

<b>cat -n</b>	display lines with line numbers (all lines are numbered)
<b>head -n N</b>	output the first N lines of each file
<b>tail -n N</b>	output the last N lines of each file
<b>tail -n +N</b>	output the end of each file beginning with line N (from the beginning)
<b>grep "pattern"</b>	output lines that contain pattern (no options) <b>-i</b> - contain pattern, ignoring case <b>-v</b> - dont contain pattern <b>-c</b> - count the number of lines that contain pattern <b>-l</b> - output the name of the files that contain at least one match
<b>sort</b>	sort lines. default: alphabetic, smallest to largest; use -n for numeric sort; use -r for reverse order (largest to smallest)
<b>type</b>	output information about where a command comes from (also useful, but more limited in bash: whereis and which)
<b>more</b>	display files page by page. Commands to control display: space - forward one page enter - forward one line ^B - back one page / <b>pattern</b> - search forward to next instance of pattern <b>q</b> quit
<b>ls</b>	<b>-L</b> - follow symlinks <b>-lu</b> long format sorted by modification date <b>-i</b> - show inode numbers <b>-d</b> - list directories themselves not contents
<b>ln orig new</b>	create a second link (named <b>new</b> ) to the file that <b>orig</b> refers to default - create hard link Use <b>-s</b> to create symbolic link
<b>man</b>	get information from the Unix manual <b>man x</b> output the first manual page found that is named <b>x</b> <b>man -k yy</b> output the names of pages whose terse description contains <b>yy</b> <b>man N yy</b> limit the search to section N of the Unix manual
<b>find dir [opts]</b>	output the path to each thing beneath directory <b>dir</b> that matches the options opts: <b>-type x</b> find objects of type <b>x</b> , where <b>x</b> is <b>f</b> (file), <b>d</b> (dir), <b>l</b> (symlink) <b>-name 'pat'</b> whose name matches the given wildcard pattern <b>pat</b> ( <b>pat</b> must be quoted) <b>-L</b> If a symlink is found, follow it and examine what it refers to. (needs other options (such as <b>-type f</b> ) to be interesting.)
<b>locate "pattern"</b>	<b>locate</b> searches a database of all files on the system looking for patterns in filenames. The database is updated daily (usually). <b>pattern</b> is a wildcard (globbing) pattern. If the pattern does not contain wildcards, leading and trailing asterisks are assumed ( <b>*pattern*</b> ) If the option <b>--regex</b> is used, <b>pattern</b> is basic regex If the option <b>--regext</b> is used, <b>pattern</b> is extended regex
<b>tee [-a] file</b>	Duplicate data going to standard output. Data still goes to standard output, but it is also copied to <b>file</b> . <b>grep 'abc' foo   tee abcs</b> would display the matches on stdout <i>and</i> save them in the file <b>abcs</b> <b>-a</b> : append <b>file</b> instead of overwriting it.
<b>uniq</b>	remove adjacent duplicate lines. Unique lines are sent to stdout.

**sort***general options*

- t****x** use **x** as delimiter. Whitespace is the default.
- f** fold (ignore) case
- u** (unique) - remove lines with duplicate keys
- o** **outfile** put the output file in **outfile** instead of stdout

*sort keys*

-**k****start** [**,****stop**]      **start** and **stop** are **f** [**.****c**]

the modifiers **r** (reverse) and **n** (numeric) can be appended to the key specifier

*Interpretation of keys. Note that [bracketed parts] are optional*

- the first character of the key is at field **f** and character offset **c** of **start**
- the last character of the key is at field **f** and character offset **c** of **stop**.
- If **.c** is missing on start, it means the first character of the field.
- If **.c** is missing on stop, it means the last character of the field.
- If **,stop** is missing, it means the last field of the record.
- Remaining ties are broken by sorting the unsorted fields alphabetically starting with field 1

**Examples**

Given a colon-delimited file **data** of four fields

**sort -t: -k2,2n data**

the key is field 2 only, numeric. Ties are broken by sorting the ties on field 1 (alphabetic) then on field 3 (alphabetic) then on field 4 (alphabetic)

**sort -t: -k2n data**

the key is field 2 numeric, but this key also includes fields 3 and 4 (numeric) to break ties. If any remaining ties exist (two records which are identical in fields 2-4), field 1 is sorted (alphabetic) to break them.

**sort -t: -k4 -u data**

the key is field 4, which is sorted alphabetically. Then the rest of the line is sorted to break any remaining ties. Since **-u** is used, only the first record that has a particular value in field 4 is kept - records with duplicate field 4's are deleted.

**sort -t: -f -k2,2n -k4 data**

the first key is field 2 numeric. Ties are broken using the second key, which is field 4 alphabetic. Remaining ties are broken by sorting on fields 1 (alphabetic) then on field 3 (alphabetic). All sorts ignore case.

**sort -t: -o data -k3.1,3.1n -k2r data**

the first key is a single character - the first character of field 3. It is a numeric sort, which is redundant for a single digit. Ties are broken by sorting alphabetically in reverse order by field 2, then by the rest of field 3, then by field 4 then by field 1.

The output of this sort is placed in a file named **data**, replacing the original file. This moves the responsibility of creating the output file to the **sort** command so that the input file can be read before the output file is created. The command below overwrites the input file before the **sort** command starts, destroying the file in the process and resulting in an empty file.

**sort -t: -k3.1,3.1n -k2r data > data # DONT DO THIS!!!!**