

```
(base) x@aris: /demo/ncbi/ncbi-con/test$ more top.txt
1      18959      98.721 gi|10313991|ref|NC_002549.1|_Zaire ebolavirus isolate Ebola_virus/H.sapiens-to
2      18935      8.808 gi|302315369|ref|NC_014372.1|_Tai_Forest_ebolavirus_isolate_Tai_Forest_virus/H
3      18940      5.104 gi|302371213|ref|NC_014373.1|_Bundibugyo_ebolavirus_isolate_Bundibugyo_virus/H
4      18891      3.280 gi|22789222|ref|NC_004161.1|_Reston_ebolavirus_isolate_Reston_virus/M.fascicu
5      18875      2.375 gi|55770807|ref|NC_006432.1|_Sudan_ebolavirus_isolate_Sudan_virus/H.sapiens-to
6      1259013    1.022 gi|363539767|ref|NC_016072.1|_Megavirus_chiliensis,_complete_genome
7      1350       0.763 gi|146411809|ref|NC_009450.1|_Honeysuckle_yellow_vein_mosaic_beta-[Japan:Miyiz
8      3057       0.505 gi|20153367|ref|NC_003619.1|_Groundnut_bud_necrosis_virus_segment_S,_complete
9      144575     0.423 gi|12084983|ref|NC_002642.1|_Yaba-like_disease_virus,_complete_genome
10     3105       0.357 gi|1013949526|ref|NC_029800.1|_Iris_yellow_spot_virus_non-structural_protein_
11     59866     0.344 gi|939482425|ref|NC_004913.3|_Stx1_converting_phage_DNA,_complete_genome
12     9130     0.237 gi|985485914|ref|NC_029052.1|_Goose_dicistrovirus_isolate_UW1,_complete_genome
13     8432     0.191 gi|9635564|ref|NC_002468.1|_Cherry_necrotic_rusty_mottle_virus,_complete_genom
14     8911     0.177 gi|20153365|ref|NC_003614.1|_Peanut_bud_necrosis_virus_L_protein_(L)_gene,_cor
15     47021     0.163 gi|148609382|ref|NC_009514.1|_Phage_cdtI_DNA,_complete_genome
```



ID



SIZE



% OF
SIMILARITY



HEADER (GI,NAME)