

# HOW TO ACCESS PROTECTED TARGET DATA



## Obtain dbGaP Authorized Access

URL: <https://dbgap.ncbi.nlm.nih.gov/aa/wga.cgi?login=&page=login>

For Assistance: <https://dbgap.ncbi.nlm.nih.gov/aa/wga.cgi?page=email&from=login>

### TARGET Data at NCI – All data except raw sequence data

- Clinical information
- All levels chip-based molecular characterization data
- Processed sequencing data

### TARGET Data at NCBI Archives – Raw Sequence Data

**Sanger**  
(Targeted re-sequencing, kinome seq)

**2<sup>nd</sup>/3<sup>rd</sup> Generation**  
Whole genome, exome, RNA-Seq, miRNA-Seq

### Requires an NCI user account

#### TARGET Data Portal

Access data directly via the Data Matrix in the portal

Access Sanger sequencing linking tables

### Requires an eRA commons account

**Trace Archive**  
Access sequence trace files

**TARGET dbGaP Project Site**  
Access BAM sequencing files stored at the Sequence Read Archive (SRA)

Link to TARGET data at NCBI via the Data Matrix



TARGET Data Portal  
<http://target.cancer.gov/dataportal>

TARGET dbGaP Project Site  
[http://www.ncbi.nlm.nih.gov/projects/gap/cgi-bin/study.cgi?study\\_id=phs000218.v3.p1](http://www.ncbi.nlm.nih.gov/projects/gap/cgi-bin/study.cgi?study_id=phs000218.v3.p1)

NCBI Trace Archive  
<http://www.ncbi.nlm.nih.gov/Traces/trace.cgi?view=search>

For assistance with all TARGET Data:

NCI – [OCG@mail.nih.gov](mailto:OCG@mail.nih.gov)

NCBI – <https://dbgap.ncbi.nlm.nih.gov/aa/wga.cgi?page=email&from=login>