

RefSeqMash 0.0.2

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Abstract

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Protocol

Step 1.

Upload your into the [Cyverse/iPlant Discovery Environment](#) either by logging in and choosing to upload or by using iRODS to 'iput' your FASTA files.

Step 2.

Open the "Apps" by clicking on the button on the left. Search for "refseq" to find the RefSeqMash-0.0.2 app.

Step 3.

Indicate a FASTA file or a directory of FASTA files for the "query." Click on "Run Analysis."