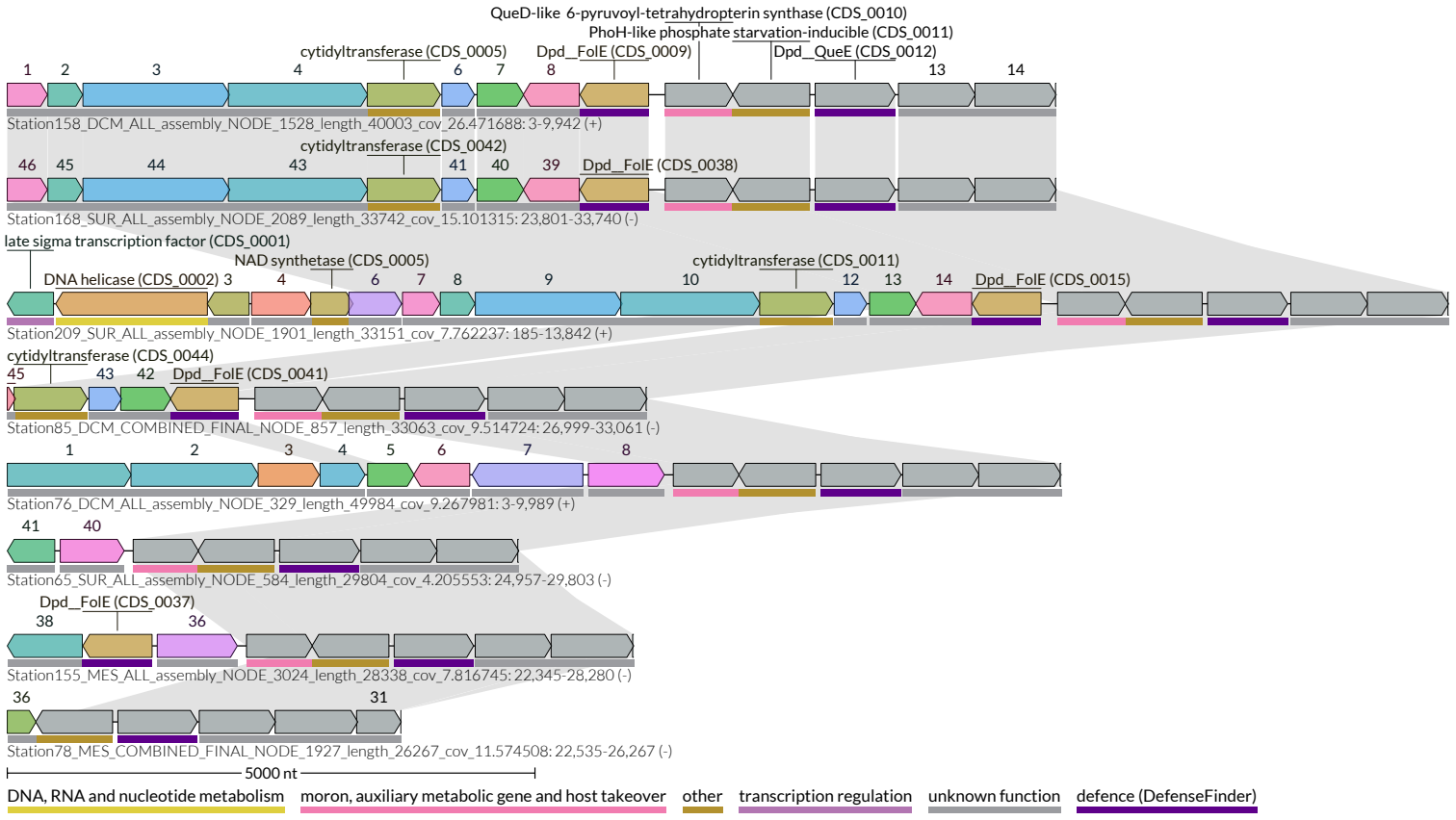
QueD-like 6-pyruvoyl-tetrahydropterin synthase (CDS_0010)
PhoH-like phosphate starvation-inducible (CDS_0011)
cytidyltransferase (CDS_0005)
Dpd__FolE (CDS_0009)
Dpd__QueE (CDS_0012)
1 2 3 4 6 7 8 13 14
Station158_DCM_ALL_assembly_NODE_1528_length_40003_cov_26.471688: 3-9,942 (+)
cytidyltransferase (CDS_0042)
46 45 44 43 41 40 39
Dpd__FolE (CDS_0038)
Station168_SUR_ALL_assembly_NODE_2089_length_33742_cov_15.101315: 23,801-33,740 (-)
late sigma transcription factor (CDS_0001)
NAD synthetase (CDS_0005)
cytidyltransferase (CDS_0011)
DNA helicase (CDS_0002) 3 4 6 7 8 9 10 12 13 14
Dpd__FolE (CDS_0015)
Station209_SUR_ALL_assembly_NODE_1901_length_33151_cov_7.762237: 185-13,842 (+)
cytidyltransferase (CDS_0044)
45 43 42
Dpd__FolE (CDS_0041)
Station85_DCM_COMBINED_FINAL_NODE_857_length_33063_cov_9.514724: 26,999-33,061 (-)
1 2 3 4 5 6 7 8
Station76_DCM_ALL_assembly_NODE_329_length_49984_cov_9.267981: 3-9,989 (+)
41 40
Station65_SUR_ALL_assembly_NODE_584_length_29804_cov_4.205553: 24,957-29,803 (-)
Dpd__FolE (CDS_0037)
38 36
Station155_MES_ALL_assembly_NODE_3024_length_28338_cov_7.816745: 22,345-28,280 (-)
36 31
Station78_MES_COMBINED_FINAL_NODE_1927_length_26267_cov_11.574508: 22,535-26,267 (-)
5000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
defence (DefenseFinder)