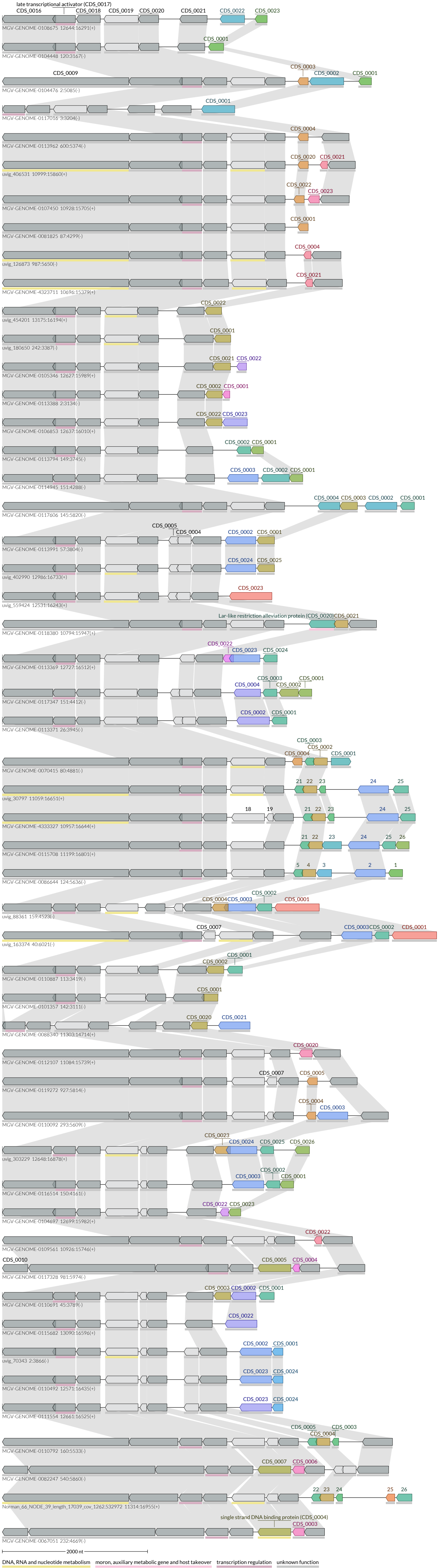
late transcriptional activator (CDS_0017)
CDS_0016
CDS_0018
CDS_0019
CDS_0020
CDS_0021
CDS_0022
CDS_0023
MGV-GENOME-0108675 12644:16291(+)
CDS_0001
MGV-GENOME-0104448 120:3167(-)
CDS_0009
CDS_0003
CDS_0002
CDS_0001
MGV-GENOME-0104476 2:5085(-)
CDS_0001
MGV-GENOME-0117016 3:3204(-)
CDS_0004
MGV-GENOME-0113962 600:5374(-)
CDS_0020
CDS_0021
uvig_406531 10999:15860(+)
CDS_0022
CDS_0023
MGV-GENOME-0107450 10928:15705(+)
CDS_0001
MGV-GENOME-0081825 87:4299(-)
CDS_0004
uvig_126873 987:5650(-)
CDS_0021
MGV-GENOME-4323711 10696:15379(+)
CDS_0022
uvig_454201 13175:16194(+)
CDS_0001
uvig_180650 242:3387(-)
CDS_0021
CDS_0022
MGV-GENOME-0105346 12627:15989(+)
CDS_0002
CDS_0001
MGV-GENOME-0113388 2:3134(-)
CDS_0022
CDS_0023
MGV-GENOME-0106853 12637:16010(+)
CDS_0002
CDS_0001
MGV-GENOME-0113794 149:3745(-)
CDS_0003
CDS_0002
CDS_0001
MGV-GENOME-0114945 151:4288(-)
CDS_0004
CDS_0003
CDS_0002
CDS_0001
MGV-GENOME-0117606 145:5820(-)
CDS_0005
CDS_0004
CDS_0002
CDS_0001
MGV-GENOME-0113991 57:3804(-)
CDS_0024
CDS_0025
uvig_402990 12986:16733(+)
CDS_0023
uvig_559424 12531:16243(+)
Lar-like restriction alleviation protein (CDS_0020)
CDS_0021
MGV-GENOME-0118380 10794:15947(+)
CDS_0022
CDS_0023
CDS_0024
MGV-GENOME-0113369 12727:16512(+)
CDS_0003
CDS_0001
CDS_0004
CDS_0002
MGV-GENOME-0117347 151:4412(-)
CDS_0002
CDS_0001
MGV-GENOME-0113371 26:3945(-)
CDS_0003
CDS_0002
CDS_0004
CDS_0001
MGV-GENOME-0070415 80:4881(-)
21
22
23
24
25
uvig_30797 11059:16651(+)
18
19
21
22
23
24
25
MGV-GENOME-4333327 10957:16644(+)
21
22
23
24
25
26
MGV-GENOME-0115708 11199:16801(+)
5
4
3
2
1
MGV-GENOME-0086644 124:5636(-)
CDS_0002
CDS_0004
CDS_0003
CDS_0001
uvig_88361 159:4523(-)
CDS_0007
CDS_0003
CDS_0002
CDS_0001
uvig_163374 40:6021(-)
CDS_0001
CDS_0002
MGV-GENOME-0110887 113:3419(-)
CDS_0001
MGV-GENOME-0101357 142:3111(-)
CDS_0020
CDS_0021
MGV-GENOME-0088340 11303:14714(+)
CDS_0020
MGV-GENOME-0112107 11084:15739(+)
CDS_0007
CDS_0005
MGV-GENOME-0119272 927:5814(-)
CDS_0004
CDS_0003
MGV-GENOME-0110092 293:5609(-)
CDS_0023
CDS_0024
CDS_0025
CDS_0026
uvig_303229 12648:16878(+)
CDS_0002
CDS_0003
CDS_0001
MGV-GENOME-0116514 150:4161(-)
CDS_0022
CDS_0023
MGV-GENOME-0104697 12699:15982(+)
CDS_0022
MGV-GENOME-0109561 10926:15746(+)
CDS_0010
CDS_0005
CDS_0004
MGV-GENOME-0117328 981:5974(-)
CDS_0003
CDS_0002
CDS_0001
MGV-GENOME-0110691 45:3789(-)
CDS_0022
MGV-GENOME-0115682 13090:16596(+)
CDS_0002
CDS_0001
uvig_70343 2:3866(-)
CDS_0023
CDS_0024
MGV-GENOME-0110492 12571:16435(+)
CDS_0023
CDS_0024
MGV-GENOME-0111554 12661:16525(+)
CDS_0005
CDS_0003
CDS_0004
MGV-GENOME-0110792 160:5533(-)
CDS_0007
CDS_0006
MGV-GENOME-0082247 540:5860(-)
22
23
24
25
26
Norman_66_NODE_39_length_17039_cov_1262.532972 11314:16955(+)
single strand DNA binding protein (CDS_0004)
CDS_0003
MGV-GENOME-0067051 232:4669(-)
2000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
transcription regulation
unknown function