++Introduction

You are invited to make additions or modifications 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.

BASH is a BourneShell compatible shell, which adds many new features to its ancestor. Most of them are available in the 'KornShell', too.

++About This Guide

This guide aims to become a starting point for people interested in learning to work with BASH. It aspires to teach its readers good practice techniques for developing scripts for the BASH interpreter, and to educate them about the internal operation of BASH.

This guide is targeted at beginning users. It assumes no basic knowledge, but rather expects you to have enough common sense to put two and two together. If something is unclear to you, you are invited to report this (use BashGuideFeedback, or the #bash channel on irc.freenode.org) 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.

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+ + 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 batch processes quickly. In no way are they required for the execution of processes; they are merely a layer between system function calls and the user.

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 the interface for you to execute statements (using BASH syntax), either at the

In The Manual: • Introduction

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.

++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* through scripts. We can use scripts to automate certain logic. Scripts are basically lists of commands that you can type on the command line. When such 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 opens documentation (so-called "man pages") on a certain topic. 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.

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 FAO:

Is there a list of which features were added to specific releases (versions) of Bash?

Interactive mode: A mode of operation where a prompt asks you for one command at a time.

++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.
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 - Variables; special parameters; parameter types; parameter expansion.
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++Commands and Arguments

BASH reads commands from its input (which is 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, but we won't worry about that just yet.)

BASH divides each line into *words* at each whitespace character (spaces and tabs). The first word it finds is the name of the command to be executed. All the remaining words become *arguments* to that command (options, filenames, etc.).

Assume you're in an empty directory. (If you want to try this code out, you can create

and go into an empty directory called test 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 all files again; this time the output
shows 'a', 'b' and 'c'.
a b c
```

touch is an application that changes the 'Last Modified'-time of a certain file to the current time. If the filename that it's given does not exist yet, it simply creates that file, 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 *  # Remove all files in the current directory.
$ ls  # List files in the current directory (no
output: no files).
$ touch a b c  # Create files 'a', 'b' and 'c'.
$ ls  # List all files again; this time the output
shows 'a', 'b' and 'c'.
a b c
```

rm is an application that removes all the files that it was given. * is a *glob*. It basically means *all files in the current directory*. You will read more about this later on.

Now, did you 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. You now know that 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.
```

In this case, 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 actually want the extra whitespace, we need to pass the sentence as one single argument. We can do this by using quotes:

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

Quotes group everything inside them into a single argument. This argument is 'This is a test.', properly 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:

```
$ ls
                                                       # List files in
the current directory.
    The secret voice in your head.mp3 secret
    $ rm The secret voice in your head.mp3
                                                       # Executes rm with
6 arguments; not 1!
    rm: cannot remove `The': No such file or directory
    rm: cannot remove `voice': No such file or directory
    rm: cannot remove `in': No such file or directory
    rm: cannot remove `your': No such file or directory
rm: cannot remove `head.mp3': No such file or directory
    $ 1s
                                                       # List files in
the current directory: It is still there.
    The secret voice in your head.mp3
                                                       # But your file
'secret' is now gone!
```

You need to make sure you quote filenames properly. If you don't you'll end up deleting the wrong things! rm takes filenames as arguments. If you do not quote filenames with spaces, rm thinks that each argument is a separate file. Since BASH splits your arguments at the spaces, rm will try to remove each word. The above example tried to

delete files for each word in the filename of the song, instead of the filename of the song. That caused our file secret to be deleted, and our song to remain behind!

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

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

You want the [command name to be separated from the arguments -f, file and]. If you do not separate [and -f from each other with whitespace, bash will think you are trying to execute the command name [-f and look in PATH for a program named [-f. Additionally, the arguments file and] need to be separated by spaces. The [command expects its last argument to be]. The correct command separates all arguments with spaces:

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

In BASH scripts and functions, arguments that were passed to the script or function are saved in *Positional Parameters*. You can read these by using \$1,\$2, and so on for the respective argument. You can also use \$@ and \$* but more about this later on.

NOTE: Please have a good look at Arguments, Quotes, WordSplitting and http://wiki.bash-hackers.org/syntax/words if all this isn't very clear to you yet. It is important that you have a good grasp of how the shell interprets the statements you give it before you continue this guide.

Good Practice:

You should *always* quote sentences or strings that belong together, even if it's not absolutely necessary. This will keep you alert and reduce the risk of human error in your scripts.

For example, you should always quote arguments to the echo command.

In the FAO:

I'm trying to put a command in a variable, but the complex cases always fail! How can I handle command-line arguments (options) to my script easily?

Arguments: These are the optional additional words you can specify when running commands. They are given after the command's name ('ls -l foo' executes ls with two arguments).

Quotes: The two forms of quotes (' and ") are used to protect certain special characters inside them from being interpreted as special by BASH. The difference between ' and " will be discussed later.

++Types of Commands

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

• Aliases: Aliases are a way of shortening commands. They are only used in **interactive** shells, not in **scripts**. An alias is a *name* that is mapped to a certain *string*. Whenever that *name* is used as a command name, it is replaced by the *string* before executing the command. So, instead of executing:

```
$ nmap -P0 -A --osscan_limit 192.168.0.1
```

You could use an alias like this:

