Alternative ways to fetch sequence data

- Kingfisher
- sra-tools (fastq-dump/fasterq-dump/prefetch)
- ena-ftp-downloader
- ENA download script
- Globus (web/command line)
- wget/curl/FTP client/Aspera from FTP site

```
wget ftp://ftp.sra.ebi.ac.uk/vol1/fastq/ERR164/ERR164407/ERR164407.fastq.gz
```

 Various tools that are wrappers for some or several of the above methods