Genomic data collected from the curated set of non-redundant studies on cBioPortal
159 studies, 42177 samples

Filtered genomic data for alteration frequency analysis
150 studies, 32568 patients, 39 cancer types

Data assembly
- Remove samples without mutations data
- Remove samples not profiled for TET1 mutation
- Assemble duplicate samples for the same patient
- Unify cancer type annotation