a line of information from stdin, it finishes and lets echo display a message. echo sends its output to stdout. stdin and stdout are connected to your terminal. When a program reads from a terminal, it receives keystrokes from your keyboard; when it writes to a terminal, characters are displayed on your monitor. As a result, you can type in your name and are then greeted with a friendly message on your monitor, offering you a cup of tea.

So what is stderr? Let's demonstrate:

```
$ rm secrets
rm: cannot remove `secrets': No such file or directory
```

Unless you have a file called secrets in your current directory, that rm command will fail and show an error message explaining what went wrong. Error messages like these are by convention displayed on stderr.

stderr is also connected to your terminal's output device, just like stdout. As a result, error messages display on your monitor just like the messages on stdout. However, the distinction between stdout and stderr makes it easy to keep errors separated from the application's normal messages. For example, a script might wish to log stderr messages in a special place for long-term storage. Some people also like to use wrappers to make all the output on stderr red, so that they can see the error messages more clearly. (This doesn't work as well as one might wish, but some people find it good enough for some tasks.)

In shell scripts, FDs are always referenced by number. In the next section, we will see some of the ways we can work with FDs using their numbers.

- In the Manual: The read builtin<sup>2</sup>
- Good Practice: Remember that when you create scripts, you should send your custom error messages to the stderr FD. This is a convention and it is very convenient when applications follow the convention. As such, so should you! You're about to learn redirection soon, but let me show you quickly how it's done:

```
echo "Uh oh. Something went really bad.." >&2
```

• *File Descriptor*: A numeric index referring to one of a process's open files. Each command has at least three basic descriptors: FD 0 is stdin, FD 1 is stdout and FD 2 is stderr.

http://mywiki.wooledge.org/BASH

 $<sup>^2</sup>$ http://www.gnu.org/software/bash/manual/bashref.html#index-read-142

#### 8.4 Redirection

The most basic form of input/output manipulation in BASH is *Redirection*. *Redirection* is used to change the data source or destination of a program's FDs. That way, you can send output to a file instead of the terminal, or have an application read from a file instead of from the keyboard.

Redirections are performed by BASH (or any other shell), *before* the shell runs the command to which the redirections are applied.

- In The Manual: Redirections<sup>1</sup>
- Redirection: the practice of changing a FD to read its input from, or send its output to, a different location.

#### 8.4.1 File Redirection

File Redirection involves changing a single FD to point to a file. Let's start with an output redirection:

```
$ echo "It was a dark and stormy night. Too dark to write." > story
$ cat story
It was a dark and stormy night. Too dark to write.
```

The > operator begins an *output redirection*. The redirection applies only to one command (in this case, an echo command). It tells BASH that when BASH runs the command, stdout should point to a file, rather than wherever it was pointing before.

As a result, the echo command will not send its output to the terminal; rather, the > story redirection changes the destination of the stdout FD so that it now points to a file called story. Be aware that this redirection occurs before the echo command is executed. By default, Bash doesn't check to see whether that file story exists first; it just opens the file, and if there was already a file by that name, its former contents are lost. If the file doesn't exist, it is created as an empty file, so that the FD can be pointed to it. This behaviour can be toggled with *Shell Options* (see later).

It should be noted that this redirection is in effect only for the single echo command it was applied to. Other commands executed after that will continue sending their output to the script's stdout location.

We then use the application cat to print out the contents of that file. cat is an application that reads the contents of all the files you pass it as arguments. It then writes each file one after another on stdout. In essence, it concatenates the contents of all the files you pass it as arguments.

Warning: Far too many code examples and shell tutorials on the Internet tell you to use cat whenever you need to read the contents of a file. This is not necessary! cat only serves well to concatenate multiple files together, or as a quick tool on the shell prompt to see what's inside a file. You should NOT use cat to pipe files to commands in your scripts. Instead, you should use a redirection. Please keep this warning in mind. Useless use of cat will result in an extra process to create, and using a pipe instead of a redirection takes away an application's ability to skip back and forth inside the input file.

When we use cat without passing any kind of arguments, it obviously doesn't know what files to read. In this case, cat will just read from stdin instead of from a file (much like read). Since stdin is normally not a regular file, starting cat without any arguments will seem to do nothing:

```
$ cat
```

<sup>1</sup>https://www.gnu.org/savannah-checkouts/gnu/bash/manual/bash.html#Redirections

It doesn't even give you back your shell prompt! What's going on? cat is still reading from stdin, which is your terminal. Anything you type on your keyboard now will be sent to cat as soon as you hit the *Enter* key. With each line you type, cat will do what it normally does: display it reads on stdout, the same way as when it displayed our story on stdout.

```
$ cat
test?
test?
```

Why does it say test? twice now? First of all, terminals are actually more complicated than they appear; they have different modes of operation. The mode we are using in this example is called *canonical mode*, and in this mode, the terminal shows you each character as you type it, and lets you perform extremely simple editing (such as using the Backspace key) on your input. The stuff you type is not actually sent to the application until you press Enter.

As you type test?, you will see it echoed on the screen by the terminal itself. Once you press Enter, the whole line becomes available to the application (cat) that's reading from the terminal. cat reads the line from stdin, and then shows it on stdout, which is also your terminal; hence, the second line: test?.

You can press *Ctrl+D* to send your terminal the *End of File* character. That'll cause cat to think stdin has closed. It will stop reading, and terminate. BASH will see that cat has terminated, and return you to your prompt.

Now let's use an *input redirection* to attach a file to stdin, so that stdin is no longer reading from our keyboard, but instead, now reads from the file:

```
$ cat < story
The story of William Tell.

It was a cold december night. Too cold to write.</pre>
```

The result of this is exactly the same as the result from our previous cat story; except this time, the way it works is a little different. In our first example, cat opened an FD to the file story and read its contents through that FD. In the second example, cat simply reads from stdin, just like it did when it was reading from our keyboard. However, this time, the < story operation has **modified** cat's stdin so that its data source is the file story rather than our keyboard.

Redirection operators can be preceded by a number. That number denotes the FD that will be changed. Let's summarize with some examples:

- command > file: Send the stdout of command to file.
- **command 1> file**: Send the stdout of command to file. Since stdout is FD 1, that's the number we put in front of the redirection operator. This is identical to the previous example, because FD 1 is the default for the > operator.
- command < file: Use the contents of file when command reads from stdin.
- command 0< file: Use the contents of file when command reads from stdin, exactly as in the previous example, since FD 0 (stdin) is the default for the < operator.

The number for the stderr FD is 2. So, let's try sending stderr to a file:

```
$ for homedir in /home/*
> do rm "$homedir/secret"
> done 2> errors
```

In this example, we're looping over each directory (or file) in /home. We then try to delete the file secret in each of them. Some homedirs may not have a secret, or we may not have permission to remove it. As a result, the rm operation will fail and send an error message on stderr.

You may have noticed that our redirection operator isn't on rm, but it's on that done thing. Why is that? Well, this way, the redirection applies to all output to stderr made inside the whole loop. Technically, what happens is BASH opens the file named errors and points stderr to it before the loop begins, then closes it when the loop ends. Any commands run inside the loop (such as rm) *inherit* the open FD from BASH.

Let's see what the result of our loop was:

```
$ cat errors
rm: cannot remove `/home/axxo/secret': No such file or directory
rm: cannot remove `/home/lhunath/secret': No such file or directory
```

Two error messages in our error log file. Two people that didn't have a secret file in their home directory.

If you're writing a script, and you expect that running a certain command may fail on occasion, but don't want the script's user to be bothered by the possible error messages that command may produce, you can silence a FD. Silencing it is as easy as normal *File Redirection*. We're just going to send all output to that FD into the system's black hole:

```
$ for homedir in /home/*
> do rm "$homedir/secret"
> done 2> /dev/null
```

The file /dev/null is **always** empty, no matter what you write to it or read from it. As such, when we write our error messages to it, they just disappear. The /dev/null file remains as empty as ever before. That's because it's not a normal file; it's a *virtual* device. Some people call /dev/null the *bit bucket*.

There is one last thing you should learn about *File Redirection*. It's interesting that you can make error log files like this to keep your error messages; but as I mentioned before, Bash destroys the existing contents of a file when it redirects to it. As a result, each time we run our loop to delete secret files, our log file will be truncated empty before we fill it up again with new error messages. What if we'd like to keep a record of any error messages generated by our loop? What if we don't want that file to be truncated each time we start our loop? The solution is achieved by doubling the redirection operator. > becomes >>. >> will not empty a file; it will just append new data to the end of it!

```
$ for homedir in /home/*
> do rm "$homedir/secret"
> done 2>> errors
```

#### Hooray!

By the way, the space between the redirection operator and the filename is optional. Some people write > file and some write > file. Both ways are correct.

• Good Practice: It's a good idea to use redirection whenever an application needs file data and is built to read data from stdin. A lot of bad examples on the Internet tell you to pipe (see later) the output of cat into processes; but this is nothing more than a very bad idea. When designing an application that could be fed data from a variety of different sources, it is often best simply to have your application read from stdin; that way, the user can use redirection to feed it whatever data she wishes. An application that reads standard input in a generalized way is called a filter.

