

Mapped cancer reads to human genome

Downloaded the human genome from NCBI (see reference directory)

Mapped Dr Normal's reads to it with **bwa** and **samtools**:

```
bwa mem reference/GCA_000001405.15_GRCh38_genomic.fna
data/glioblastoma.fastq | samtools view -Sb - >
results/glioblastoma_map_GRCh38.bam
```