```
$ alias nmapp='nmap -P0 -A --osscan_limit'
$ nmapp 192.168.0.1
```

• Functions: Functions in BASH are somewhat like aliases, but more powerful.

Unlike aliases they can be used in **scripts**. A function contains shell commands, very much like a small script. When a function is called, the commands in it are executed. Functions will be covered in depth later in the guide.

- **Builtins**: BASH has some basic commands built into it, such as cd (change directory), echo (write output), and so on. You can think of them as functions that are provided already.
- **Keywords**: Keywords are quite like builtins, but the main difference is that special parsing rules apply to them. For example, [is a bash builtin, while [[is a bash keyword. They are both used for testing stuff, but since [[is a keyword rather than a builtin, it benefits from a few special parsing rules which make it a lot easier:

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

The first example returns an error because bash tries to redirect the file b to the command [a] (See File Redirection). The second example actually does what you expect it to. The character < no longer has its special meaning of *File Redirection* operator.

• Executables: The last kind of command that can be executed by bash is an executable, also called an external command or application. Executables are invoked by using a pathname. If the executable is in the current directory, use ./myprogram. If it's in the /usr/local/bin directory, use /usr/local/bin/myprogram.

To make life a little easier for you, though, BASH uses a variable that tells it where to find applications in case you just know the name of the application but not its full pathname. This variable is called PATH, and it is a set of directory names separated by colons — for example, /bin:/usr/bin. When a command is specified in BASH without a pathname (e.g. myprgram, or ls), and it isn't an alias, function, builtin or keyword, BASH searches through the directories in PATH, in order from left to right, to see whether they contain an executable of the name you typed.

Tip:

You can use the type command to figure out the type of a command. For example:

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

In The Manual: • Simple Commands

In the FAQ:

What is the difference between test, [and [[? How can I make an alias that takes an argument?

Alias: A name that is mapped to a string. Whenever that name 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 that name is used as a command, the function is called with the arguments provided to the function's name on the command line. Functions are the basic method of making new commands.

Builtin: Certain commands have been built into BASH. These are handled internally whenever they are executed on the command line (and do not create a new process).

Application: A program that can be executed by referring to its pathname (/bin/ls), or simply by its name if its location is in your PATH variable.

++Scripts

A script is basically a sequence of commands in a file. BASH reads the file and processes the commands in order. It only moves on to the next command when the current one has ended, unless the current one has been executed asynchronously (in the background). Don't worry too much about the latter case yet -- you'll learn about how that works later on.

Virtually any example that you see in this guide can be used in a script just as well as on the command line.

Making a script is easy. You just make a new file, and put this in it at the top:

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

This header makes sure that whenever your script is executed, BASH will be used as its interpreter. **Please** do not be fooled by examples on the Internet that use /bin/sh as interpreter. **sh** is **not** bash. Even though sh's syntax and bash's look very much alike and even though most bash scripts will run in sh, a lot of the examples in this guide only apply to bash and will just break or cause unexpected behaviour in sh. Also, please refrain from giving your scripts that stupid . sh extension. It serves no purpose, and it's completely misleading (since it's going to be a bash script, not an sh script).

And by the way, it's perfectly fine if you use *Windows* to write your scripts, but if at all possible, **avoid using** *Notepad* **for writing scripts**. *Microsoft Notepad* can only make files with DOS-style line-endings. That means that each line you make in notepad will be ended by two characters: a *Carriage Return* and a *Newline* character. BASH reads lines as terminated by *Newline* characters only. As a result, the *Carriage Return* character will cause you incredible headache if you don't know it's there (very weird error messages). If at all possible, *use a decent editor* like Vim, Emacs, kate, GEdit, GVIM or xemacs. If you don't, then you will need to remove the carriage returns from your scripts before running them.

Once your script file has been made, you can run it like this:

```
$ bash myscript
```

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

Alternatively, you can give your script executable permissions. When you do this, you can actually execute the script as an application instead of calling BASH manually:

```
$ chmod +x myscript
$ ./myscript
```

When executed in this way, the #! line tells the operating system (OS) what interpreter to use. The OS runs /usr/bin/env, which in turn runs bash, which reads our script.

Some people like to keep their scripts in a personal directory. Others like to keep their scripts somewhere in the PATH variable. Most like to do both at once. Here's what I suggest you do:

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

The first command will make a directory called bin in your home directory. The second command will add a line to your .bashrc file which adds the directory we just made to the beginning of the PATH variable. Every new instance of BASH will now check for executable scripts in your bin directory. Finally, the third line replaces our current instance of BASH with a new one, which reads the .bashrc file.

Changes to DotFiles (such as .bashrc) never have an immediate effect. You have to

take some step to re-read the files. In the example above, we used exec bash to replace the running shell. If you wanted, you could close your existing terminal and open a new one. BASH would then initialize itself again by reading .bashrc (and possibly other files). Or, you could just execute that line of code on the command line (PATH="\$HOME/bin:\$PATH") or manually process your .bashrc file in the running shell by running source "\$HOME/.bashrc".

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 its path, which was the . / part in the previous examples):

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

Tip:

While you're defining the interpreter in your header, you might want to take the time to explain your script's function and expected arguments a little too:

```
#! /usr/bin/env bash
#
# scriptname argument [argument] ...
#
# A short explanation of your script's purpose.
#
# Copyright [date], [name]
```

Tip:

You can use this header to specify up to one word of optional arguments that you want to pass to the interpreter. For example, the following arguments will turn on some verbose debugging:

```
#! /bin/bash -xv
```

Unfortunately, you can't use this:

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

because that would require two words of arguments in the header line. Unix doesn't allow that. You could do this instead:

