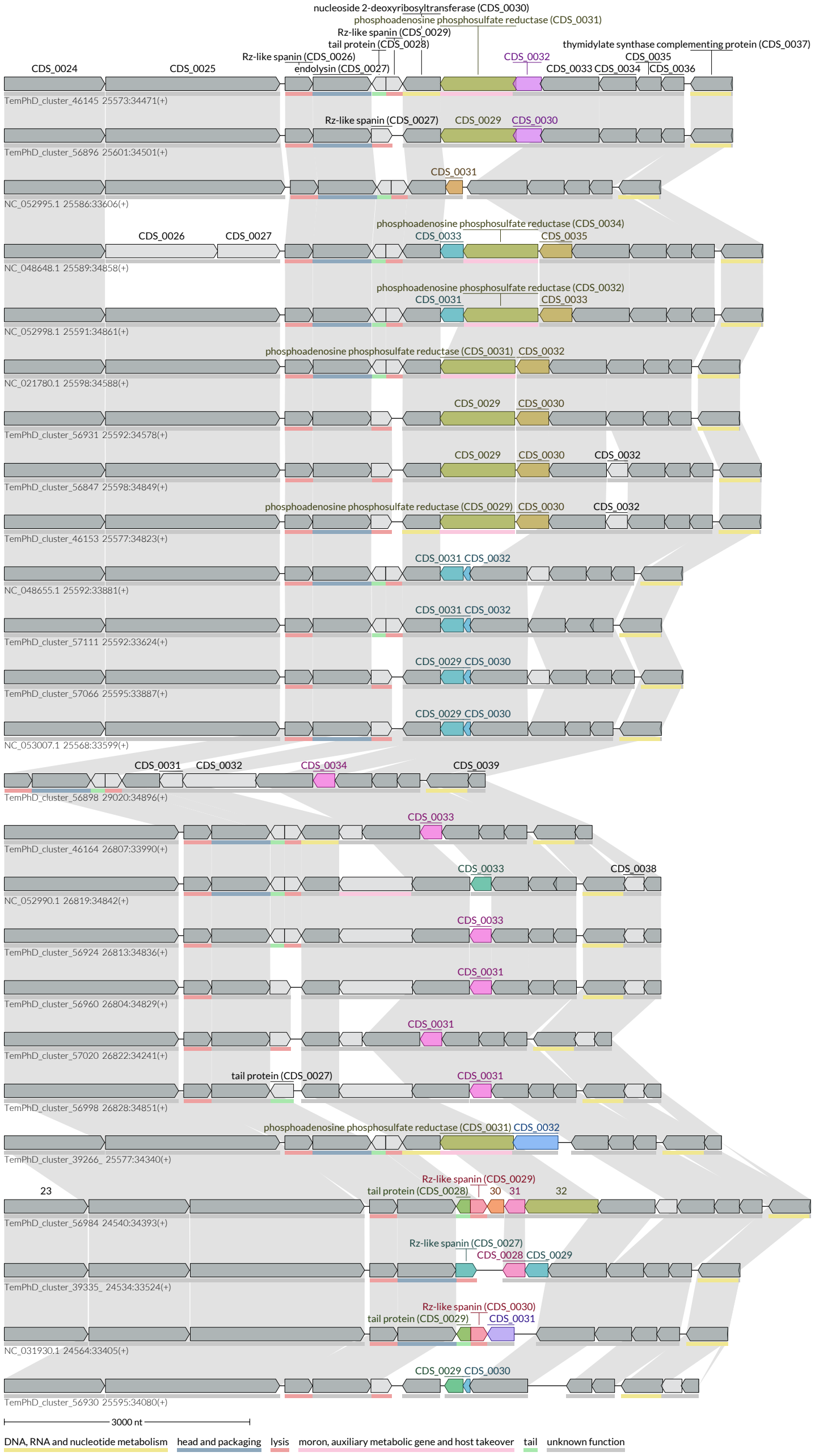
nucleoside 2-deoxyribosyltransferase (CDS_0030)
phosphoadenosine phosphosulfate reductase (CDS_0031)
Rz-like spanin (CDS_0029)
tail protein (CDS_0028)
thymidylate synthase complementing protein (CDS_0037)
Rz-like spanin (CDS_0026)
CDS_0032
CDS_0035
CDS_0024
CDS_0025
endolysin (CDS_0027)
CDS_0033 CDS_0034
CDS_0036
TemPhD_cluster_46145 25573:34471(+)
Rz-like spanin (CDS_0027)
CDS_0029
CDS_0030
TemPhD_cluster_56896 25601:34501(+)
CDS_0031
NC_052995.1 25586:33606(+)
phosphoadenosine phosphosulfate reductase (CDS_0034)
CDS_0026
CDS_0027
CDS_0033
CDS_0035
NC_048648.1 25589:34858(+)
phosphoadenosine phosphosulfate reductase (CDS_0032)
CDS_0031
CDS_0033
NC_052998.1 25591:34861(+)
phosphoadenosine phosphosulfate reductase (CDS_0031)
CDS_0032
NC_021780.1 25598:34588(+)
CDS_0029
CDS_0030
TemPhD_cluster_56931 25592:34578(+)
CDS_0029
CDS_0030
CDS_0032
TemPhD_cluster_56847 25598:34849(+)
phosphoadenosine phosphosulfate reductase (CDS_0029)
CDS_0030
CDS_0032
TemPhD_cluster_46153 25577:34823(+)
CDS_0031 CDS_0032
NC_048655.1 25592:33881(+)
CDS_0031 CDS_0032
TemPhD_cluster_57111 25592:33624(+)
CDS_0029 CDS_0030
TemPhD_cluster_57066 25595:33887(+)
CDS_0029 CDS_0030
NC_053007.1 25568:33599(+)
CDS_0031
CDS_0032
CDS_0034
CDS_0039
TemPhD_cluster_56898 29020:34896(+)
CDS_0033
TemPhD_cluster_46164 26807:33990(+)
CDS_0033
CDS_0038
NC_052990.1 26819:34842(+)
CDS_0033
TemPhD_cluster_56924 26813:34836(+)
CDS_0031
TemPhD_cluster_56960 26804:34829(+)
CDS_0031
TemPhD_cluster_57020 26822:34241(+)
tail protein (CDS_0027)
CDS_0031
TemPhD_cluster_56998 26828:34851(+)
phosphoadenosine phosphosulfate reductase (CDS_0031)
CDS_0032
TemPhD_cluster_39266_ 25577:34340(+)
Rz-like spanin (CDS_0029)
23
tail protein (CDS_0028)
30
31
32
TemPhD_cluster_56984 24540:34393(+)
Rz-like spanin (CDS_0027)
CDS_0028
CDS_0029
TemPhD_cluster_39335_ 24534:33524(+)
Rz-like spanin (CDS_0030)
tail protein (CDS_0029)
CDS_0031
NC_031930.1 24564:33405(+)
CDS_0029 CDS_0030
TemPhD_cluster_56930 25595:34080(+)
3000 nt
DNA, RNA and nucleotide metabolism
head and packaging
lysis
moron, auxiliary metabolic gene and host takeover
tail
unknown function