Center for Genomic Epidemiology

Input Files: *Vibrio_unknown_sample.fasta.txt*

Antibiotic(s): *Database*

Database

Fasta header	Identity	Query / Template length	Contig	Position in contig
7thVC_VC2346_AE003852	100	687 / 687	NODE_10_length_341876_cov_21.186171	172807..173493
ace_AE003852	100	294 / 294	NODE_29_length_4900_cov_21.245306	2625..2918
ctxA_CP001235	100	777 / 777	NODE_29_length_4900_cov_21.245306	554..1330
ctxBet_AE003852	100	375 / 375	NODE_29_length_4900_cov_21.245306	183..557
hlyA_HQ452880	100	217 / 217	NODE_13_length_637077_cov_20.154480	573780..573996
mshA_X77217	100	537 / 537	NODE_102_length_354682_cov_22.107241	28401..28937
ompU_KF434513	100	1026 / 1026	NODE_102_length_354682_cov_22.107241	267014..268039
ompW_FJ462451	100	416 / 416	NODE_13_length_637077_cov_20.154480	82735..83150
rfvBO1_AE003852	100	1233 / 1233	NODE_100_length_124700_cov_23.600674	58534..59766
rstRet_AE003852	100	339 / 339	NODE_36_length_2005_cov_60.292767	485..823
rtxA_AY101181	100	203 / 203	NODE_44_length_796052_cov_20.188923	410..612
tcpAet_AF325734	100	675 / 675	NODE_44_length_796052_cov_20.188923	672805..673479
toxR_HM042639	98.2	721 / 720	NODE_44_length_796052_cov_20.188923	516366..517086
zot_AE003852	100	1200 / 1200	NODE_29_length_4900_cov_21.245306	1429..2628

[extended output]

[Results as text] [Results tsv] [Hits in genome seqs] [Virulence factor seqs]