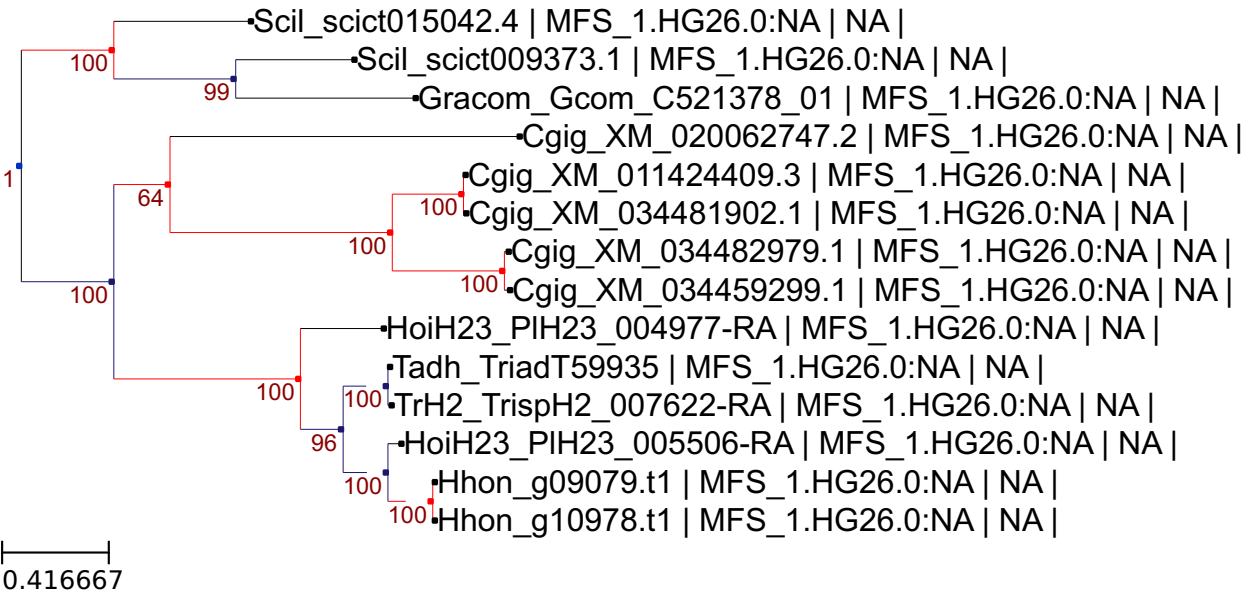
Scil_scict015042.4 | MFS_1.HG26.0:NA | NA |
Scil_scict009373.1 | MFS_1.HG26.0:NA | NA |
Gracom_Gcom_C521378_01 | MFS_1.HG26.0:NA | NA |
Cgig_XM_020062747.2 | MFS_1.HG26.0:NA | NA |
Cgig_XM_011424409.3 | MFS_1.HG26.0:NA | NA |
Cgig_XM_034481902.1 | MFS_1.HG26.0:NA | NA |
Cgig_XM_034482979.1 | MFS_1.HG26.0:NA | NA |
Cgig_XM_034459299.1 | MFS_1.HG26.0:NA | NA |
HoiH23_PIH23_004977-RA | MFS_1.HG26.0:NA | NA |
Tadh_TriadT59935 | MFS_1.HG26.0:NA | NA |
TrH2_TrispH2_007622-RA | MFS_1.HG26.0:NA | NA |
HoiH23_PIH23_005506-RA | MFS_1.HG26.0:NA | NA |
Hhon_g09079.t1 | MFS_1.HG26.0:NA | NA |
Hhon_g10978.t1 | MFS_1.HG26.0:NA | NA |
100
99
1
64
100
100
100
100
100
100
96
100
100
0.416667