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UNIX pipes and redirection

Redirection within UNIX allows a user to control where input comes from and output goes to. Pipes allow a user to pass the output from one command directly to the input of another command.

Redirection

>

The > symbol redirects the output from standard out (This is what you see on the screen) to a file. **Beware** if the file already exist, it will be erased and overwritten without warning, so be careful.

An example of this is for the commonly used mapping tool bwa. By default bwa mem will print the output to the screen. However you can write this to a file using >

```
bwa mem reference.fasta sample1 R1.fastq.gz sample1 R2.fastq.gz > sample1.sam
```

This command writes the output from bwa mem (in SAM format) to a file called sample1.sam

>>

The >> symbol redirects the output from standard out to a file but rather than overwriting the file, appends the output to the existing file. If the file does not exist it will just create a new one

<

The < symbol takes input from a file. An example below using the sickle trimming software that does not take gzipped fastqs is as follows

```
sickle se -f <(gunzip -c sample1 R1.fastq.gz) -t sanger -o trimmed.fa
```

Here gunzip -c writes the unzipped output to the STDOUT and rather than write to disc we convert it on the fly and read it into sickle

Piping |

The | symbol passed the out from one command onto another for further processing. An example using bwa mem is

```
bwa mem reference.fasta sample1_R1.fastq.gz sample1_R2.fastq.gz | samtools view
-bS -F 4 | samtools sort -o sample1.sorted.bam
```

Here rather than running bwa mem, followed by samtools view to convert to a bam file keeping only mapped reads (-F 4) and then sorting, these can be joined together through the pipe command into a single line. This has the advantage that the intermediate file are not written to disc.