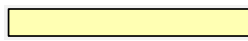


Function of Matched Query Gene



Modification



Precursor Synthesis

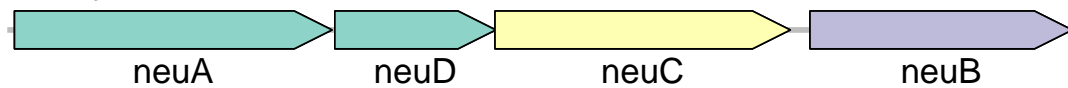


Synthase

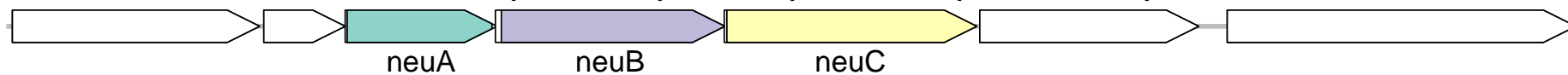


NA

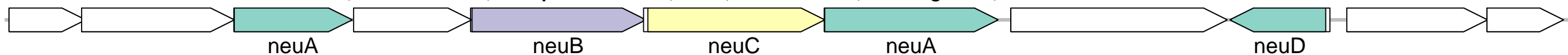
Query



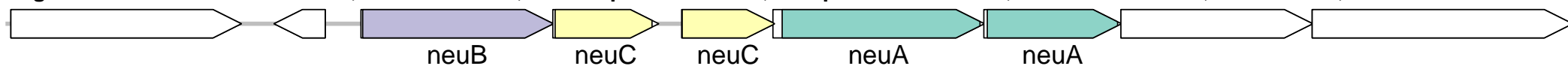
EsbW_18-Q3-R4-48_MAXAC.012, Nitrospirae, Nitrospira, Nitrospirales, Nitrospiraceae, *Nitrospira*, *defluvi*



EsbW_18-Q3-R4-48_MAXAC.279_cIn, Proteobacteria, Deltaproteobacteria, PB19, midas_f_2377, *midas_g_9396*, *midas_s_939*



Ega_18-Q3-R5-49_MAXAC.062, Proteobacteria, Gammaproteobacteria, Betaproteobacteriales, Burkholderiaceae, *Rhodoferrax*, *saidenbachensi*



0

2500

5000

7500