#### 8.4.2 File Descriptor Manipulation

Now that you know how to manipulate process input and output by sending it to and reading it from files, let's make it a little more interesting still.

It's possible to change the source and destination of FDs to point to or from files, as you know. It's also possible to copy one FD to another. Let's prepare a simple testbed:

```
$ echo "I am a proud sentence." > file
```

We've made a file called file, and written a proud sentence into it.

There's an application called grep that we've seen briefly in a previous chapter. grep is like duct tape: you can use it in almost any project (whether it's a good idea or not). It basically takes a *search pattern* as its first argument and maybe some filenames as extra arguments. Just like cat, grep also uses stdin if you don't specify any files. grep reads the files (or stdin if none were provided) and searches for the *search pattern* you gave it. Most versions of grep even support a -r switch, which makes it take directories as well as files as extra arguments, and then searches all the files and directories in those directories that you gave it. Here's an example of how grep can work:

```
$ ls house/
drawer closet dustbin sofa
$ grep -r socks house/
house/sofa:socks
```

In this silly example we have a directory called house with several pieces of furniture in it as files. If we're looking for our socks in each of those files, we send grep to search the directory house/. grep will search everything in there, open each file and look through its contents. In our example, grep finds socks in the file house/sofa; presumably tucked away under a pillow. You want a more realistic example? Sure:

```
$ grep "$HOSTNAME" /etc/*
/etc/hosts:127.0.0.1 localhost Lyndir
```

Here we instruct grep to search for whatever \$HOSTNAME expands to in whatever files /etc/\* expands to. It finds my hostname, which is Lyndir, in the file /etc/hosts, and shows me the line in that file that contains the *search pattern*.

OK, now that you understand grep, let's continue with our *File Descriptor Manipulation*. Remember that we created a file called file, and wrote a proud sentence to it? Let's use grep to find where that proud sentence is now:

```
$ grep proud *
file:I am a proud sentence.
```

Good! grep found our sentence in file. It writes the result of its operation to stdout which is shown on our terminal. Now let's see if we can make grep send an error message, too:

```
$ grep proud file 'not a file'
file:I am a proud sentence.
grep: not a file: No such file or directory
```

This time, we instruct grep to search for the string proud in the files 'file' and 'not a file'. file exists, and the sentence is in there, so grep happily writes the result to stdout. It moves on to the next file to scan, which is 'not a

file'. grep can't open this file to read its content, because it doesn't exist. As a result, grep emits an error message on stderr which is still connected to our terminal.

Now, how would you go about silencing this grep statement completely? We'd like to send all the output that appears on the terminal to a file instead; let's call it proud.log:

```
# Not quite right....
$ grep proud file 'not a file' > proud.log 2> proud.log
```

Does that look about right? We first use > to send stdout to proud.log, and then use 2> to send stderr to proud.log as well. Almost, but not quite. If you run this command (at least on some computers), and then look in proud.log, you'll see there's only an error message, not the output from stdout. We've created a very bad condition here. We've created two FDs that both point to the same file, independently of each other. The results of this are not well-defined. Depending on how the operating system handles FDs, some information written via one FD may clobber information written through the other FD.

```
$ echo "I am a very proud sentence with a lot of words in it, all for you." > file2
$ grep proud file2 'not a file' > proud.log 2> proud.log
$ cat proud.log
grep: not a file: No such file or directory
of words in it, all for you.
```

What happened here? grep opened file2 first, found what we told it to look for, and then wrote our very proud sentence to stdout (FD 1). FD 1 pointed to proud.log, so the information was written to that file. However, we also had another FD (FD 2) pointed to this same file, and specifically, pointed to the *beginning* of this file. When grep tried to open 'not a file' to read it, it couldn't. Then, it wrote an error message to stderr (FD 2), which was pointing to the beginning of proud.log. As a result, the second write operation overwrote information from the first one!

We need to prevent having two independent FDs working on the same destination or source. We can do this by *duplicating* FDs:

```
$ grep proud file 'not a file' > proud.log 2>&1
```

In order to understand these, you need to remember: always read file redirections from left to right. This is the order in which Bash processes them. First, stdout is changed so that it points to our proud.log. Then, we use the >& syntax to duplicate FD 1 and put this duplicate in FD 2.

A duplicate FD works differently from having two independent FDs pointing to the same place. Write operations that go through either one of them are exactly the same. There won't be a mix-up with one FD pointing to the start of the file while the other has already moved on.

Be careful not to confuse the order:

```
$ grep proud file 'not a file' 2>&1 > proud.log
```

This will duplicate stdout (which points to the terminal) and put the duplicate in stderr, and then stdout will be redirected to proud.log. As a result, stdout's messages will be logged, but the error messages will still go to the terminal. *Oops*.

Note: For convenience, Bash also makes yet another form of redirection available to you. The &> redirection operator is actually just a shorter version of what we did here; redirecting both stdout and stderr to a file:

```
$ grep proud file 'not a file' &> proud.log
```

This is the same as > proud.log 2>&1, but not portable to BourneShell<sup>1</sup>. It is not recommended practice, but you should recognize it if you see it used in someone else's scripts.

TODO: Moving FDs and Opening FDs RW.

- In the FAQ: How can I redirect the output of multiple commands at once?<sup>2</sup>
- How can I redirect the output of 'time' to a variable or file?<sup>3</sup>
- How do I use dialog to get input from the user?<sup>4</sup>
- How can I redirect stderr to a pipe?<sup>5</sup>
- Tell me all about 2>&1 -- what's the difference between 2>&1 >foo and >foo 2>&1, and when do I use which?<sup>6</sup>

#### 8.4.3 Heredocs And Herestrings

Sometimes storing data in a file is overkill. We might only have a tiny bit of it -- enough to fit conveniently in the script itself. Or we might want to redirect the contents of a variable into a command, without having to write it to a file first.

```
$ grep proud <<END
> I am a proud sentence.
> END
I am a proud sentence.
```

This is a *Heredoc* (or *Here Document*). *Heredocs* are useful if you're trying to embed short blocks of multi-line data inside your script. (Embedding larger blocks is bad practice. You should keep your logic (your code) and your input (your data) separated, preferably in different files, unless it's a small data set.)

In a *Heredoc*, we choose a word to act as a sentinel. It can be any word; we used END in this example. Choose one that won't appear in your data set. All the lines that follow the *first* instance of the sentinel, up to the *second* instance, become the stdin for the command. The second instance of the sentinel word has to be a line all by itself.

There are a few different options with *Heredocs*. Normally, you can't indent them -- any spaces you use for indenting your script will appear in the stdin. The terminator string (in our case END) must be at the beginning of the line.

```
echo "Let's test abc:"
if [[ abc = a* ]]; then
    cat <<END
        abc seems to start with an a!
END
fi</pre>
```

Ihttp://mywiki.wooledge.org/BourneShell
2http://mywiki.wooledge.org/BashFAQ/014
3http://mywiki.wooledge.org/BashFAQ/032
4http://mywiki.wooledge.org/BashFAQ/040
5http://mywiki.wooledge.org/BashFAQ/047
6http://mywiki.wooledge.org/BashFAQ/055

#### Will result in:

```
Let's test abc:

abc seems to start with an a!
```

You can avoid this by temporarily removing the indentation for the lines of your *Heredocs*. However, that distorts your pretty and consistent indentation. There is an alternative. If you use <<-END instead of <<END as your *Heredoc* operator, Bash removes any tab characters in the beginning of each line of your *Heredoc* content before sending it to the command. That way you can still use tabs (but not spaces) to indent your *Heredoc* content with the rest of your code. Those tabs will not be sent to the command that receives your *Heredoc*. You can also use tabs to indent your sentinel string.

Bash substitutions are performed on the contents of the *Heredoc* by default. However, if you quote the word that you're using to delimit your *Heredoc*, Bash won't perform any substitutions on the contents. Try this example with and without the quote characters, to see the difference:

```
$ cat <<'XYZ'
> My home directory is $HOME
> XYZ
My home directory is $HOME
```

The most common use of *Heredocs* is dumping documentation to the user:

```
usage() {
    cat <<EOF
usage: foobar [-x] [-v] [-z] [file ...]
A short explanation of the operation goes here.
It might be a few lines long, but shouldn't be excessive.
EOF
}</pre>
```

Now let's check out the very similar but more compact *Herestring*:

```
$ grep proud <<<"I am a proud sentence"
I am a proud sentence.</pre>
```

This time, stdin reads its information straight from the string you put after the <<< operator. This is very convenient to send data that's in variables into processes:

```
$ grep proud <<<"$USER sits proudly on his throne in $HOSTNAME."
lhunath sits proudly on his throne in Lyndir.</pre>
```

*Herestring*s are shorter, less intrusive and overall more convenient than their bulky *Heredoc* counterpart. However, they are not portable to the Bourne shell.

