

ANiB value computed with PyANI using "anvi-compute-genome-similarity" from Anvi'o

	HBD_Gamma_02	<i>V. diazotrophicus</i> NS1
HBD_Gamma_02	1	0.687
<i>V. diazotrophicus</i> NS1	0.686	1

<i>V. diazotrophicus</i>	NS1	60.18M	60.27F	60.6B	60.6F	65.10M	65.7M	99A	HF9B
NS1	1	0.975	0.977	0.942	0.975	0.974	0.973	0.966	0.903
60.18M	0.975	1	0.977	0.94	0.975	0.974	0.972	0.965	0.904
60.27F	0.977	0.976	1	0.942	0.977	0.976	0.975	0.968	0.904
60.6B	0.942	0.94	0.942	1	0.942	0.94	0.94	0.953	0.898
60.6F	0.974	0.975	0.977	0.942	1	0.974	0.973	0.966	0.903
65.10M	0.974	0.973	0.975	0.939	0.974	1	0.974	0.974	0.903
65.7M	0.973	0.973	0.974	0.94	0.973	0.975	1	0.968	0.903
99A	0.965	0.965	0.967	0.952	0.966	0.974	0.966	1	0.902
HF9B	0.903	0.904	0.903	0.899	0.903	0.903	0.904	0.902	1