```
#! /usr/bin/env bash
set -xv
```

For more hints, see Debugging.

Header: The header of a script determines the application that will function as its interpreter (e.g. bash, sh, perl, ...). Colloquially, this is also called a *shebang* - a slang term derived by combining *hash* (#) and *bang* (!). You will probably see the word *shebang* more often than *header*.

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++Special Characters

There are several special characters in BASH that have a non-literal meaning. When we use these characters, BASH evaluates these characters and their meaning, but usually does not pass them on to the underlying commands. These are also called *metacharacters*.

Here are a few of those special characters, and what they do:

- [whitespace]: Whitespace (spaces, tabs and newlines). BASH uses whitespace to determine where words begin and end. The first word of each command is used as the command name; any additional words become arguments to that command.
- \$: Expansion character. This character is used in most substitutions, including parameter expansion (variable substitution). More about this later.
- 'text': Single quotes protect the text inside from any kind of expansion by the shell and keep it from being split into multiple words or arguments. They also prevent the special meaning of all special characters inside.
- "text": Double quotes protect the text inside from being split into multiple words or arguments, but they permit substitutions to occur. They prevent the special meaning of *most* special characters inside them -- basically, all except for the \$.
- #: Comment character. Any word beginning with # begins a *comment* that extends to the next newline. Comments are not processed by the shell.
- ;: Command separator. The semicolon is used to separate multiple commands from each other if the user chooses to keep them on the same line. It's *basically* the same thing as a newline.
- \: Escape character. The backslash prevents the next character from being used in any special sort of way. This works inside double quotes, or outside of all quotes, but *not* inside single quotes.
- > or <: Redirection characters. These characters are used to modify (redirect) the input and/or output of a command. Redirections will be covered later.
- I: Pipelines allow you to send the output of one command as the input to another command.
- [[expression]]: Test expression. This evaluates the conditional expression as a logical statement, to determine whether it's "true" or "false".
- { commands; } : Command grouping. The commands inside the braces are treated as though they were only one command. It is convenient for places where BASH syntax requires only one command to be present, and you don't feel a function is warranted.
- `command`, \$ (command): Command substitution (The latter form is highly preferred.) Command substitution executes the inner command first, and then replaces the whole `...` or \$ (...) with that command's standard output.
- (command): Subshell Execution. This executes the command in a new bash shell, instead of in the current one. If the command causes side effects (like changing variables), those changes will have no effect on the current shell.
- ((expression)): Arithmetic Command. Inside the parentheses, operators such as +, -, * and / are seen as mathematical operators. This can be used for assignments like ((a=\$b+7)) as well as tests like if ((\$a < \$b)). More on this later.
- \$((expression)): Arithmetic Substitution. Comparable to the above, but this expression is replaced with the result of its arithmetic evaluation. Example: echo "The average is \$(((\$a+\$b)/2))".

Some 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 bigger than zero."
Five is bigger than zero.
```

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their meaning is interpreted and then they are removed from the command before executing it.

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- 2. Special Parameters and Variables
- 3. Variable Types
- 4. Parameter Expansion

+ + Parameters

Parameters should be seen as a sort of named space in memory where you can store your data. Generally speaking, they will store string data, but can also be used to store integers or arrays.

Parameters can be variables or special parameters. Special parameters are read-only parameters pre-set by BASH and report 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 alphanumeric characters and underscores, and beginning with an alphabetic character or an underscore. Also referred to as an *identifier*.

To assign to variables, 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 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 that is stored in variables, 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 it to use the contents of the variable. After that, BASH still performs 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 deleting it from the command. Then, it replaces 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, because of the following case which relies on you having understood the chapter on argument splitting above:

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

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!**

```
$ rm "$song"
```

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

++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 parameters: 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 was started with these options: $-"
My shell is -bash, and was started with these options: 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 it 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*:

- 0: Contains the name of the script. (This is not always reliable.)
- **Positional Parameters**: 1, 2, ...; They contain the arguments that were passed to the current script.
- *: Expands to all the words of all the positional parameters. If double quoted, it expands to a single string containing all the positional parameters separated by the first character of the **IFS** variable (which will be discussed later).
- @: Expands to all the words of all the positional parameters. If double quoted, it expands to a list of all the positional parameters 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 (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 some examples of *Variables* that the shell initializes 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 lines.)
- **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).

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 well 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 FAQ:

How can I concatenate two variables? How do I append a string to a variable?

How can I access positional parameters after \$9?

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, 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.

+ + 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 have. They are stored in the variable's attributes.

Attributes are settings for a variable. They define the way the variable will behave. Here are the attributes you can assign to a variable:

- 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 subshell or child process.

Arrays are basically indexed lists of strings. They are very convenient for their ability to store multiple elements together without relying on a delimiter. That way, you don't need to worry about the fact that this delimiter could possibly end up being part of an element's content and thus split that element up.

(If you want a more comprehensive look at arrays, see arrays in the FAQ.)

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. For the most part, experienced shell scripters prefer to use explicit arithmetic commands (let or ((...))) when they want to perform arithmetic.

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 subshell or child process.

In the FAQ: How can I use array variables?

+ + 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 parameter 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 "'$USER', '$USERs', '${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 can also be used to modify the string that will be expanded. These operations are 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 PEs that are available:

- **\${parameter:-word}**: Use Default Values. If 'parameter' is unset or null, the expansion of 'word' is substituted. Otherwise, the value of 'parameter' is substituted.
- **\${parameter:=word}**: Assign Default Values. If 'parameter' is unset or null, the expansion of 'word' is assigned to 'parameter'. The value of 'parameter' is then substituted.
- **\${parameter:?word}**: Display Error if Null or Unset. If 'parameter' is null or unset, the expansion of 'word' (or a message to that effect if 'word' is not present) is written to the standard error and the shell, if it is not interactive, exits. Otherwise, the value of 'parameter' is substituted.
- **\${parameter:+word}**: Use Alternate Value. If 'parameter' is null or unset, nothing is substituted, otherwise the expansion of 'word' is substituted.
- **\${parameter:offset:length}** Substring Expansion. Expands to up to 'length' characters of 'parameter' starting at the character specified by 'offset'. If 'length' is omitted, expands to the substring of 'parameter' starting at the character specified by 'offset'.
- **\${#parameter}**: The length in characters of the value of 'parameter' is substituted.
- **\${parameter#pattern}**: The 'pattern' is anchored to the beginning of 'parameter'. The result of the expansion is the expanded value of 'parameter' with the shortest match deleted.
- **\${parameter##pattern}**: The 'pattern' is anchored to the beginning of 'parameter'. The result of the expansion is the expanded value of 'parameter' with the longest match deleted.

