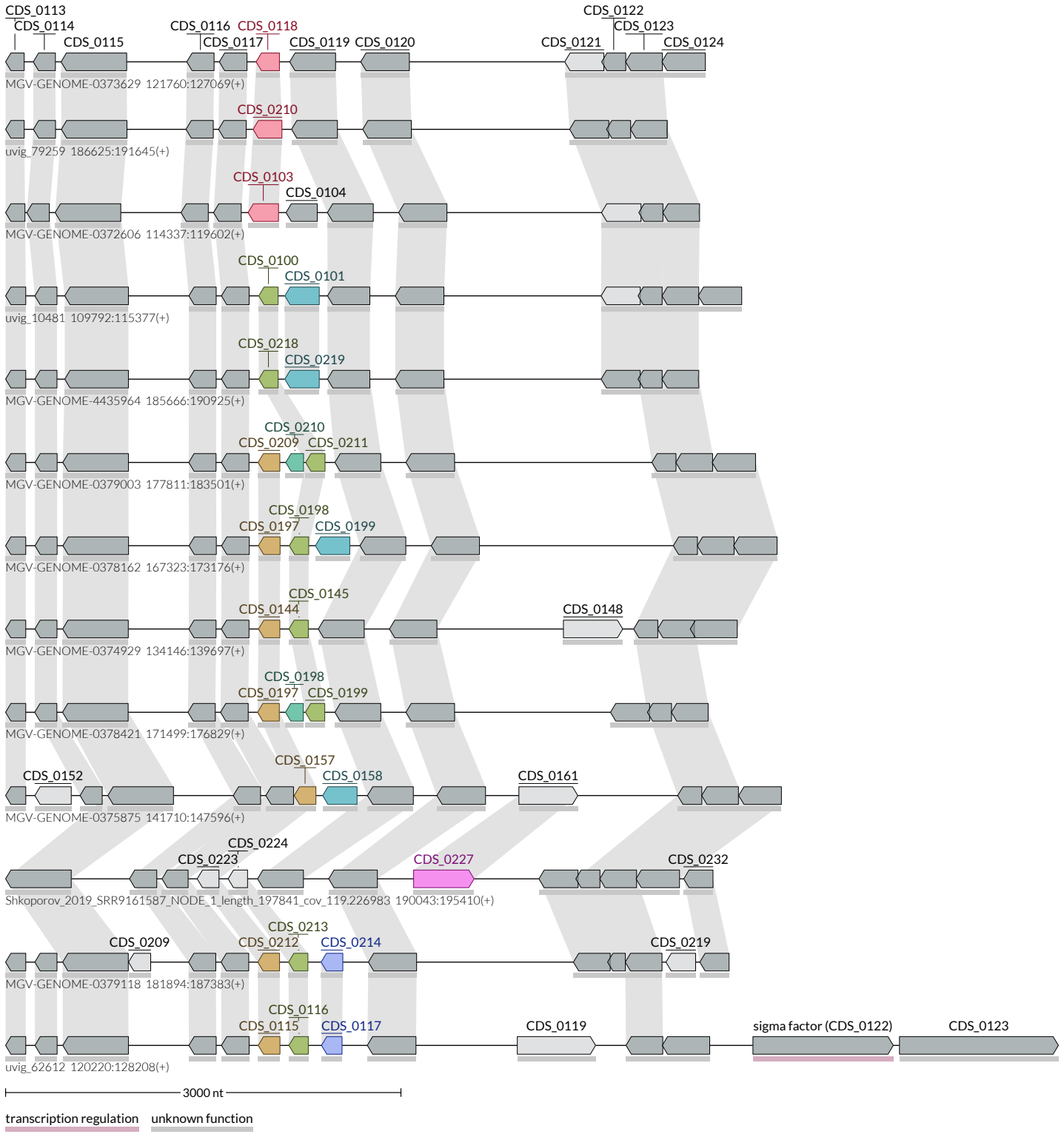
CDS_0113
CDS_0114
CDS_0115
CDS_0116
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CDS_0122
CDS_0123
CDS_0121
CDS_0124
MGV-GENOME-0373629 121760:127069(+)
CDS_0210
uvig_79259 186625:191645(+)
CDS_0103
CDS_0104
MGV-GENOME-0372606 114337:119602(+)
CDS_0100
CDS_0101
uvig_10481 109792:115377(+)
CDS_0218
CDS_0219
MGV-GENOME-4435964 185666:190925(+)
CDS_0210
CDS_0209
CDS_0211
MGV-GENOME-0379003 177811:183501(+)
CDS_0198
CDS_0197
CDS_0199
MGV-GENOME-0378162 167323:173176(+)
CDS_0145
CDS_0144
CDS_0148
MGV-GENOME-0374929 134146:139697(+)
CDS_0198
CDS_0197
CDS_0199
MGV-GENOME-0378421 171499:176829(+)
CDS_0157
CDS_0152
CDS_0158
CDS_0161
MGV-GENOME-0375875 141710:147596(+)
CDS_0224
CDS_0223
CDS_0227
CDS_0232
Shkoporov_2019_SRR9161587_NODE_1_length_197841_cov_119.226983 190043:195410(+)
CDS_0213
CDS_0209
CDS_0212
CDS_0214
CDS_0219
MGV-GENOME-0379118 181894:187383(+)
CDS_0116
CDS_0115
CDS_0117
CDS_0119
sigma factor (CDS_0122)
CDS_0123
uvig_62612 120220:128208(+)
3000 nt
transcription regulation
unknown function