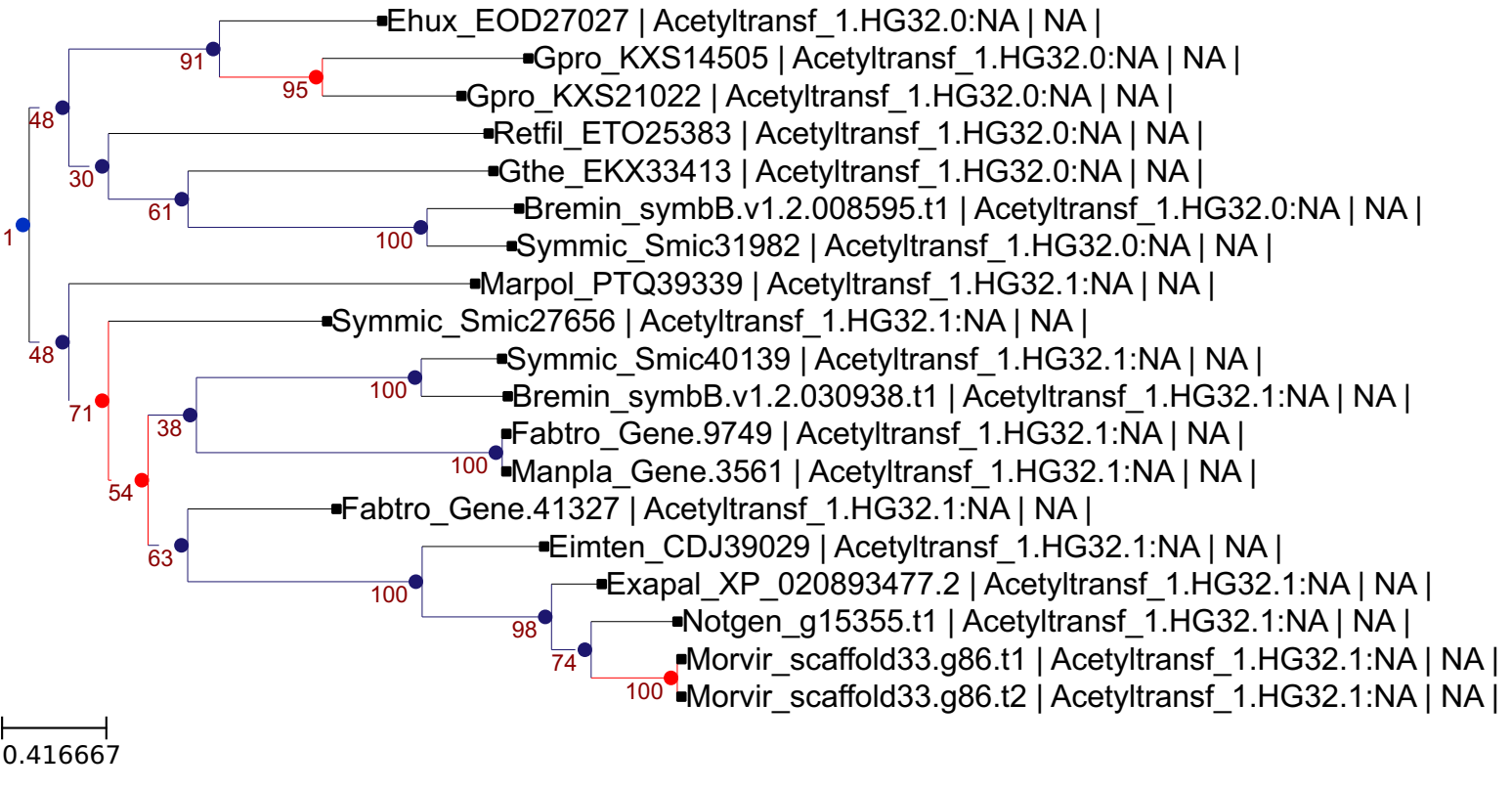
Ehux_EOD27027 | Acetyltransf_1.HG32.0:NA | NA |
Gpro_KXS14505 | Acetyltransf_1.HG32.0:NA | NA |
Gpro_KXS21022 | Acetyltransf_1.HG32.0:NA | NA |
Retfil_ETO25383 | Acetyltransf_1.HG32.0:NA | NA |
Gthe_EKX33413 | Acetyltransf_1.HG32.0:NA | NA |
Bremin_symbB.v1.2.008595.t1 | Acetyltransf_1.HG32.0:NA | NA |
Symmic_Smic31982 | Acetyltransf_1.HG32.0:NA | NA |
Marpol_PTQ39339 | Acetyltransf_1.HG32.1:NA | NA |
Symmic_Smic27656 | Acetyltransf_1.HG32.1:NA | NA |
Symmic_Smic40139 | Acetyltransf_1.HG32.1:NA | NA |
Bremin_symbB.v1.2.030938.t1 | Acetyltransf_1.HG32.1:NA | NA |
Fabtro_Gene.9749 | Acetyltransf_1.HG32.1:NA | NA |
Manpla_Gene.3561 | Acetyltransf_1.HG32.1:NA | NA |
Fabtro_Gene.41327 | Acetyltransf_1.HG32.1:NA | NA |
Eimten_CDJ39029 | Acetyltransf_1.HG32.1:NA | NA |
Exapal_XP_020893477.2 | Acetyltransf_1.HG32.1:NA | NA |
Notgen_g15355.t1 | Acetyltransf_1.HG32.1:NA | NA |
Morvir_scaffold33.g86.t1 | Acetyltransf_1.HG32.1:NA | NA |
Morvir_scaffold33.g86.t2 | Acetyltransf_1.HG32.1:NA | NA |
91
95
48
30
61
100
1
48
71
100
38
100
54
63
100
98
74
100
0.416667