- **\${parameter*pattern}**: The 'pattern' is anchored to the end of 'parameter'. The result of the expansion is the expanded value of 'parameter' with the shortest match deleted.
- **\${parameter%*pattern}**: The 'pattern' is anchored to the end of 'parameter'. The result of the expansion is the expanded value of 'parameter' with the longest match deleted.
- \${parameter/pattern/string}: The 'pattern' is not anchored but evaluated from left to right in the value of 'parameter'. The result of the expansion is the expanded value of 'parameter' with the first match of 'pattern' replaced by 'string'.
- **\${parameter//pattern/string}**: As above, but every match of 'pattern' is replaced.

You will learn them all through experience. They will come in handy far more often than you think they might. Here's 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 $\{v\#p\}$ and $\{v\#p\}$. 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

In the FAQ:

Is there a function to return the length of a string?

How can I rename all my *.foo files to *.bar, or convert spaces to underscores, or convert upper-case file names to lower case?

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?

How do I get the effects of those nifty Bash Parameter Expansions in older shells?

How do I determine whether a variable is already defined? Or a function?

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.

- <- Special Characters | Patterns ->
- <- Parameters | Tests and Conditionals ->

+ + Patterns

BASH 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.

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

+ + Glob Patterns

Globs 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 meta characters. Meta characters are characters that have a special meaning. These are the basic meta characters:

- *: 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 enumerated and used in place of the glob. As a result, the statement echo a* is

replaced by the statement echo a abc, and is then executed.

BASH performs filename expansions *after* word splitting has already been done; therefore, filenames generated by a glob 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. It is important to understand that 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 file "a b.txt", which for takes as a single argument.

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 shell options:

```
$ 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 except one of the given patterns.

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

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

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

In addition to filename expansion, globs may also be used to check 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 builtin command (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.

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

```
$ 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
```

Good Practice:

You should always use globs instead of 1s (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

In the FAO:

How can I use a logical AND/OR/NOT in a shell pattern (glob)?

Glob: A glob is a string that can match certain strings or filenames.

+ ↑Regular Expressions

Regular expressions (regex) are similar to *Glob Patterns* but cannot be used for filename matching in BASH. Since 3.0, BASH supports the =~ 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, or Extended Regular Expressions.

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:

```
$ if [[ $LANG =~ (..)_(..) ]]
> then echo "You live in ${BASH_REMATCH[2]} and speak
${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.

```
$ [[ "My sentence" =~ My\ sentence ]]
```

Be careful to escape any characters that the shell could misinterpret, such as whitespace, dollar signs followed by text, braces, etc.

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, section E14.

In The Manual: ● **Regex(3)**

In the FAQ:

I want to check if [[\$var == foo or \$var == bar or \$var == more ... without repeating \$var n times.

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).

- <- Parameters | Tests and Conditionals ->
- <- Tests and Conditionals | Input and Output ->

++Arrays

As mentioned earlier, BASH provides three types of parameters: Strings, Integers and 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.

→ ↑ 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 1s. The first would create a **string** with the output of 1s. 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. 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 -a ip_elements <<< "127.0.0.1"</pre>
```

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 – print0, which we'll use in this example:

```
files=()
while read -r -d $'\0'; do
    files+=("$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 that runs a read command each time. The read command uses the $-d \$ '\0' option, which means that instead of reading a line at a time (up to a newline), we're reading up to a NUL byte (\0). 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.

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.

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

In the FAQ:

How can I use array variables?

How can I use variables (indirect variables, pointers, references) or associative arrays?

How can I find and deal with file names containing newlines, spaces or both? I set variables in a loop. Why do they suddenly disappear after the loop terminates? Or, why can't I pipe data to read?

+ † Using Arrays

properly quoted.

Walking over array elements is really easy. Because an array is such a safe medium of storage, we can simply use a for loop to iterate over its elements:

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

Notice the syntax used to **expand** the array here. We use the **quoted** form: "\${myfiles[@]}". Bash replaces this syntax with every single element in the array,

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:

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

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

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 spare, 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 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 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).

The printf command deserves special mention here, because it's a supremely elegant way to dump an array:

```
$ names=("Bob" "Peter" "$USER" "Big Bad John")
$ printf "%s\n" "${names[@]}"
Bob
Peter
lhunath
Big Bad John
```

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
```

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.

+ ↑ 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.

To create an associative array, you need to declare it as such (using declare -A). This is to guarantee backward compatibility with the standard indexed arrays. Here's how you do that:

```
$ 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.
```

With the same syntax as for indexed arrays, you can iterate over the keys 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.

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 actually an arithmetic context (really, you can do math there without an explicit \$ ((. . .)) markup). 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 + 1]}"
```

As you can see, both \$index and index work fine with indexed arrays. They both evaluate to 0. You can even do math on it to increase it to 1 and get the second value. No go with associative arrays, though. Here, we need to use \$key; the others fail.

- <- Tests and Conditionals | Input and Output ->
- <- Patterns | Arrays ->

++Tests and Conditionals

Contents

- 1. Exit Status
- 2. Control Operators (&& and ||)
- 3. Grouping Statements
- 4. Conditional Blocks (if, test and [])
- 5. Conditional Loops (while, until and for)
- 6. Choices (case and select)

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.

++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 God
ping: unknown host God
$ 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!'; exit 1; }
```

In The Manual: • 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.

++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 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
```

| | 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 | | 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 himself.

In The Manual: Lists of Commands

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.

++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:

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

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:

(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 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:

```
cd "$appdir" \mid\,\mid { echo "Please create the appdir and try again"; exit 1; }
```

++4. Conditional Blocks (if, test and [[)

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

```
$ 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
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 [[. [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 we know this is not the case, [will not exit successfully (its exit code will be 1). if sees that [terminated unsuccessfully and executes the code in the else block.

Now, to see why [[is so much more interesting and trustworthy than [, let us highlight some possible problems with [:

```
$ 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 which are missing from [.

