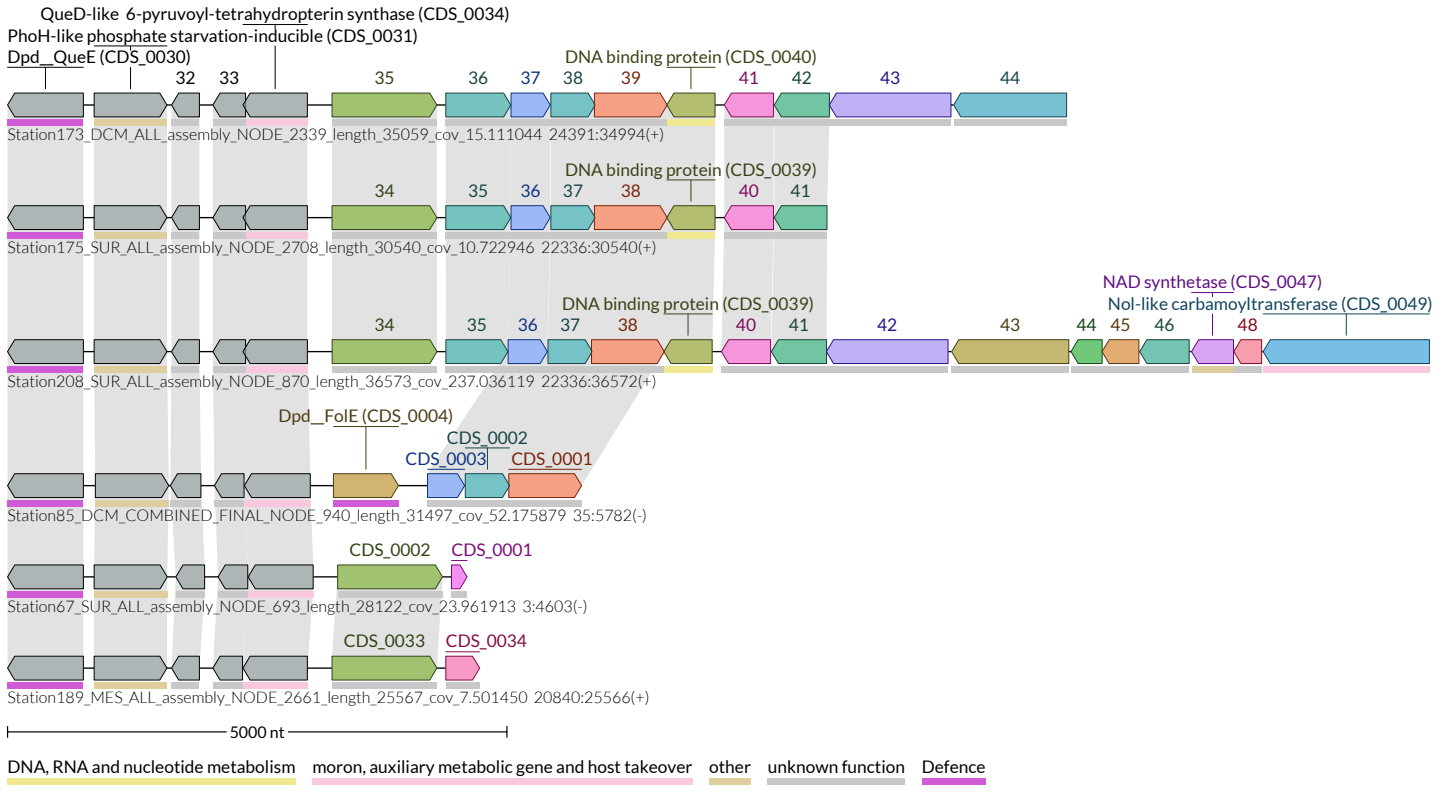
QueD-like 6-pyruvoyl-tetrahydropterin synthase (CDS_0034)
PhoH-like phosphate starvation-inducible (CDS_0031)
Dpd__QueE (CDS_0030)
32
33
35
36
37
38
39
DNA binding protein (CDS_0040)
41
42
43
44
Station173_DCM_ALL_assembly_NODE_2339_length_35059_cov_15.111044 24391:34994(+)
34
35
36
37
38
DNA binding protein (CDS_0039)
40
41
Station175_SUR_ALL_assembly_NODE_2708_length_30540_cov_10.722946 22336:30540(+)
NAD synthetase (CDS_0047)
Nol-like carbamoyltransferase (CDS_0049)
34
35
36
37
38
DNA binding protein (CDS_0039)
40
41
42
43
44
45
46
48
Station208_SUR_ALL_assembly_NODE_870_length_36573_cov_237.036119 22336:36572(+)
Dpd__FolE (CDS_0004)
CDS_0002
CDS_0003
CDS_0001
Station85_DCM_COMBINED_FINAL_NODE_940_length_31497_cov_52.175879 35:5782(-)
CDS_0002
CDS_0001
Station67_SUR_ALL_assembly_NODE_693_length_28122_cov_23.961913 3:4603(-)
CDS_0033
CDS_0034
Station189_MES_ALL_assembly_NODE_2661_length_25567_cov_7.501450 20840:25566(+)
5000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
unknown function
Defence