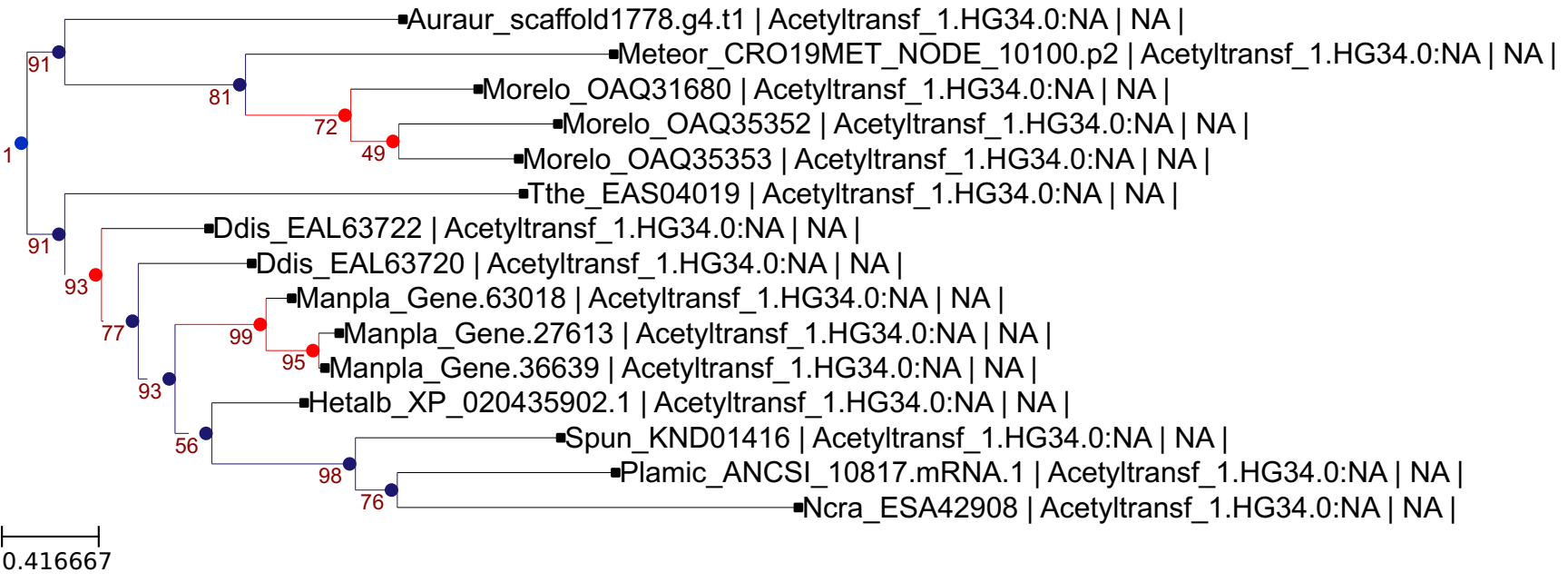
Auraur_scaffold1778.g4.t1 | Acetyltransf_1.HG34.0:NA | NA |
Meteor_CRO19MET_NODE_10100.p2 | Acetyltransf_1.HG34.0:NA | NA |
Morelo_OAQ31680 | Acetyltransf_1.HG34.0:NA | NA |
Morelo_OAQ35352 | Acetyltransf_1.HG34.0:NA | NA |
Morelo_OAQ35353 | Acetyltransf_1.HG34.0:NA | NA |
Tthe_EAS04019 | Acetyltransf_1.HG34.0:NA | NA |
Ddis_EAL63722 | Acetyltransf_1.HG34.0:NA | NA |
Ddis_EAL63720 | Acetyltransf_1.HG34.0:NA | NA |
Manpla_Gene.63018 | Acetyltransf_1.HG34.0:NA | NA |
Manpla_Gene.27613 | Acetyltransf_1.HG34.0:NA | NA |
Manpla_Gene.36639 | Acetyltransf_1.HG34.0:NA | NA |
Hetalb_XP_020435902.1 | Acetyltransf_1.HG34.0:NA | NA |
Spun_KND01416 | Acetyltransf_1.HG34.0:NA | NA |
Plamic_ANCSI_10817.mRNA.1 | Acetyltransf_1.HG34.0:NA | NA |
Ncra_ESA42908 | Acetyltransf_1.HG34.0:NA | NA |
91
81
72
49
1
91
93
77
99
95
93
56
98
76
0.416667