
HH-SEARCH

Final

HOW TO USE IT

- Location of HH-Database

- `/mnt/project/pp2_hhblits_db/pp2_hhm_db`

- Command

- `hhsearch -i <input-file> -d' <db-filename> -o <output-file>`

EXAMPLE USAGE

- `hhsearch -i testSequence -d /mnt/project/pp2_hhblits_db/pp2_hhm_db -o newOutfile.txt`
 - optional parameter for speed-up
 - `-cpu #of-cpus-to use`
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