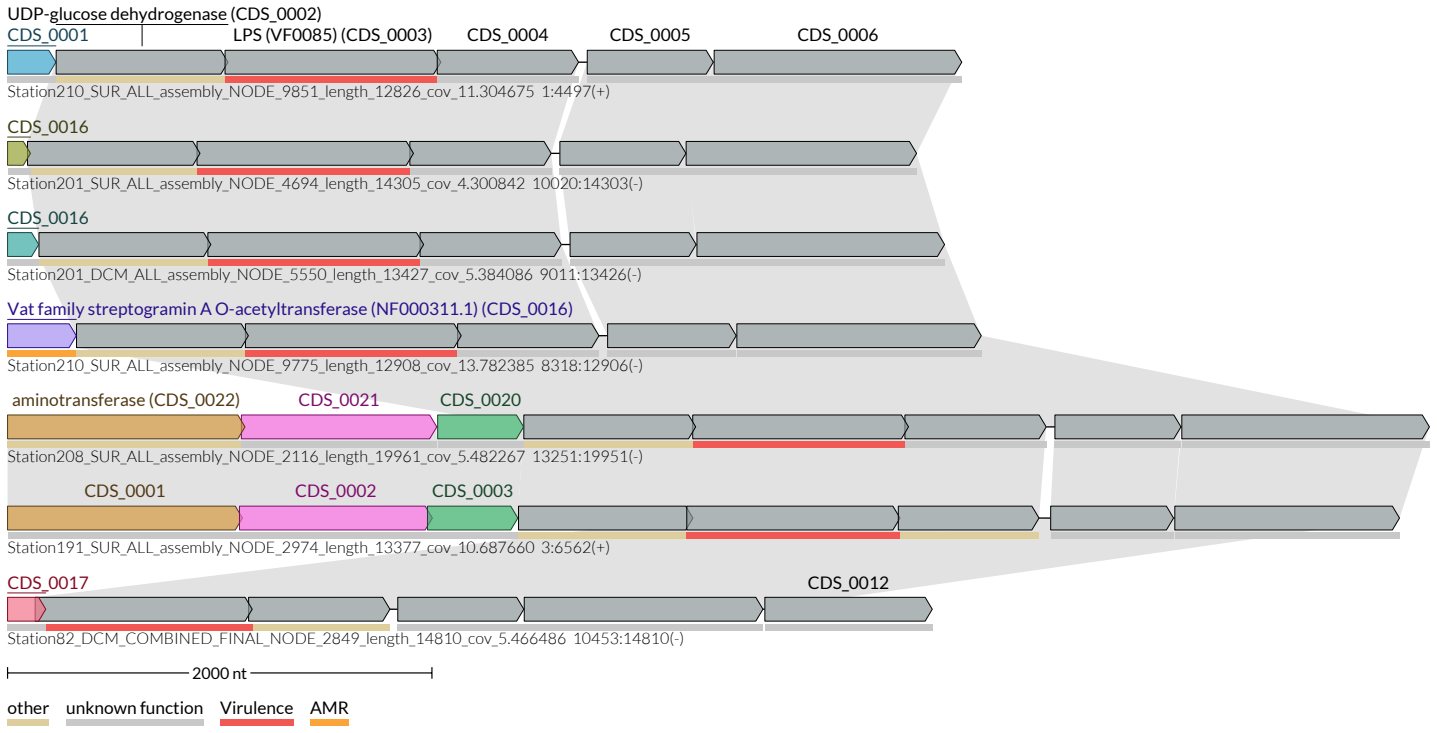
UDP-glucose dehydrogenase (CDS_0002)
CDS_0001
LPS (VF0085) (CDS_0003)
CDS_0004
CDS_0005
CDS_0006
Station210_SUR_ALL_assembly_NODE_9851_length_12826_cov_11.304675 1:4497(+)
CDS_0016
Station201_SUR_ALL_assembly_NODE_4694_length_14305_cov_4.300842 10020:14303(-)
CDS_0016
Station201_DCM_ALL_assembly_NODE_5550_length_13427_cov_5.384086 9011:13426(-)
Vat family streptogramin A O-acetyltransferase (NF000311.1) (CDS_0016)
Station210_SUR_ALL_assembly_NODE_9775_length_12908_cov_13.782385 8318:12906(-)
aminotransferase (CDS_0022)
CDS_0021
CDS_0020
Station208_SUR_ALL_assembly_NODE_2116_length_19961_cov_5.482267 13251:19951(-)
CDS_0001
CDS_0002
CDS_0003
Station191_SUR_ALL_assembly_NODE_2974_length_13377_cov_10.687660 3:6562(+)
CDS_0017
CDS_0012
Station82_DCM_COMBINED_FINAL_NODE_2849_length_14810_cov_5.466486 10453:14810(-)
2000 nt
other
unknown function
Virulence
AMR