Later on, you will learn about pipes and how they can be used to send the output of a command into another command's stdin. Many people use pipes to send the output of a variable as stdin into a command. However, for this purpose, *Herestrings* should be preferred. They do not create a subshell and are lighter both to the shell and to the style of your shell script:

```
$ echo 'Wrap this silly sentence.' | fmt -t -w 20
Wrap this silly
```

```
sentence.
$ fmt -t -w 20 <<< 'Wrap this silly sentence.'
Wrap this silly
sentence.</pre>
```

Technically, *Heredocs* and *Herestrings* are themselves redirects just like any other. As such, additional redirections can occur on the same line, all evaluated in the usual order.

```
$ cat <<EOF > file
> My home dir is $HOME
> EOF
$ cat file
My home dir is /home/greg
```

• Good Practice: Long heredocs are usually a bad idea because scripts should contain logic, not data. If you have a large document that your script needs, you should ship it in a separate file along with your script. Herestrings, however, come in handy quite often, especially for sending variable content (rather than files) to filters like grep or sed.

# 8.5 Pipes

Now that you can effortlessly manipulate *File Descriptors* to direct certain types of output to certain files, it's time you learn some more ingenious tricks available through I/O redirection.

You can use *File Redirection* to write output to files or read input from files. But what if you want to connect the output of one application directly to the input of another? That way, you could build a sort of chain to process output. If you already know about FIFOs, you could use something like this to that end:

```
$ ls
$ mkfifo myfifo; ls
myfifo
$ grep bea myfifo &
[1] 32635
$ echo "rat
> cow
> deer
> bear
> snake" > myfifo
bear
```

We use the mkfifo command to create a new file in the current directory named 'myfifo'. This is no ordinary file, however, but a FIFO (which stands for *First In*, *First Out*). FIFOs are special files that serve data on a First In, First Outbasis. When you read from a FIFO, you will only receive data as soon as another process writes to it. As such, a FIFO never really contains any data. So long as no process writes to it, any read operation on the FIFO will **block** as it waits for data to become available. The same works for writes to the FIFO -- they will block until another process reads from the FIFO.

In our example, the FIFO called myfifo is read from by grep. grep waits for data to become available on the FIFO. That's why we append the grep command with the & operator, which puts it in the background. That way, we can continue typing and executing commands while grep runs and waits for data. Our echo statement feeds data to the FIFO. As soon as this data becomes available, the running grep command reads it in and processes it. The result is displayed. We have successfully sent data from the echo command to the grep command.

But these temporary files are a real annoyance. You may not have write permissions. You need to remember to clean up any temporary files you create. You need to make sure that data is going in and out, or the FIFO might just end up blocking for no reason.

For these reasons, another feature is made available: *Pipes*. A pipe basically just connects the stdout of one process to the stdin of another, effectively *piping* the data from one process into another. The entire set of commands that are piped together is called a *pipeline*. Let's try our above example again, but using pipes:

```
$ echo "rat
> cow
> deer
> bear
> snake" | grep bea
bear
```

The pipe is created using the | operator between two commands that are connected with the pipe. The former command's stdout is connected to the latter command's stdin. As a result, grep can read echo's output and display the result of its operation, which is bear.

Pipes are widely used as a means of post-processing application output. FIFOs are, in fact, also referred to as named pipes. They accomplish the same results as the pipe operator, but through a filename.

Note: The pipe operator creates a subshell environment for each command. This is important to know because any variables that you modify or initialize inside the second command will appear unmodified outside of it. Let's illustrate:

```
$ message=Test
$ echo 'Salut, le monde!' | read message
$ echo "The message is: $message"
The message is: Test
$ echo 'Salut, le monde!' | { read message; echo "The message is: $message"; }
The message is: Salut, le monde!
$ echo "The message is: $message"
The message is: Test
```

Once the pipeline ends, so do the subshells that were created for it. Along with those subshells, any modifications made in them are lost. So be careful!

• Good Practice: Pipes are a very attractive means of post-processing application output. You should, however, be careful not to over-use pipes. If you end up making a pipeline that consists of three or more applications, it is time to ask yourself whether you're doing things a smart way. You might be able to use more application features of one of the post-processing applications you've used earlier in the pipe. Each new command in a pipeline causes a new subshell and a new application to be loaded. It also makes it very hard to follow the logic in your script!

- In The Manual: Pipelines<sup>1</sup>
- In the FAQ: I set variables in a loop. Why do they suddenly disappear after the loop terminates? Or, why can't I pipe data to read?<sup>2</sup>
- How can two processes communicate using named pipes (fifos)?<sup>3</sup>
- How can I redirect stderr to a pipe?<sup>4</sup>
- How can I read a file line-by-line?<sup>5</sup>
- Tell me all about 2>&1 -- what's the difference between 2>&1 >foo and >foo 2>&1, and when do I use which?<sup>6</sup>

# 8.6 Miscellaneous Operators

Aside from the standard I/O operators, bash also provides a few more advanced operators that make life on the shell that much nicer.

#### 8.6.1 Process Substitution

A cousin of the pipe is the *Process Substitution* operator, which comes in two forms: < (cmd) and > (cmd). It's a convenient way to get the benefits of temporary files or named pipes without having to create them yourself. Whenever you think you need a temporary file to do something, process substitution might be a better way to handle it.

The > () form is relatively rare, and we won't cover it here, because it'll just confuse things. Once you understand how < () works, > () just does the same thing in reverse (writing instead of reading).

What < () does, is run the command inside the parentheses, and gives you a temporary filename that you can use to read the command's output. The advantage over pipes is that you can use the filename as an argument to a command that expects to see a filename.

For example, let's say you have two files that you'd like to diff, but they aren't sorted yet. You could generate two temporary files, to hold the sorted versions of your original files, and then run diff on those. Or, you could use process substitutions:

```
$ diff <(sort file1) <(sort file2)</pre>
```

With the < () operator, the command's output is sent through a named pipe (or something similar) that's created by bash. The operator itself in your command is replaced by the filename of that file. After your whole command finishes, the file is cleaned up.

The same thing can be done with any commands whose output you'd like to pass to diff. Imagine you want to see the difference between the output of two commands. Ordinarily, you'd have to put the two outputs in two files and diff those:

```
http://www.gnu.org/software/bash/manual/bashref.html#Pipelines
```

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/024

<sup>&</sup>lt;sup>3</sup>http://mywiki.wooledge.org/BashFAQ/027

<sup>4</sup>http://mywiki.wooledge.org/BashFAQ/047

<sup>5</sup>http://mywiki.wooledge.org/BashFAQ/001

 $<sup>^{6}</sup>$ http://mywiki.wooledge.org/BashFAQ/055

Using the process substitution operator, we can do all that with a one-liner and no need for manual cleanup:

The < (...) part is replaced by the temporary FIFO created by bash, so diff actually sees something like this:

```
$ diff -y /dev/fd/63 /dev/fd/62
```

Here we see how bash runs diff when we use process substitution. It runs our head and tail commands, sending their respective outputs through the "files" /dev/fd/63 and /dev/fd/62. Then it runs the diff command, passing those filenames where originally we had put the process substitution operators.

The actual implementation differs from system to system. In fact, you can see what the above would actually look like to diff on your box by putting an echo in front of our command:

```
box1$ echo diff -y <(head -n 1 .dictionary) <(tail -n 1 .dictionary)
diff -y /dev/fd/63 /dev/fd/62

box2$ echo <(cat /dev/null)
/var/tmp//sh-np-605454726</pre>
```

For examples using the > () form, see ProcessSubstitution<sup>1</sup> and Bash FAQ 106<sup>2</sup>.

• Good Practice: Process Substitution gives you a concise way to create temporary FIFOs automatically. They're less flexible than creating your own named pipes by hand, but they're perfect for common short commands like diff that need filenames for their input sources.

http://mywiki.wooledge.org/ProcessSubstitution

<sup>2</sup>http://mywiki.wooledge.org/BashFAQ/106

# 9 Compound Commands

BASH<sup>1</sup> offers numerous ways to combine simple commands to achieve our goals. We've already seen some of them in practice, but now let's look at a few more.

Bash has constructs called *compound commands*, which is a catch-all phrase covering several different concepts. We've already seen some of the compound commands Bash has to offer -- if statements, for loops, while loops, the [[keyword, case and select. We won't repeat that information again here. Instead, we'll explore the other compound commands we haven't seen yet: subshells, command grouping, and arithmetic evaluation.

In addition, we'll look at functions and aliases, which aren't compound commands, but which work in a similar way.

• In the manual: Compound Commands<sup>2</sup>

### 9.1 Subshells

A SubShell<sup>3</sup> is similar to a child process, except that more information is inherited. Subshells are created implicitly for each command in a pipeline. They are also created explicitly by using parentheses around a command:

```
$ (cd /tmp || exit 1; date > timestamp)
$ pwd
/home/lhunath
```

When the subshell terminates, the cd command's effect is gone -- we're back where we started. Likewise, any variables that are set during the subshell are not remembered. You can think of subshells as temporary shells. See SubShell<sup>4</sup> for more details.

Note that if the cd failed in that example, the exit 1 would have terminated the subshell, but *not* our interactive shell. As you can guess, this is quite useful in real scripts.

• In the manual: Command Grouping<sup>5</sup>

# 9.2 Command grouping

We've already touched on this subject in Grouping Statements<sup>6</sup>, though it pays to repeat it in the context of this chapter.

