

Mapping glioblastoma Illumina sequences to the human genome sequence GRCh38.

Downloaded the human genome from

(https://ftp.ncbi.nlm.nih.gov/genomes/all/GCA/000/001/405/GCA_000001405.15_GRCh38/) and put it in the reference directory.

Dr Normal gave me the sequencing reads in `glioblastoma.fastq`. See the data directory.

Used `bwa` (v0.7.17) and `samtools` to map the reads:

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