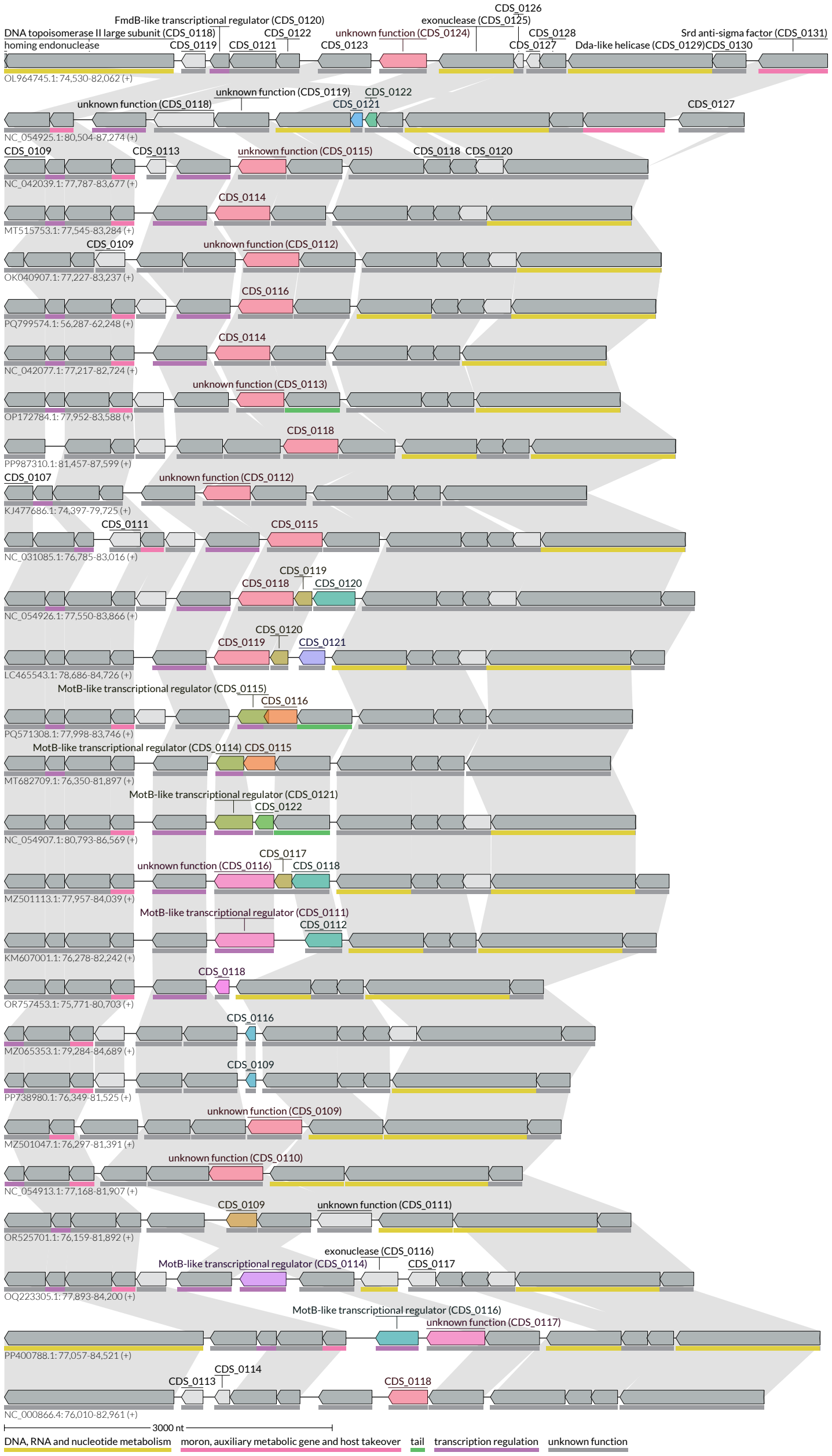
DNA topoisomerase II large subunit (CDS_0118)
homing endonuclease
FmdB-like transcriptional regulator (CDS_0120)
CDS_0119
CDS_0121
CDS_0122
CDS_0123
unknown function (CDS_0124)
exonuclease (CDS_0125)
CDS_0126
CDS_0127
CDS_0128
Dda-like helicase (CDS_0129)
CDS_0130
Srd anti-sigma factor (CDS_0131)
OL964745.1: 74,530-82,062 (+)
unknown function (CDS_0118)
unknown function (CDS_0119)
CDS_0121
CDS_0122
CDS_0127
NC_054925.1: 80,504-87,274 (+)
CDS_0109
CDS_0113
unknown function (CDS_0115)
CDS_0118
CDS_0120
NC_042039.1: 77,787-83,677 (+)
CDS_0114
MT515753.1: 77,545-83,284 (+)
CDS_0109
unknown function (CDS_0112)
OK040907.1: 77,227-83,237 (+)
CDS_0116
PQ799574.1: 56,287-62,248 (+)
CDS_0114
NC_042077.1: 77,217-82,724 (+)
unknown function (CDS_0113)
OP172784.1: 77,952-83,588 (+)
CDS_0118
PP987310.1: 81,457-87,599 (+)
CDS_0107
unknown function (CDS_0112)
KJ477686.1: 74,397-79,725 (+)
CDS_0111
CDS_0115
NC_031085.1: 76,785-83,016 (+)
CDS_0119
CDS_0118
CDS_0120
NC_054926.1: 77,550-83,866 (+)
CDS_0120
CDS_0119
CDS_0121
LC465543.1: 78,686-84,726 (+)
MotB-like transcriptional regulator (CDS_0115)
CDS_0116
PQ571308.1: 77,998-83,746 (+)
MotB-like transcriptional regulator (CDS_0114)
CDS_0115
MT682709.1: 76,350-81,897 (+)
MotB-like transcriptional regulator (CDS_0121)
CDS_0122
NC_054907.1: 80,793-86,569 (+)
CDS_0117
unknown function (CDS_0116)
CDS_0118
MZ501113.1: 77,957-84,039 (+)
MotB-like transcriptional regulator (CDS_0111)
CDS_0112
KM607001.1: 76,278-82,242 (+)
CDS_0118
OR757453.1: 75,771-80,703 (+)
CDS_0116
MZ065353.1: 79,284-84,689 (+)
CDS_0109
PP738980.1: 76,349-81,525 (+)
unknown function (CDS_0109)
MZ501047.1: 76,297-81,391 (+)
unknown function (CDS_0110)
NC_054913.1: 77,168-81,907 (+)
CDS_0109
unknown function (CDS_0111)
OR525701.1: 76,159-81,892 (+)
exonuclease (CDS_0116)
MotB-like transcriptional regulator (CDS_0114)
CDS_0117
OQ223305.1: 77,893-84,200 (+)
MotB-like transcriptional regulator (CDS_0116)
unknown function (CDS_0117)
PP400788.1: 77,057-84,521 (+)
CDS_0114
CDS_0113
CDS_0118
NC_000866.4: 76,010-82,961 (+)
3000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
tail
transcription regulation
unknown function