Commands may be grouped together using curly braces. Command groups allow a collection of commands to be considered as a whole with regards to redirection and control flow. All compound commands such as if statements and while loops do this as well, but command groups do *only* this. In that sense, command groups can be thought of as "null compound commands" in that they have no effect other than to group commands. They look a bit like subshells, with the difference being

<sup>1</sup>http://mywiki.wooledge.org/BASH

 $<sup>^2 \</sup>verb|http://www.gnu.org/s/bash/manual/bash.html#Compound-Commands|$ 

<sup>&</sup>lt;sup>3</sup>http://mywiki.wooledge.org/SubShell

<sup>4</sup>http://mywiki.wooledge.org/SubShell

<sup>5</sup>http://www.gnu.org/s/bash/manual/bash.html#Command-Grouping

 $<sup>^{6}</sup>$ http://mywiki.wooledge.org/BashGuide/TestsAndConditionals#Grouping\_Statements

that command groups are executed in the same shell as everything else, rather than a new one. This is both faster and allows things like variable assignments to be visible outside of the command group.

All commands within a command group are within the scope of any redirections applied to a command group (or any compound command):

```
$ { echo "Starting at $(date)"; rsync -av . /backup; echo "Finishing at $(date)"; } >backup.log 2>&1
```

The above example truncates and opens the file backup.log on stdout, then points stderr at where stdout is currently pointing (backup.log), then runs each command with those redirections applied. The file descriptors remain open until all commands within the command group complete before they are automatically closed. This means backup.log is only opened a single time, not opened and closed for each command. The next example demonstrates this better:

```
$ echo "cat
> mouse
> dog" > inputfile
$ for var in {a..c}; do read -r "$var"; done < inputfile
$ echo "$b"
mouse</pre>
```

Notice how we didn't actually use a command group here. As previously explained, "for" being a compound command behaves just like a command group. It would have been extremely difficult to read the second line from a file without allowing for multiple read commands to read from a single open FD without rewinding to the start each time. Contrast with this:

```
$ read -r a < inputfile
$ read -r b < inputfile
$ read -r c < inputfile
$ echo "$b"
cat</pre>
```

That's not what we wanted at all!

Command groups are also useful to shorten certain common tasks:

```
$ [[ -f $CONFIGFILE ]] || { echo "Config file $CONFIGFILE not found" >&2; exit 1; }
```

The logical "or" now executes the command group if \$CONFIGFILE doesn't exist rather than just the first simple command. A subshell would not have worked here, because the exit 1 in a command group terminates the entire shell -- which is what we want here.

Compare that with a differently formatted version:

```
$ if [[ ! -f $CONFIGFILE ]]; then
> echo "Config file $CONFIGFILE not found" >&2
> exit 1
> fi
```

If the command group is on a single line, as we've shown here, then there *must* be a semicolon before the closing }, ( i.e { ...; last command; } ) otherwise, Bash would think } is an argument to the final command in the group. If the command group is spread across multiple lines, then the semicolon may be replaced by a newline:

```
$ {
> echo "Starting at $(date)"
> rsync -av . /backup
> echo "Finishing at $(date)"
> } > backup.log 2>&1
```

If redirections are used on a simple command, they only apply to the command itself, not parameter or other expansions. Command groups make all contents including expansions apply even in the case of a single simple command:

```
$ { echo "$(cat)"; } <<<'hi'
hi
$ { "$(</dev/stdin)" <<<"$_"; } <<<'cat'
hi</pre>
```

The second command (which you don't need to fully understand) would require killing the shell if the command group weren't present, since the shell would be reading from the tty to determine the command to execute. It also illustrates that the command still gets the redirect applied to it, while the expansion gets that of the command group.

• In the manual: Command Grouping<sup>1</sup>

### 9.3 Arithmetic Evaluation

Bash has several different ways to say we want to do arithmetic instead of string operations. Let's look at them one by one. The first way is the let command:

```
$ unset a; a=4+5
$ echo $a
4+5
$ let a=4+5
$ echo $a
9
```

You may use spaces, parentheses and so forth, if you quote the expression:

```
$ let a='(5+2)*3'
```

For a full list of operators availabile, see help let or the manual.

Next, the actual arithmetic evaluation compound command syntax:

```
$ ((a=(5+2) *3))
```

This is equivalent to let, but we can also use it as a *command*, for example in an if statement:

```
$ if (($a == 21)); then echo 'Blackjack!'; fi
```

 $<sup>^{\</sup>rm I}{\rm http://www.gnu.org/s/bash/manual/bash.html\#Command-Grouping}$ 

Operators such as ==, <, > and so on cause a comparison to be performed, inside an arithmetic evaluation. If the comparison is "true" (for example, 10 > 2 is true in arithmetic -- but not in strings!) then the compound command exits with status 0. If the comparison is false, it exits with status 1. This makes it suitable for testing things in a script.

Although not a compound command, an arithmetic substitution (or arithmetic expression) syntax is also available:

```
$ echo "There are $(($rows * $columns)) cells"
```

Inside \$((...)) is an *arithmetic context*, just like with ((...)), meaning we do arithmetic (multiplying things) instead of string manipulations (concatenating \$rows, space, asterisk, space, \$columns). \$((...)) is also portable to the POSIX shell, while ((...)) is not.

Readers who are familiar with the C programming language might wish to know that ((...)) has many C-like features. Among them are the ternary operator:

```
((abs = (a >= 0) ? a : -a))
```

and the use of an integer value as a truth value:

```
$ if ((flag)); then echo "uh oh, our flag is up"; fi
```

Note that we used variables inside ((...)) without prefixing them with \$-signs. This is a special syntactic shortcut that Bash allows inside arithmetic evaluations and arithmetic expressions.

There is one final thing we must mention about ((flag)). Because the inside of ((...)) is C-like, a variable (or expression) that evaluates to zero will be considered false for the purposes of the arithmetic evaluation. Then, because the evaluation is false, it will exit with a status of 1. Likewise, if the expression inside ((...)) is non-zero, it will be considered true; and since the evaluation is true, it will exit with status 0. This is potentially very confusing, even to experts, so you should take some time to think about this. Nevertheless, when things are used the way they're intended, it makes sense in the end:

```
$ flag=0  # no error
$ while read line; do
> if [[ $line = *err* ]]; then flag=1; fi
> done < inputfile
$ if ((flag)); then echo "oh no"; fi</pre>
```

• In the manual: Arithmetic Expansion<sup>1</sup>, Shell Arithmetic<sup>2</sup>

### 9.4 Functions

Functions are very nifty inside Bash scripts. They are blocks of commands, much like normal scripts you might write, except they don't reside in separate files, and they don't cause a separate process to be executed. However, they take arguments just like scripts -- and unlike scripts, they can affect variables inside your script, if you want them to. Take this for example:

<sup>1</sup>http://www.gnu.org/s/bash/manual/bash.html#Arithmetic-Expansion

<sup>2</sup>http://www.gnu.org/s/bash/manual/bash.html#Shell-Arithmetic

```
$ sum() {
> echo "$1 + $2 = $(($1 + $2))"
> }
```

This will do absolutely nothing when run. This is because it has only been stored in memory, much like a variable, but it has not yet been called. To run the function, you would do this:

```
$ sum 1 4
1 + 4 = 5
```

Amazing! We now have a basic calculator, and potentially a more economic replacement for a five year-old.

A note on scope: if you choose to embed functions within script files, as many will find more convenient, then you need to understand that the parameters you pass to the script are not necessarily the parameters that are passed to the function. To wrap this function inside a script, we would write a file containing this:

```
#!/bin/bash
sum() {
    echo "$1 + $2 = $(($1 + $2))"
}
sum "$1" "$2"
```

As you can see, we passed the script's two parameters to the function within, but we could have passed anything we wanted (though, doing so in this situation would only confuse users trying to use the script).

Functions serve a few purposes in a script. The first is to isolate a block of code that performs a specific task, so that it doesn't clutter up other code. This helps you make things more readable, when done in moderation. (Having to jump all over a script to track down 7 functions to figure out what a single command does has the opposite effect, so make sure you do things that make sense.) The second is to allow a block of code to be reused with slightly different arguments.

Here's a slightly less silly example:

Here, we define a *function* named open. This function is a block of code that takes a single argument, and based on the *pattern* of that argument, it will either run xmms, display or mplayer with that argument. Then, a for loop iterates over all of the *script's* positional parameters. (Remember, for file is equivalent to for file in "\$@" and both of them iterate over the full set of positional parameters.) The for loop calls the open function for each parameter.

As you may have observed, the function's parameters are different from the script's parameters.

Functions may also have *local variables*, declared with the local or declare keywords. This lets you do work without potentially overwriting important variables from the caller's namespace. For example,

```
#!/bin/bash
count() {
    local i
    for ((i=1; i<=$1; i++)); do echo $i; done
    echo 'Ah, ah, ah!'
6 }
for ((i=1; i<=3; i++)); do count $i; done</pre>
```

The local variable i inside the function is stored differently from the variable i in the outer script. This allows the two loops to operate without interfering with each other's counters.

Functions may also call themselves recursively, but we won't show that today. Maybe later!

• In the manual: Shell Functions<sup>1</sup>

### 9.5 Aliases

Aliases are superficially similar to functions at first glance, but upon closer examination, they have entirely different behavior.

