

When answering Linux command questions on this side or the back side of this page, refer to the following Inverted Tree diagram. The **linux** directory is contained in your home directory. Assume that you just logged into your **Matrix** account. Directories are underlined.

```
linux
|-- content
|   |-- assignments
|   |-- tests
|       |-- .answers.txt
|       |-- questions.txt
|-- projects
```

Questions:

1. Write a single Linux command to provide a detailed listing of all files in the /bin directory, sending the output to a file called listing.txt in the “projects” directory.
(append output to existing file and use a relative pathname)

```
ls -l /bin >> projects/listing.txt
```

2. Write a single Linux command to redirect the stderr from the command:
cat a.txt b.txt c.txt to a file called error.txt contained in the “assignments” directory.
(overwrite previous file’s contents and use only relative pathnames)

```
cat a.txt b.txt c.txt 2> assignments/error.txt
```

3. Write a single Linux command: cat ~/a.txt ~/b.txt ~/c.txt and redirect stdout to a file called “good.txt” to the “tests” directory and stderr to a file called “bad.txt” to the “tests” directory.
(overwrite previous contents for both files and use only relative-to-home pathnames)

```
cat ~/a.txt ~/b.txt ~/c.txt > tests/good.txt 2> tests/bad.txt
```

4. Write a single Linux command to redirect the stdout from the command:
cat a.txt b.txt c.txt to a file called wrong.txt contained in the “projects” directory and throw-out any standard error messages so they don’t appear on the screen.
(append output to existing file and use only relative pathnames)

```
cat a.txt b.txt c.txt >> projects/wrong.txt
```

5. Write a single Linux pipeline command to display a detailed listing of the “projects” directory but pause one screen at a time to view and navigate through all of the directory contents.

Use a relative-to-home pathname.

```
ls -l ~/projects | more
```

6. Write a single Linux pipeline command to display the sorted contents (in reverse alphabetical order) of the “linux” directory. Use a relative pathname.

```
ls linux | sort -r
```

7. Assume that the text file called “.answers.txt” contains 10 lines. Write a single Linux pipeline command to only displays lines 5 through 8 for this file. Use only relative pathnames.

```
1
2
3
4
5
6
7
8
9
10
```

```
head -8 linux/content/tests/.answers.txt | tail -4
```

8. Write a single Linux pipeline command to only display the contents of the “assignments” directory whose filenames match the pattern “murray” (both upper or lowercase). Use an absolute pathname.

```
ls /home/murray.saul/linux/content/assignments | grep -i murray
```

9. Write a single Linux pipeline command to display the number of characters contained in the file called “.answers.txt”. Use a relative-to-home pathname.

```
cat ~/murray.saul/linux/content/tests/.answers.txt | wc -c
```

10. Write a single Linux pipeline command to display the number of lines contained in the file called “questions.txt”. Use a relative pathname.

```
cat linux/content/tests/questions.txt | wc -l
```

11. Write a single Linux pipeline command to display only the first 10 characters of each filename contained in your current directory. Also, there is will be a lot of output, so also pause at each screenful so you can navigate throughout the display contents. Use a relative pathname.
- ```
ls | cut -c1-10 | more
```