### F25A430000757\_HOMukwhX1

**Project Information**

The basic information of the project is shown below:

* Project ID: F25A430000757\_HOMukwhX
* Product name: filter - DNBSEQ whole exome sequencing
* Sample size: 6
* Library type: Whole Exome library
* Sequencing Platform: DNBSEQ
* Sequencing read Length: PE100
* Clean fastq phred quality score encoding: Phred+33

**Normal Reference of Kuromochi downloaded from GEO database**

Whole Exome Sequencing of Kuromochi Cell Line from a study

https://trace.ncbi.nlm.nih.gov/Traces/?view=run\_browser&acc=SRR26823042&display=metadata

**Normal Reference of OAW28 is download from GEO database**

Targeted Sequencing of OAW28 from DepMap

https://trace.ncbi.nlm.nih.gov/Traces/?view=run\_browser&acc=SRR8658026&display=metadata