- Aliases do not work in scripts, at all. They only work in interactive shells.
- Aliases cannot take arguments.
- Aliases will not invoke themselves recursively.
- Aliases cannot have local variables.

Aliases are essentially keyboard shortcuts intended to be used in .bashrc files to make your life easier. They usually look like this:

```
$ alias ls='ls --color=auto'
```

Bash checks the first word of every simple command to see whether it's an *alias*, and if so, it does a simple text replacement. Thus, if you type

```
$ ls /tmp
```

Bash acts as though you had typed

```
$ ls --color=auto /tmp
```

If you wanted to duplicate this functionality with a function, it would look like this:

```
$ unalias ls
$ ls() { command ls --color=auto "$@"; }
```

 $<sup>^{1} \</sup>verb|http://www.gnu.org/s/bash/manual/bash.html # Shell-Functions|$ 

As with a *command group*, we need a ; before the closing } of a function if we write it all in one line. The special built-in command command tells our function **not** to call itself recursively; instead, we want it to call the ls command that it would have called if there hadn't been a function by that name.

Aliases are useful as long as you don't try to make them work like functions. If you need complex behavior, use a function instead.

# 9.6 Destroying Constructs

To remove a function or variable from your current shell environment use the *unset* command. It is usually a good idea to be explicit about whether a function or variable is to be unset using the -f or -v flags respectively. If unspecified, variables take precedence over functions.

```
$ unset -f myfunction
$ unset -v 'myArray[2]' # unset element 2 of myArray. The quoting is important to prevent globbing.
```

To remove an alias, use the unalias command.

```
$ unalias rm
```

- In the manual: Bourne Shell Builtins<sup>1</sup>
- In the FAQ: ...

<sup>1</sup>http://www.gnu.org/s/bash/manual/bash.html#Bourne-Shell-Builtins

# 10 Sourcing

When you call one script from another, the new script inherits the environment of the original script, just like running any other program in UNIX. Explaining what this means is out of the scope of this guide, but for the most part you can consider the environment to be the current working directory, open file descriptors, and environment variables, which you can view using the export command.

When the script that you ran (or any other program, for that matter) finishes executing, its environment is discarded. The environment of the first script will be same as it was before the second script was called, although of course some of bash's special parameters may have changed (such as the value of \$?, the return value of the most recent command). This means, for example, you can't simply run a script to change your current working directory for you.

#### • In the FAO:

I'm trying to write a script that will change directory (or set a variable), but after the script finishes, I'm back where I started (or my variable isn't set)!<sup>1</sup>

What you can do is to source the script, instead of running it as a child. You can do this using the . (dot) command:

```
. ./myscript  #runs the commands from the file myscript in this environment
```

This is often called *dotting in* a script. The . tells BASH<sup>2</sup> to read the commands in ./myscript and run them in the current shell environment. Since the commands are run in the current shell, they can change the current shell's variables, working directory, open file descriptors, functions, etc.

Note that Bash has a second name for this command, source, but since this works identically to the . command, it's probably easier to just forget about it and use the . command, as that will work everywhere.

http://mywiki.wooledge.org/BashFAQ/060

<sup>&</sup>lt;sup>2</sup>http://mywiki.wooledge.org/BASH

# 11 Job Control

Though not typically used in scripts, job control is very important in interactive shells. Job control allows you to interact with background jobs, suspend foreground jobs, and so on.

# 11.1 Theory

On Posix systems, jobs are implemented as "process groups", with one process being the leader of the group. Each tty (terminal) has a single "foreground process group" that is allowed to interact with the terminal. All other process groups with the same controlling tty are considered background jobs, and can be either running or suspended.

A job is suspended when its process group leader receives one of the signals SIGSTOP, SIGTSTP, SIGTTIN, or SIGTTOU. SIGTTIN (and SIGTTOU, if the user issued stty tostop) are automatically sent whenever a background job tries to read from or write to the terminal---this is why cat & is immediately suspended rather than running in the background.

Certain keypresses at the terminal cause signals to be sent to all processes in the foreground process group. These can be configured with stty, but are usually set to the defaults:

- Ctrl-Z sends SIGTSTP to the foreground job (usually suspending it)
- Ctrl-C sends SIGINT to the foreground job (usually terminating it)
- Ctrl-\ sends SIGQUIT to the foreground job (usually causing it to dump core and abort)

#### 11.2 Practice

Job control is on by default in interactive shells. It can be turned on for scripts with set -m or set -o monitor.

A foreground job can be suspended by pressing Ctrl-Z. There is no way to refer to the foreground job in bash: if there is a foreground job other than bash, then bash is waiting for that job to terminate, and hence cannot execute any code (even traps are delayed until the foreground job terminates). The following commands, therefore, work on background (and suspended) jobs only.

Job control enables the following commands:

- **fg** [jobspec]: bring a background job to the foreground.
- **bg** [jobspec ...]: run a suspended job in the background.
- **suspend**: suspend the shell (mostly useful when the parent process is a shell with job control).

Other commands for interacting with jobs include:

• **jobs** [*options*] [*jobspec* ...]: list suspended and background jobs. Options include -p (list process IDs only), -s (list only suspended jobs), and -r (list only running background jobs). If one or more jobspecs are specified, all other jobs are ignored.

- kill can take a jobspec instead of a process ID.
- disown tells bash to forget about an existing job. This keeps bash from automatically sending SIGHUP to the processes in that job, but also means it can no longer be referred to by jobspec.

So, what does all that mean? Job control allows you to have multiple things going on within a single terminal session. (This was terribly important in the old days when you only had one terminal on your desk, and no way to create virtual terminals to add more.) On modern systems and hardware, you have more choices available -- you could for example run screen or tmux within a terminal to give you virtual terminals. Or within an X session, you could open more xterm or similar terminal emulators (and you can mix the two together as well).

But sometimes, a simple job control "suspend and background" comes in handy. Maybe you started a backup and it's taking longer than you expected. You can suspend it with Ctrl-Z and then put it in the background with bg, and get your shell prompt back, so that you can work on something else in the same terminal session while that backup is running.

# 11.3 Job Specifications

A job specification or "jobspec" is a way of referring to the processes that make up a job. A jobspec may be:

- n to refer to job number n.
- %str to refer to a job which was started by a command beginning with str. It is an error if there is more than one such job.
- %?str to refer to a job which was started by a command containing str. It is an error if there is more than one such job.
- %% or %+ to refer to the current job: the one most recently started in the background, or suspended from the foreground. fg and bg will operate on this job if no **jobspec** is given.
- %- for the previous job (the job that was %% before the current one).

It is possible to run an arbitrary command with a jobspec, using jobs -x ''cmd args...''. This replaces arguments that look like jobspecs with the PIDs of the corresponding process group leaders, then runs the command. For example, jobs -x strace -p %% will attach strace to the current job (most useful if it is running in the background rather than suspended).

Finally, a bare jobspec may be used as a command: %1 is equivalent to fg %1, while %1 & is equivalent to bg %1.

#### 11.4 See Also

ProcessManagement<sup>1</sup> discusses practices for working with multiple processes. There's also an example of the use of job control within a script.

http://mywiki.wooledge.org/ProcessManagement

# 12 Practices

#### 12.1 Choose Your Shell

The first thing you should do before starting a shell script, or any kind of script or program for that matter, is enumerate the requirements and the goal of that script. Then evaluate what the best tool is to accomplish those goals.

Keep in mind that BASH<sup>1</sup> may not always be the best tool for the job.

If you need complex text processing, awk, perl, ruby are generally well suited for the task, though the latter is rarely installed by default.

All of them can be given the code to interpret on the command line (awk -- awk-script-code arguments for the inline script, perl/ruby -e perl/ruby-script-code -- arguments for the inline script). As many of the characters in the syntax of those languages are also special in the syntax of the shell, you'll want to put the code argument inside single quotes. It's perfectly fine (and recommended for legibility) to write that code on several lines. You may also prefer to write separate perl/ruby scripts, but note that for awk, the script interface is broken by design as with a script starting with #! /usr/bin/awk -f, any argument passed to the awk script is considered for options for the awk interpreter first (not to mention the fact that foo=bar arguments are interpreted as variable assignments). That's fixed by the GNU implementation of awk, where you can use a #! /usr/bin/gawk -E shebang instead.

If you need to retrieve data from an HTML or XML file in a reliable manner, consider XPath/XSLT, or a language that has a library available for parsing XML or HTML.

If you decide to use a shell script, evaluate the following questions:

- In the foreseeable future, might your script be needed in an environment where Bash is not available by default?
  - If so, then consider **limiting the script to POSIX sh syntax only.** POSIX defines a specification for a minimum standard language for sh and a few basic utilities and virtually all systems supply a shell called sh (whether it is or is derived from Bash or the Bourne, Almquist, Forsyth, Korn, Yash or Zsh shells) and the most common of those utilities (sometime built-in to their sh) that try to conform to that standard. It you write your script to that standard, you can reasonably expect that script to run on any POSIX system. However, you will have to balance the need for portability against the cost of doing away with the more advanced features from shells such as Bash that you'd be unable to use.
  - Keep in mind that this guide does not apply to sh See Bashism<sup>2</sup> page gives some guidance how to convert Bash constructs into standard sh, but it is not complete.
- Are you considering supporting multiple operating systems to run the script? Then, not all the features found in the latest Bash will be available.
  - For example in macOS, the default Bash version is permanently 3.2 due to GPL 2.0
  - Some systems have 10 or 20 years supported lifespan and would keep the same version of bash during that span. For example, Solaris 10 was released in 2005 when the current version of bash was 3.2 and will keep supplying that version (in an optional package) until it's end of life in 2027.

