



- 1) User creates metadata file and place the genome files to upload as shown in the Overture Client examples
- 2) User calls the Overture Client to login
- 3) Overture Client logs in to Keycloak with user credentials to get an auth token
- 4) Overture Client Stores the auth token on the filesystem
- 5) User calls the Overture Client to batch create the analyses defined in the metadata file and batch upload the related genome files