One of the features of [[is pattern matching:

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

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. [[can't know whether your literal whitespace in the statement is intentional or not; so it splits it up 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

```
$ foo=[a-z]* name=lhunath
$ [[ $name = $foo ]] && echo "Name $name matches pattern $foo"
Name lhunath matches pattern [a-z]*
$ [[ $name = "$foo" ]] || echo "Name $name is not equal to the
string $foo"
Name lhunath is not equal to the string [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: Always quote stuff if you are unsure. If foo **really** contains a pattern instead of a string (a **rare** thing to want -- you would normally write the pattern out literally: [[name = [a-z]*]]), you will get a safe error here and you can come and fix it. If you neglect to quote, bugs can become very hard to find, since the broken code may not fail immediately.

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

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):
 - **-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.
 - **-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 (it's length is zero).
 - **-n STRING**: True if the string is not empty (it's length is not zero).
 - **STRING = STRING**: True if the first string is identical to the second.
 - **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 -a EXPR**: True if both expressions are true (logical AND).
 - **EXPR -o EXPR**: True if either expression is true (logical OR).
 - ! EXPR: Inverts the result of the expression (logical NOT).
 - 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 =~ REGEX**: True if the string matches the regex pattern.
 - (EXPR): Parantheses 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** II **EXPR**: Much like the '-o' operator of test, but does not evaluate the second expression if the first already turns out to be true.

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

In the FAO:

How can I group expressions, e.g. (A AND B) OR C?

What is the difference between the old and new test commands ([and [[)?

How do I determine whether a variable contains a substring?

How can I tell whether a variable contains a valid number?

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.

++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

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.
- **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). 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 `world.mp3': No such file or directory
```

You should already know to quote the \$file in the rm statement; but what's going wrong here? BASH expands the command substitution (\$(ls *.mp3)), 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:

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

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
```

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
```

```
$ # 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.

```
# A simple menu:
while true; do
    echo "Welcome to the Menu"
    echo " 1. Say hello"
    echo " 2. Say good-bye"

read -p "-> " response
    case $response in
        1) echo 'Hello there!';;
        2) echo 'See you later!'; break;;
        *) echo 'What was that?';;
    esac
done
```

In The Manual: • Looping Constructs

In the FAQ:

How can I run a command on all files with the extention .gz?

How can I use numbers with leading zeros in a loop, e.g. 01, 02?

How can I find and deal with file names containing newlines, spaces or both?

How can I rename all my *.foo files to *.bar, or convert spaces to underscores, or convert upper-case file names to lower case?

Can I do a spinner in Bash?

I want to check to see whether a word is in a list (or an element is a member of a set).

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. Choices (case and select)

Sometimes you want to build application logic depending on the content of a variable, and you already know all of the possible values the variable can contain. You no longer want if's two *yes or no* blocks; instead, you want a block of code for each possible

value. You could obviously implement this behaviour using several if-statements:

```
$ if [[ $LANG = en* ]]
> then echo 'Hello!'; fi
$ if [[ $LANG = fr* ]]
> then echo 'Salut!'; fi
$ if [[ $LANG = de* ]]
> then echo 'Guten Tag!'; fi
$ if [[ $LANG = nl* ]]
> then echo 'Hallo!'; fi
$ if [[ $LANG = it* ]]
> then echo 'Ciao!'; fi
```

But let's say we like to actually make clean code, shall we? 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';;
> *) echo 'I do not speak your language.';;
> 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). 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.

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."
> done
```

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.

Good Practice:

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

In the FAQ:

I want to check if [[\$var == foo or \$var == bar or \$var = more ... without repeating \$var n times.

How can I handle command-line arguments (options) to my script easily?

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.

- <- Patterns | Arrays ->
- <- Tests and Conditionals | Compound Commands ->

++Input And Output

Contents

- 1. Command-line Arguments
- 2. The Environment
- 3. File Descriptors
- 4. Redirection
 - 1. File Redirection
 - 2. File Descriptor Manipulation
 - 3. Heredocs And Herestrings
- 5. Pipes
- 6. Miscellaneous Operators
 - 1. Process Substitution

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 the bread and butter of 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.

++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?

++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.) 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). Environment variables can be set in each user's DotFiles, 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
```

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.

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)!

++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 being read from or written to the FD's data source.

By default, every new process starts with three FDs. They are referred to by the names *Standard Input*, *Standard Output* and *Standard Error*. In short, they are respectively called stdin, stdout and stderr. 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. Be aware that GUI applications work in the same way; but the actual GUI doesn't work via these FDs. GUI applications can still read and write from and to the standard FDs, but they usually don't. Usually, they do all the user interaction via that GUI; making it hard for BASH to control. As a result, we'll stick to simple terminal applications. Those, we can easily feed data on their "Standard Input", and read data from 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 uses stdout to send its output to. stdin is connected to your terminal's input device, which is probably your keyboard. stdout is connected to your terminal's output device, which we assume is your computer 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, this separation makes it easy to keep errors separated from the application's normal messages. Some people like to use wrappers to make all the output on stderr red, so that they can see the error messages more clearly. This is not generally advisable, but it is a simple example of the many options this separation provides us with. Another common practice is writing stderr to a special log file.

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.

++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 an application's FDs. That way, you can send the application's output to a file instead of the terminal, or have the application read from a file instead of from the keyboard.

Redirection, too, comes in different shapes. There's *File Redirection*, *File Descriptor manipulation*, *Heredocs* and *Herestrings*.

Redirection: This is the practice of changing a certain FD to read its input from or send its output to elsewhere.

+ + 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 outputs 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. There will almost always be far better ways to do this. Please keep this warning in mind. Useless use of cat will merely result in an extra process to create, and often results in poorer read speed because cat cannot determine the context of what it's reading and the purpose for that data.

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
```

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

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

Why does it say test? twice now? Well, as you type, your terminal shows you all the characters that you send to stdin before sending them there. That results in the first test? that you see. As soon as you hit *Enter*, cat reads a line from stdin, and 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 from it and return you to your prompt.