<sup>!</sup>http://mywiki.wooledge.org/BASH

<sup>2</sup>http://mywiki.wooledge.org/Bashism

If the above questions do not limit your choices, use all the Bash features you require, and then make note of which version of Bash is required to run your script.

Using Bash means you can use arrays and avoid some awkward sh scripting techniques.

- Examine scripting examples on the Web carefully before giving in to the temptation to use their code or other constructs. Some may be broken or may not work universally.
- Remember to use the correct shebang. For Bash script, write #!/usr/bin/env bash at the top of your script. Omitting this or using the #!/bin/sh header is **incorrect**. The sh limits the script to run under the POSIX shell features even if the operating system's /bin/sh is a symlink to Bash, like is the case RedHat<sup>1</sup> Enterprise Linux or a special build of bash like on macos.
- In Bash scripts, consider to use [[...]] construct over the [command unless you're doing decimal integer comparisons. Bash's [[...]] is more forgiving in many ways, like safeguarding against *Word Splitting* or *globbing* when you forget to quote variables. It's specific micro-syntax also supports glob pattern and regular expression matching capabilities.
- Skip over the over-30-years-old obsolete command substitution `...` syntax. The modern POSIX version \$ (...) is more consistent with the syntax of *Expansion*, supports multiple quoting, and is easier to nest. An example: fullpath=\$ (CDPATH= cd -P -- "\$ (readlink -- "\$dir") " && pwd)
- First and foremost, remember to "use more quotes". This will protect the strings and parameter expansions from WordSplitting<sup>2</sup> and globbing<sup>3</sup>. Word splitting and the filename generation aka globbing that is are done upon unquoted expansions may cause major problems if things are not quoted properly. A problem example: file='file \* name.txt'; rm \$file
- Learn to use parameter expansions instead of sed or cut to manipulate simple strings in Bash. If you want to remove the extension from a filename, use \${filename%.\*} instead of `echo "\$filename" | sed 's/\.[^.]\*\$//'` or some other dinosaur.
- Use built-in POSIX math operations instead of calling external expr to do simple calculations. If the some script uses x=`expr \$x+1\$` then it's *not* something you want to mimic. The faster and portable POSIX syntax is: x=\$ ( (x+1))

# 12.2 Quoting

While in most other languages, quotes are used to introduce string or character literals, most things that shells deal with are strings and quotes are only used to remove the special meaning of characters that are special in the syntax of the shell so they be treated as parts of strings like regular characters.

```
http://mywiki.wooledge.org/RedHat
```

<sup>&</sup>lt;sup>2</sup>http://mywiki.wooledge.org/WordSplitting

<sup>&</sup>lt;sup>3</sup>http://mywiki.wooledge.org/globbing

<sup>&</sup>lt;sup>4</sup>http://mywiki.wooledge.org/BashGuide/Parameters#Parameter\_Expansion

For instance, space is used to separate command line arguments and quoting it whether with singe, double quotes or preceding it with a backslash make it a regular character that can be included inside a command argument for instance. Same goes for ;, newline, &, |... which otherwise separate commands or <, > which introduce redirections.

But quoting with double-quotes is also used to work around a misfeature introduced by the Bourne shell (though to be fair as an attempt to provide some level of backward compatibility with its predecessor the Thompson shell which did even worse).

That *Word Splitting* feature is the demon inside Bash that is out to get unsuspecting newcomers or even veterans who let down their guard.

If you do not understand how *Word Splitting* or *globbing* work or when it is applied you should be very careful with your strings and your *Parameter Expansions*. I suggest you read up on WordSplitting<sup>1</sup> if you doubt your knowledge.

As mentioned above, quotes can be used to remove the special meaning of that space character. Compare:

If the first command, echo received 6 arguments, which it joined with spaces; in the second and third, it received only one argument which included literal space characters as all but the space character that separated echo from its argument where quoted, all at once with '...' or each individually with backslash.

Word Splitting is something unrelated and applies on *expansions* only (parameter expansion, arithmetic expansion and command substitution) and is not about whitespace but about characters of the \$IFS special parameter (which so happens to contain 3 whitespace characters by default).

That (and the fact that expansions are subject to globbing) is the one that is pernicious as it is hidden and easy to overlook. As a result, if you missed a cup of coffee this morning you may find yourself making this mistake:

```
$ sentence='Push that word away from me.'
$ echo $sentence
Push that word away from me.
$ echo "$sentence"
Push that word away from me.
```

#### Or worse:

As you can see, in the first echo command, we neglected to add quotes. Bash expanded our sentence and then used *Word Splitting* (here assuming \$IFS was not modified from its default) to split the resulting expansion up into arguments to use for

http://mywiki.wooledge.org/WordSplitting

echo, which had the result of destroying our carefully thought-out formatting. In our second example, the quotes around the *Parameter Expansion* of \$sentence make sure Bash **does not** split it up into multiple arguments around the characters of \$IFS.

The \*worse\* section shows it's not just spaces you need to worry about. Word Splitting occurs on all characters in \$IFS (space, tab and newline by default), and  $\star$ , ?, [...], \ in some versions of bash (and more if the extglob option is enabled) trigger globbing. Here's another example to show you just how badly you can break things if you neglect to quote when necessary:

In some very rare occasions it may be desired to leave out the quotes. That's if you **need** Word Splitting to take place:

```
$ friends="Marcus JJ Thomas Michelangelo"
$ IFS=$' \t\n' # make sure IFS is set to the value you want in case it has been modified before
$ set -o noglob # needed if you can't guarantee the text will contain glob characters
$ for friend in $friends
> do echo "$friend is my friend!"; done
Marcus is my friend!
JJ is my friend!
Thomas is my friend!
Michelangelo is my friend!
```

But, honestly? You should use arrays for nearly all of these cases. Arrays have the benefit that they separate strings without the need for an explicit delimiter. That means your strings in the array can contain **any** valid (non-NUL) character, without the worry that it might be your string delimiter (like the space is in our example above). Using arrays in our example above gives us the ability to add middle or last names of our friends:

```
$ friends=( 'Marcus The Rich' 'JJ The Short' 'Timid Thomas' 'Michelangelo The Mobster' )
$ for friend in "${friends[@]}"
> do echo "$friend is my friend"; done
```

Note that in our previous for we used an *unquoted* \$friends. This allowed Bash to split our friends string up into words. In this last example, we quoted the \${friends[@]} Parameter Expansion. Quoting an expansion of an array through the @ index makes Bash expand that array into a sequence of its elements, where each element is wrapped in quotes.

# 12.3 Readability

Almost as important as the result of your code is the **readability of your code**.

Chances are that you aren't just going to write a script once and then forget about it. If so, you might as well delete it after using it. If you plan to continue using it, you should also plan to continue maintaining it. Unlike your house your code won't get dirty over time, but you will learn new techniques and new approaches constantly. You will also gain insight about how your script is used. All that new information you gather since the completion of your initial code should be used to maintain your code in such a way that it constantly improves. Your code should keep growing more user-friendly and more stable.

• Trust me when I say, no piece of code is ever 100% finished, with the exception of some very short and useless chunks of nothingness.

To make it easier for yourself to keep your code healthy and improve it regularly you should keep an eye on the readability of what you write. When you return to a long loop after a year has passed since your last visit to it and you wish to improve it, add a feature, or debug something about it, consider whether you'd rather see this:

```
friends=(
       'Marcus The Rich'
2
        'JJ The Short'
       'Timid Thomas'
4
       'Michelangelo The Mobster'
7
   # Say something significant about my friends.
   for name in "${friends[@]}"; do
10
       # Mv first friend (in the list).
11
       if [[ $name = "${friends[0]}" ]]; then
12
           echo "$name was my first friend."
13
14
       # My friends whose names start with M.
15
       elif [[ $name = M* ]]; then
16
           echo "$name starts with an M"
17
18
       # My short friends.
19
       elif [[ " $name " = *" Short "* ]]; then
           echo "$name is a shorty."
21
22
       # Friends I kind of didn't bother to remember.
23
24
           echo 'I kind of forgot what $name is like.'
25
26
       fi
27
28
   done
```

Than something like this:

```
1 x=( Marcus\ The\ Rich JJ\ The\ Short
2 Timid\ Thomas Michelangelo\ The\ Mobster)
```

```
for name in "${x[@]}"
do if [ "$name" = "$x" ]; then echo $name was my first friend.
elif
echo $name | \
grep -qw Short
then echo $name is a shorty.
elif [ "x${name:0:1}" = "xM" ]
then echo $name starts with an M; else
echo I kind of forgot what $name \
is like.; fi; done
```

And yes, I know this is an exaggerated example, but I've seen some authentic code that actually has a lot in common with that last example.

