Site ENA : <https://www.ebi.ac.uk/ena/browser/view/PRJEB24932>

Télécharger doc ENA : <https://ena-docs.readthedocs.io/en/latest/retrieval/file-download.html>

Aide commande linux : <https://github.com/LeCoupa/awesome-cheatsheets/blob/master/languages/bash.sh>

Mapping BWA: <https://github.com/lh3/bwa> + <http://bio-bwa.sourceforge.net/>

Génome de référence: <http://sgd-archive.yeastgenome.org/sequence/S288C_reference/genome_releases/>

GATK: <https://gatk.broadinstitute.org/hc/en-us/categories/360002302312>

Samtools: <http://www.htslib.org/>

**BWA :**

git clone https://github.com/lh3/bwa.git

cd bwa; make

./bwa index ref.fa

./bwa mem ref.fa read-se.fq.gz | gzip -3 > aln-se.sam.gz

./bwa mem ref.fa read1.fq read2.fq | gzip -3 > aln-pe.sam.gz

Autre :

https://www.southgreen.fr/sites/southgreen.fr/files/TP-LINUX-NGS-Mapping\_0.pdf

Sam → Bam : page 5