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** 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 it changes.

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 was open on FD 1, that's the number we precede the redirection operator with. Note, the effect of 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 the output of 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 an 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:

```
> 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*.

In The Manual: ● Redirecting Input, ● Redirecting Output, ● Appending Redirected Output, ● Redirecting Standard Output and Standard Error

+ +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 desination 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 stderr to where stdout points (which is the terminal), 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. 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 Manual: • Duplicating File Descriptors, • Moving File Descriptors, • Opening File Descriptors for Reading and Writing

In the FAQ:

How can I redirect the output of multiple commands at once?

How can I redirect the output of 'time' to a variable or file?

How do I use dialog to get input from the user?

How can I redirect stderr to a pipe?

Tell me all about 2>&1 -- what's the difference between 2>&1 >foo and >foo 2>&1, and when do I use which?

+ + 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>
```

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>
```

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.

In The Manual: • Here Documents, • Here Strings

+ + 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

In the FAO:

I set variables in a loop. Why do they suddenly disappear after the loop terminates? Or, why can't I pipe data to read?

How can two processes communicate using named pipes (fifos)?

How can I redirect stderr to a pipe?

How can I read a file line-by-line?

Tell me all about 2>&1 -- what's the difference between 2>&1 >foo and >foo 2>&1, and when do I use which?

++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.

+ + 6.1. Process Substitution

A cousin of the pipe is the process substitution operator, which comes in two forms: <() and >(). It's a convenient way to use named pipes without having to create temporary files. Whenever you think you need a temporary file to do something, process substitution might be a better way to handle things.

What it does, is basically run the command inside the parentheses. With the < () operator, the command's output is put in 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.

Here's how we can put that into action: Imagine a situation where 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:

```
$ head -n 1 .dictionary > file1
$ tail -n 1 .dictionary > file2
$ diff -y file1 file2
Aachen
zymurgy
$ rm file1 file2
```

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

```
$ diff -y <(head -n 1 .dictionary) <(tail -n 1 .dictionary)
Aachen
zymurgy</pre>
```

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, redirecting their respective outputs to 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 of the temporary files 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:

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

The > (..) operator is much like the < (..) operator, but instead of redirecting the command's output to a file, we redirect the file to the command's input. It's used for cases where you're running a command that writes to a file, but you want it to write to another command instead:

```
$ tar -cf > (ssh host tar x) .
```

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.

- <- Tests and Conditionals | Compound Commands ->
- <- Input and Output | Sourcing ->

++Compound Commands

Contents

- 1. Compound Commands
 - 1. Subshells
 - 2. Command grouping
 - 3. Arithmetic Evaluation
 - 4. Functions
 - 5. Aliases
 - 6. Destroying Constructs

BASH 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.

+ ↑1. Subshells

A SubShell 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 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.

+ ↑2. Command grouping

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

Commands may be grouped together using curly braces. This looks somewhat like a subshell, but it isn't. Command groups are executed in the same shell as everything else, rather than a new one.

Command groups can be used to run multiple commands and have a single redirection affect all of them:

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

A subshell would have been overkill in this case, because we did not require a temporary environment. However, a subshell would also have worked.

Command groups are also useful to shorten certain common tasks:

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

Compare that with the more formal version:

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

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.

Command groups can also be used for setting variables in unusual cases:

```
$ echo "cat
> mouse
> dog" > inputfile
$ { read a; read b; read c; } < inputfile
$ echo "$b"
mouse</pre>
```

It would have been extremely difficult to read the second line from a file without a command group, which allowed multiple read commands to read from a single open FD without rewinding to the start each time. Contrast with this:

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

That's not what we wanted at all!

If the command group is on a single line, as we've shown here, then there *must* be a semicolon before the closing }; 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
```

+ +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
```

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. We can also do that inside \$((...)) in BASH, but not in the POSIX shell.

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>
```

+ † 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. 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:

```
$ 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:

```
Toggle line numbers

1 #!/bin/bash
2 sum() {
3 echo "$1 + $2 = $(($1 + $2))"
4 }
5 sum 5 7
```

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:

```
Toggle line numbers
   1 #!/bin/bash
   2 open() {
         case "$1" in
            *.mp3|*.ogg|*.wav|*.flac|*.wma) xmms "$1";;
   4
             *.jpg|*.gif|*.png|*.bmp) display "$1";;
   5
             *.avi | *.mpg | *.mp4 | *.wmv)
                                              mplayer "$1";;
   6
  7
         esac
  8 }
  9 for file; do
 10
         open "$file"
 11 done
```

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,

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

The local variable i 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!*

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 "$0"; }
```

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 1s 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.

+ ↑6. Destroying Constructs

To remove a function or variable from your current shell environment use the *unset* command.

```
$ unset myfunction
```

In the manual: ...

In the FAQ: ...

- <- Input and Output | Sourcing ->
- <- Compound Commands | Job Control ->

++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 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)!

What you can do is to *source* the script, instead of running it as a child. You can do this using the . (dot) command:

This is often called *dotting in* a script. The . tells BASH 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.

- <- Compound Commands | Job Control ->
- <- Sourcing | Practices ->

++,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.

+ † 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 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)

→ ↑2. Practice

Job control can be turned on 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.

+ +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.

- <- Sourcing | Practices ->
- <- Job Control

++Practices

Contents

- 1. Choose Your Shell
- 2. Quoting
- 3. Readability
- 4. Bash Tests
- 5. Don't Ever Do These
- 6. Debugging

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.