For your own health, keep these few points in mind:

- *Healthy whitespace gives you breathing space*. Indent your code properly and consistently. Use blank lines to separate paragraphs or logic blocks.
- *Avoid backslash-escaping*. It's counter-intuitive and boggles the mind when overused. Even in small examples it takes your mind more effort to understand a\b\c than it takes to understand 'a b c'.
- Comment your way of thinking before you forget. You might find that even code that looks totally common sense right now could become the subject of "What the hell was I thinking when I wrote this?" or "What is this supposed to do?".
- Consistency prevents mind boggles. Be consistent in your naming style. Be consistent in your use of capitals. Be consistent in your use of shell features. In coding, unlike in the bedroom, it's good to be simple and predictable.

#### 12.4 Bash Tests

The test command, also known as [, is an application that usually resides somewhere in /usr/bin or /bin and is used a lot by shell programmers to perform certain tests on files and variables. In a number of shells, including Bash, test is also implemented as a shell builtin.

It can produce surprising results, especially for people starting shell scripting that think [ ] is part of the shell syntax.

If you use sh, you have little choice but to use test as it is the only way to do most of your testing.

If however you are using Bash to do your scripting (and I presume you are since you're reading this guide), then you can also use the <code>[[...]]</code> Korn-style construct. As it comes with its own micro-syntax dedicated to performing tests, it presents several advantages over the traditional <code>[/test</code> command.

Let me illustrate how [[...]] can be used to replace test, and how it can help you to avoid some of the common mistakes made by using test:

```
$ var=''
$ [ $var = '' ] && echo True
-bash: [: =: unary operator expected
$ [ "$var" = '' ] && echo True
True
$ [[ $var = '' ]] && echo True
True
```

[ \$var = '' ] expands into [ = '' ]. The first thing test does is count its arguments. Since we're using the [ form, we'll just strip off the mandatory ] argument at the end. In the first example, as we forgot to quote \$var, test sees two arguments: = and ''. It knows that if it has two arguments, the first one has to be a *unary operator* (an operator that takes one operand). But = is not a unary operator (it's binary -- it requires *two* operands), so test blows up.

Yes, test did not see our empty \$var because Bash expanded it into nothingness before test could even see it. Moral of the story? **Use more quotes!** Using quotes, [ "\$var" = '' ] expands into [ "" = '' ] and test has no problem.

Now, [[...]] can see the whole command before it's being expanded. It sees \$var, and not the expansion of \$var. As a result, there is no need for the quotes at all! In that case, [[...]] is safer.

```
$ var=
$ [ "$var" < a ] && echo True
-bash: a: No such file or directory
$ [ "$var" '<' a ] && echo True
True
$ [[ $var < a ]] && echo True</pre>
True
```

In this example we attempted a string comparison between an empty variable and 'a'. We're surprised to see the first attempt does not yield True even though we think it should. Instead, we get some weird error that implies Bash is trying to open a file called 'a'.

We've been bitten by *File Redirection*. Since test is just an application, the < character in our command is interpreted (as it should) as a *File Redirection* operator instead of the string comparison operator of test. Bash is instructed to open a file 'a' and connect it to stdin for reading. To prevent this, we need to quote < so that test receives a literal < argument that it treats as its own string comparison operator rather than Bash treating it as its redirection operator. This makes our second attempt work.

Using [[...]] we can avoid the mess altogether. [[...]] has its own micro-syntax where redirection doesn't apply and < is treated as a comparison operator when literal -- problem fixed. Once again, [[...]] is safer.

Even more dangerous is using the > operator instead of our previous example with the < operator. Since > triggers output *Redirection* it will create a file called 'a'. As a result, **there will be no error message warning us that we've committed a sin!** Instead, our script will just break. Even worse, we might overwrite some important file! It's up to us to *guess* where the problem is:

```
$ var=a
$ [ "$var" > b ] && echo True || echo False
True
$ [[ "$var" > b ]] && echo True || echo False
False
```

Two different results, not good. The lesson is to trust [[...]] more than [.["\$var" > b] is expanded into ["a"] and the output of that is being redirected into a new file called 'b'. Since ["a"] is the same as [-n "a"] and that basically tests whether the "a" string is non-empty, the result is a success and the echo True is executed.

Using [[...]] we get our expected scenario where "a" is tested against "b" and since we all know "a" sorts before "b" this triggers the echo False statement. And this is how you can break your script without realizing it. You will however have a suspiciously empty file called 'b' in your current directory.

Yes it adds a few characters, but if you're not very familiar with the shell syntax, [[...]] is likely to be safer than [. Everybody inevitably makes programming errors. Even if you try to use [ safely and carefully avoid these mistakes, I can assure you that you will make them. And if other people are reading your code, you can be sure that they'll absolutely mess things up.

Besides [[...]] provides the following features over [:

• [[...]] can do glob pattern matching:

```
[[ abc = a* ]]
```

• [[...]] can do regex pattern matching (since Bash 3.1; while its [, contrary to that of other shells such as yash or zsh still can't):

```
[[ abb =~ ^ab+$ ]]
```

The main advantage of [/test is its portability.

Another one is the fact that test's arithmetic operators (-eq, -ne, -lt, -gt, -le, -ge) work with decimal integer literals only while [[...]]'s work with any arithmetic expressions which make test's ones safer to use for arbitrary values and avoid numbers with leading 0s from being treated as octal.

```
if [ "$external_data" -gt 0 ]; then
  echo OK.
fi
```

is fine. Worse that can happen is a syntax error by [ if \$external\_data doesn't contain a valid decimal integer number.

```
# DO NOT USE
if [[ $external_data -gt 0 ]; then
# same as if [[ external_data -gt 0 ]]; then
# or if (( external_data > 0 )); then
echo OK.
fi
```

Which would introduce an arbitrary command execution vulnerability.

A few ground rules (some of which apply to most commands, not just [) can make [ safe to use:

- quote all expansions
- quote <, > so they're not treated as expansions. ( and ) would also need to be quoted but they're not needed anyway because of the next point:
- do **not** use the -a/-o (and/or) binary operators, instead use several [ invocations chained with & &/| | shell operators:

```
# Use:
if [ "$var" -ge 0 ] && [ "$var" -le 10 ]; then
  echo OK.
fi
# DO NOT use:
if [ "$var" -ge 0 -a "$var" -le 10 ]
```

A common pitfall with [[...]] is to forget the quotes in [[\$a = \$b"]] when an equality comparison is intended instead of pattern matching.

#### 12.5 Don't Ever Do These

The Bash shell allows you to do quite a lot of things, offering you considerable flexibility. Unfortunately, it does very little to discourage misuse<sup>1</sup> and other ill-advised behavior. It hopes people will find out for themselves that certain things should be avoided at all costs.

Unfortunately many people don't care enough to want to find out for themselves. They write without thinking things through and many awful and dangerous scripts<sup>2</sup> end up in production environments or in Linux distributions. The result of these, and even your very own scripts written in a time of neglect, can often be **DISASTROUS**.

That said, for the good of your scripts and for the rest of mankind, Never Ever Do Anything Along These Lines:

```
• ls -1 | awk '{ print $8 }'
```

- DON'T EVER parse the output<sup>3</sup> of 1s! The 1s command's output cannot be trusted for a number of reasons.
  - 1. For one, ls will mangle the filenames of files if they contain characters unsupported by your locale. As a result, parsing filenames out of ls is NEVER guaranteed to actually give you a file that you will be able to find. ls might have replaced certain characters in the filename by question marks.
  - 2. Secondly, 1s separates lines of data by newlines. This way, every bit of information on a file is on a new line. UNFORTUNATELY filenames themselves can ALSO contain newlines. This means that if you have a filename that contains a newline in the current directory, it will completely break your parsing and as a result, break your script!
  - 3. Last but not least, the output format of ls -l is NOT guaranteed consistent across platforms. Some systems omit the group ID by default, for instance, and reverse the effect of the -g switch. Some systems use two fields for the modification time, and some use three. On the systems that use three, the third field can be either a year, or an HH:MM concatenation, depending on how old the file is.

There are alternatives to using ls<sup>4</sup> in many cases. If you need to work with file modification times, you can typically use Bash Tests. If none of those things is possible, *I recommend you switch to a different language, like python or perl*.

```
if echo "$file" | fgrep .txt; thenls *.txt | grep story
```

 DON'T EVER test or filter filenames with grep! Unless your grep pattern is really smart it will probably not be trustworthy.

http://mywiki.wooledge.org/BashPitfalls

<sup>2</sup>http://mywiki.wooledge.org/ShellHallOfShame

<sup>3</sup>http://mywiki.wooledge.org/ParsingLs

 $<sup>^{4}</sup>$ http://mywiki.wooledge.org/BashFAQ/087

- In the first example above, the test would match both story.txt AND story.txt.exe. If you make grep patterns that are smart enough, they'll probably be so ugly, massive and unreadable that you shouldn't use them anyway.
- The alternative is called globbing<sup>1</sup>. Bash has a feature called *Pathname Expansion*. This will help you enumerate all files that match a certain pattern. Also, you can use globs to test whether a filename matches a certain glob pattern (in a case...esac or [[...]] construct).