BASH may be easy to learn and write in, but it isn't always fit for the job.

There are a lot of tools in the basic toolset that can help you. If you just need AWK, then don't make a shell script that invokes AWK. Just make an AWK script. If you need to retrieve data from an HTML or XML file in a reliable manner, Bash is also **the wrong tool for the job**. You should consider XPath/XSLT instead, or a language that has a library available for parsing XML or HTML.

If you decide that a shell script is what you want, then first ask yourself these questions:

- In the foreseeable future, might your script be needed in an environment where Bash is not available *by default*?
 - If so, **then consider sh instead.** sh is a POSIX shell and its features are available in any shell that complies with the POSIX standad. You can rely on the fact that any POSIX system will be able to run your script. You will have to balance the needs of portability against the Bash features you'd be unable to use.
 - (Keep in mind that this guide does not apply to sh!)
- Are you certain that in all environments you will run and might want to run this script in the future Bash 3.x (or 4.x) will be available to you?
 - If not, you should limit yourself to features of Bash 2.x ONLY.

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 3 or higher means you can avoid ancient scripting techniques. They have been replaced with far better alternatives for *very good reasons*.

- Stay away from scripting examples you see on the Web unless you understand them completely. Mostly any scripts you will find on the web are broken in some way. Do not copy/paste from them.
- If you're writing a Bash script, you **need** to put #!/usr/bin/env bash at the top of your script. Omitting this header or using the #!/bin/sh for header is **wrong**. In the latter case you will no longer be able to use any Bash features. You will be limited to the POSIX shell scripting standard (even if your /bin/sh links to Bash).
- When writing Bash scripts, do **not** use the [command. Bash has a far better alternative: [[.Bash's [[is more reliable in many ways and there are no advantages whatsoever to sticking with the old relic. If you do not heed this advice, you will find your scripts will break unless you are **very** careful with quotes. You'll also miss out on a lot of features provided by [[that [does not have (like pattern matching).
- It's also time you forgot about `...`. It isn't consistent with the syntax of *Expansion* and is terribly hard to nest without a dose of painkillers. Use \$ (...) instead.
- And for heaven's sake, "Use more quotes!" Protect your strings and parameter expansions from WordSplitting. Word splitting will eat your babies if you don't quote things properly.
- 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 math instead of expr to do simple calculations, *especially* when just incrementing a variable. If the script you're reading uses x=`expr \$x + 1` then it's not something you want to mimic.

++2. Quoting

Word Splitting is the demon inside BASH that is out to get unsuspecting newcomers or any other scripters off guard.

If you do not understand how *Word Splitting* works or when it is applied you should be very careful with your strings and your *Parameter Expansions*. I suggest you read up on WordSplitting if you doubt your knowledge.

The best way to protect yourself from this beast is to quote all your strings. Quotes keep your strings in one piece and prevent *Word Splitting* from tearing them open. Allow me to illustrate:

```
$ echo Push that word away from me.
Push that word away from me.
$ echo "Push that word away from me."
Push that word away from me.
```

Now, don't think *Word Splitting* is about collapsing spaces. What really happened in this example is that the first command passed each word in our sentence as a separate argument to echo. BASH split our sentence up in words using the whitespace between them to determine where each argument begins and ends. In the second example BASH is forced to keep the whole quoted string together. This means it's not split up into arguments and the whole string is passed to echo as the first argument. echo prints out each argument it gets with a space in between them. You should understand the basics of *Word Splitting* now.

This is where it gets dangerous! *Word Splitting* does not just happen on literal strings, it also happens **after** *Parameter Expansion*! As a result, on a dull and tired day, you might just be stupid enough to make 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.
```

As you can see, in the first echo command, we were negligent and left out the quotes. That was a mistake. BASH expanded our sentence and then used *Word Splitting* to split the resulting expansion up into arguments to use for echo. 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 whitespace.

It's not just spaces you need to protect. *Word Splitting* occurs on spaces, tabs and newlines, or whatever characters are in the IFS variable. Here's another example to show you just how badly you can break things if you neglect to use quotes:

```
$ echo "$(ls -al)"
   total 8
   drwxr-xr-x 4 lhunath users 1 2007-06-28 13:13 "."/
   drwxr-xr-x 102 lhunath users 9 2007-06-28 13:13 ".."/
   -rw-r--r-- 1 lhunath users 0 2007-06-28 13:13 "a"
   -rw-r--r-- 1 lhunath users 0 2007-06-28 13:13 "b"
                1 lhunath users 0 2007-06-28 13:13 "c"
   drwxr-xr-x 2 lhunath users 1 2007-06-28 13:13 "d"/
   drwxr-xr-x 2 lhunath users 1 2007-06-28 13:13 "e"/
   $ echo $(ls -al)
   total 8 drwxr-xr-x 4 lhunath users 1 2007-06-28 13:13 "."/
drwxr-xr-x 102 lhunath users 9 2007-06-28 13:13 ".."/ -rw-r--r-- 1
lhunath users 0 2007-06-28 13:13 "a" -rw-r--r-- 1 lhunath users 0
2007-06-28 13:13 "b" -rw-r--r- 1 lhunath users 0 2007-06-28 13:13
"c" drwxr-xr-x 2 lhunath users 1 2007-06-28 13:13 "d"/ drwxr-xr-x 2
lhunath users 1 2007-06-28 13:13 "e"/
```

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"
$ for friend in $friends
> do echo "$friend is my friend!"; done
Marcus is my friend!
JJ is my friend!
Thomas 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.

++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% ready, 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, believe me when I say you'd rather see this:

```
Toggle line numbers
         friends=( "Marcus The Rich" "JJ The Short" "Timid Thomas"
"Michelangelo The Mobster" )
         # Say something significant about my friends.
         for name in "${friends[@]}"; do
   4
   6
             # My first friend (in the list).
   7
             if [[ $name = ${friends[0]} ]]; then
                 echo $name was my first friend.
   8
  9
  10
             # My friends those names start with M.
             elif [[ $name = M* ]]; then
 11
                 echo "$name starts with an M"
 12
 13
             # My short friends.
 14
             elif [[ " $name " = *" Short "* ]]; then
 15
                 echo "$name is a shorty."
 16
 17
             # Friends I kind of didn't bother to remember.
 18
 2.0
                 echo "I kind of forgot what $name is like."
  21
             fi
 22
 23
         done
```

Than be confronted with something like this:

```
Toggle line numbers
         x = (
                    Marcus\ The\ Rich JJ\ The\ Short
           Timid\ Thomas Michelangelo\ The\ Mobster)
         for name in "\{x[@]\}"
   3
           do if [ "$name" = "$x" ]; then echo $name was my first
   4
friend.
   5
          elif
   6
           echo $name
                           7
           grep -qw Short
             then echo $name is a shorty.
          elif [ "x${name:0:1}" = "xM" ]
    then echo $name starts with an M; else
  9
  10
         echo I kind of forgot what name \
  11
  12
          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. Whatever your partner may say, it's good to be predictable and transparent.

++4. Bash Tests

The test application, 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 variables. In a number of shells, including bash, [is implemented as a shell builtin.

It can produce surprising results, especially for people starting shell scripting that consider [] as 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 newer? keywords that, though [[still behaves in many ways like a command, present several advantages over the traditional test 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 sees, is the equals sign. It takes this to be the left hand string of the test. The next thing test sees is the ''. It expects an operator to perform some kind of test on = and explodes.

Yes, test did not see our empty \$var because BASH expanded it into nothingness before test could even see it. Morale 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! [[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 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. (Furthermore, the proper operator would have been -lt instead of < anyway.) BASH is instructed to open a file 'a' and connect it to stdin for reading. To prevent this, we need to escape < so that test receives the operator rather than BASH. This makes our second attempt work.

Using [[we can avoid the mess altogether. [[sees the < operator before BASH gets to use it for *Redirection*; 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. 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, great. Trust me when I say you can always 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.

So believe me when I say, [[is safer than [. Because everybody inevitably makes programming errors. People usually don't intend to introduce bugs in their code. It just happens. So don't pretend you can use [and "You'll be careful not to make these mistakes", because I can assure you that you will.

Besides [[provides the following features over [:

```
• [[ can do glob pattern matching:
```

```
[[abc = a*]]
```

• [[can do regex pattern matching (Bash 3.1+):

```
[[abb =~ ab+]]
```

The only advantage of test is its portability.

++5. Don't Ever Do These

Of course use your common sense, if you know the data set you are working with and how it is produced then using these methods can be 100% reliable, this is for when you don't know exactly what data set you are working with, generally when you are writing code for other people.

The Bash shell allows you to do quite a lot of things, offering you considerable flexibility. Unfortunately, it does very little to discourage misuse 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 end up in production environments or in Linux distributions. The result of these, but even your very own scripts written in a time of neglect, can very 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 of ls! The ls command's output cannot be trusted for a number of reasons.

- For one, 1s will mangle the filenames of files if they contain characters unsupported by your locale. As a result, parsing filenames out of 1s is NEVER guaranteed to actually give you a file that you will be able to find. 1s 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 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; then ls *.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. In my 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 it

The alternative is called globbing. Bash has a feature called *Pathname Expansion*. This will help you enumerate all files that match a certain pattern! Alternatively you can use globs to test whether a certain filename matches a glob pattern.

• cat file | grep pattern

anyway.

Don't use cat as a default method for feeding file content to a process. 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 process

(grep 'pattern' /my/file,
sed 'expression' /my/file,...).

If the manual of the process does not specify any way to do this, you should use redirection (read column1 column2 < /my/file, tr ' '\n' < /my/file,...).

• 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 to do some counting and then pass that application's output to Bash for word splitting. Learn the syntax of for already! You should use this in Bash 3.x: for number in $\{1..10\}$, or this in Bash 2.x: for $((i=1; i\leq 10; i++))$.

• 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.

You should use this in Bash: let i++

++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.

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.

If you must, try and formulate the problem as a sentence. This will also be vital if you're going to ask for help with other people about your problem. You don't want them to rewrite your whole script. You also 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*.

Minimize The Codebase

If staring at your code doesn't give you a divine intervention, the next thing you should do is try to minimize your codebase while maintaining the same 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 by copying 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.

As soon as you delete something that makes the problem go away, you'll have isolated the problem. If not, at least you're not staring at a massive script anymore, but hopefully at a stub of no more than 5-7 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 in it:

```
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 ran 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 1s into words. That's because 1s is

UNPARSABLE in a bugless manner (do not ever use ls in scripts, unless if you want to show its output to a user).

The fix would be:

```
find "$imgFolder" -print0 | while read -d $'\0' image; do
    echo "$image"
done
```

Now that you've fixed the problem in this tiny example, it's easy to merge it back into the original script.

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 x option turned on, it prints out every command it executes before executing it. This 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:

```
#! /bin/bash
[..irrelevant code..]
set -x
[..relevant code..]
set +x
[..irrelevant code..]
```

If you have a complicated mess of scripts, you might find it helpful to change PS4 before setting -x:

```
export PS4='+$BASH_SOURCE:$LINENO:$FUNCNAME: '
```

```
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.

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:

 $tr -d '\r' < myscript > tmp && mv tmp myscript$

Read the FAQ / Pitfalls

The BashFAQ and BashPitfalls 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.

Ask Us On IRC

There are people in the #bash channel almost 24/7. This channel resides on the freenode IRC network. To reach us, you need an IRC client. Connect it to irc.freenode.net, 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.

<- Job Control

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