### • cat file | grep pattern

- Don't use cat to feed a single file's content to a filter. cat is a utility used to concatenate the contents of several files together.
- To feed the contents of a file to a process you will probably be able to pass the filename as an argument to the program (grep 'pattern' /my/file, sed 'expression' /my/file, ...).
- If the manual of the program does not specify any way to do this, you should use redirection (read column1 column2 < /my/file, tr ' '\n' < /my/file, ...).</p>

### • for line in \$(<file); do

- Don't use a for loop<sup>2</sup> to read the lines of a file. Use a while read<sup>3</sup> loop instead.

### • for number in \$(seq 1 10); do

- For the love of god and all that is holy, do not use seq to count.
- Bash is able enough to do the counting for you. You do not need to spawn an external application (especially a single-platform one) to do some counting and then pass that application's output to Bash for word splitting. Learn the syntax of for already!
- In general, C-style for loops are the best method for implementing a counter for  $((i=1; i \le 10; i++))$ .
- If you actually wanted a stream of numbers separated by newlines as test input, consider printf '%d\n' {1..10}. You can also loop over the result of a sequence expansion if the range is constant, but the shell needs to expand the full list into memory before processing the loop body. This method can be wasteful and is less versatile than other arithmetic loops.

# • i=`expr \$i + 1`

- expr is a relic of ancient Rome. Do not wield it.
- It was used in scripts written for shells with very limited capabilities. You're basically spawning a new process, calling another C program to do some math for you and return the result as a string to bash. Bash can do all this itself and so much faster, more reliably (no numbers->string->number conversions) and in all, better.

http://mywiki.wooledge.org/glob

<sup>2</sup>http://mywiki.wooledge.org/DontReadLinesWithFor

<sup>3</sup>http://mywiki.wooledge.org/BashFAQ/001

- You should use this in Bash: let i++ or (( i++ ))
- Even POSIX sh can do arithmetic: i=\$((i+1)). It only lacks the ++ operator and the ((...)) command (it has only the \$((...)) substitution).

# 12.6 Debugging

Very often you will find yourself clueless as to why your script isn't acting the way you want it to. Resolving this problem is always just a matter of common sense and debugging techniques.

#### 12.6.1 Diagnose the Problem

Unless you know what exactly the **problem** is, you most likely won't come up with a *solution* anytime soon. So make sure you understand what exactly goes wrong. Evaluate the symptoms and/or error messages.

Try to formulate the problem as a sentence. This will also be vital if you're going to ask other people for help with your problem. You don't want them to have to go through your whole script or run it so that they understand what's going on. No; **you** need to make the problem perfectly clear to **yourself** and to anybody trying to help you. *This requirement stands until the day the human race invents means of telepathy*.

#### 12.6.2 Minimize the Codebase

If staring at your code doesn't give you a divine inspiration, the next thing you should do is try to minimize your codebase to isolate the problem.

Don't worry about preserving the functionality of your script. The only thing you want to preserve is the logic of the code block that seems buggy.

Often, the best way to do this is to copy your script to a new file and start deleting everything that seems irrelevant from it. Alternatively, you can make a new script that does something similar in the same code fashion, and keep adding structure until you duplicate the problem.

As soon as you delete something that makes the problem go away (or add something that makes it appear), you'll have found where the problem lies. Even if you haven't precisely pinpointed the issue, at least you're not staring at a massive script anymore, but hopefully at a stub of no more than 3-7 lines.

For example, if you have a script that lets you browse images in your image folder by date, and for some reason you can't manage to iterate over your images in the folder properly, it suffices to reduce the script to this part:

```
for image in $(ls -R "$imgFolder"); do
    echo "$image"
done
```

Your actual script will be far more complex, and the inside of the for loop will also be far longer. But the essence of the problem is this code. Once you've reduced your problem to this it may be easier to see the problem you're facing. Your echo spits out parts of image names; it looks like all whitespace is replaced by newlines. That must be because echo is run once for each chunk terminated by whitespace, not for every image name (as a result, it seems the output has split open image names with whitespace in them). With this reduced code, it's easier to see that the cause is actually your for statement that splits up

the output of ls into words. That's because ls is **UNPARSABLE** in a bugless manner (do not ever use ls in scripts, unless if you want to show its output to a user).

We can't use a recursive glob (unless we're in bash 4), so we have to use find to retrieve the filenames. One fix would be:

```
find "$imgFolder" -print0 | while IFS= read -r -d '' image; do
    echo "$image"
done
```

Now that you've fixed the problem in this tiny example, it's easy to merge it back into the original script.

### 12.6.3 Activate Bash's Debug Mode

If you still don't see the error of your ways, Bash's debugging mode might help you see the problem through the code.

When Bash runs with the  $\times$  option turned on, it prints out every command it executes before executing it (to standard error). That is, **after any expansions have been applied**. As a result, you can see exactly what's happening as each line in your code is executed. Pay very close attention to the quoting used. Bash uses quotes to show you exactly which strings are passed as a single argument.

There are three ways of turning on this mode.

• Run the script with bash -x:

```
$ bash -x ./mybrokenscript
```

• Modify your script's header:

```
#!/bin/bash -x
[.. script ..]
```

• Or:

```
#!/usr/bin/env bash
set -x
```

• Or add set -x somewhere in your code to turn on this mode for only a specific block of your code:

```
#!/usr/bin/env bash
[..irrelevant code..]
set -x
[..relevant code..]
set +x
[..irrelevant code..]
```

Because the debugging output goes to stderr, you will generally see it on the screen, if you are running the script in a terminal. If you would like to log it to a file, you can tell Bash to send all stderr to a file:

```
exec 2>> /path/to/my.logfile
set -x
```

A nice feature of bash version >= 4.1 is the variable BASH\_XTRACEFD. This allows you to specify the file descriptor to write the set -x debugging output to. In older versions of bash, this output always goes to stderr, and it is difficult if not impossible to keep it separate from normal output (especially if you are logging stderr to a file, but you need to see it on the screen to operate the program). Here's a nice way to use it:

```
# dump set -x data to a file
# turns on with a filename as $1
# turns off with no params
setx_output()

{
    if [[ $1 ]]; then
        exec {BASH_XTRACEFD}>>"$1"
        set -x

else
        set +x
        unset -v BASH_XTRACEFD

fi
}
```

If you have a complicated mess of scripts, you might find it helpful to change PS4 before setting -x. If the value assigned to PS4 is surrounded by double quotes it will be expanded during variable assignment, which is probably not what you want; with single quotes, the value will be expanded when the PS4 prompt is displayed.

```
PS4='+$BASH_SOURCE:$LINENO:$FUNCNAME: '
```

# 12.6.4 Step Your Code

If the script goes too fast for you, you can enable code-stepping. The following code uses the DEBUG trap to inform the user about what command is about to be executed and wait for his confirmation do to so. Put this code in your script, at the location you wish to begin stepping:

```
debug_prompt () { read -p "[$BASH_SOURCE:$LINENO] $BASH_COMMAND?" _ ;}
trap 'debug_prompt "$_"' DEBUG
```

### 12.6.5 The Bash Debugger

The Bash Debugger Project is a gdb-style debugger for bash, available from http://bashdb.sourceforge.net/ The Bash Debugger will allow you to walk through your code and help you track down bugs.

### 12.6.6 Reread the Manual

If your script still doesn't seem to agree with you, maybe your perception of the way things work is wrong. Try going back to the manual (or this guide) to re-evaluate whether commands do exactly what you think they do, or the syntax is what you think it is. Very often people misunderstand what for does, how *Word Splitting* works, or how often they should use quotes.

Keep the tips and good practice guidelines in this guide in mind as well. They often help you avoid bugs and problems with scripts.

I mentioned this in the *Scripts* section of this guide too, but it's worth repeating it here. First of all, make sure your script's **header** is actually **#! /bin/bash**. If it is **missing** or if it's something like **#! /bin/sh** then you deserve the problems you're having. That means you're probably not even using Bash to run your code. Obviously that'll cause issues. Also, make sure you have no *Carriage Return* characters at the ends of your lines. This is the cause of scripts written in *Microsoft Windows(tm)*. You can get rid of these fairly easily like this:

```
$ tr -d '\r' < myscript > tmp && mv tmp myscript
```

### 12.6.7 Read the FAQ / Pitfalls

The BashFAQ<sup>1</sup> and BashPitfalls<sup>2</sup> pages explain common misconceptions and issues encountered by other Bash scripters. It's very likely that your problem will be described there in some shape or form.

To be able to find your problem in there, you'll obviously need to have *Diagnosed* it properly. You'll need to know what you're looking for.

### 12.6.8 Ask Us on IRC

There are people in the #bash channel almost 24/7. This channel resides on the Libera Chat<sup>3</sup> IRC network. To reach us, you need an IRC client. Connect it to irc.libera.chat, and /join #bash.

Make **sure** that you know what your real problem is and have stepped through it on paper, so you can explain it well. We don't like having to guess at things. Start by explaining what you're trying to do with your script.

Either way, please have a look at this page before entering #bash: XyProblem<sup>4</sup>.

http://mywiki.wooledge.org/BashFAQ

<sup>2</sup>http://mywiki.wooledge.org/BashPitfalls

<sup>3</sup>https://libera.chat

<sup>4</sup>http://mywiki.wooledge.org/XyProblem