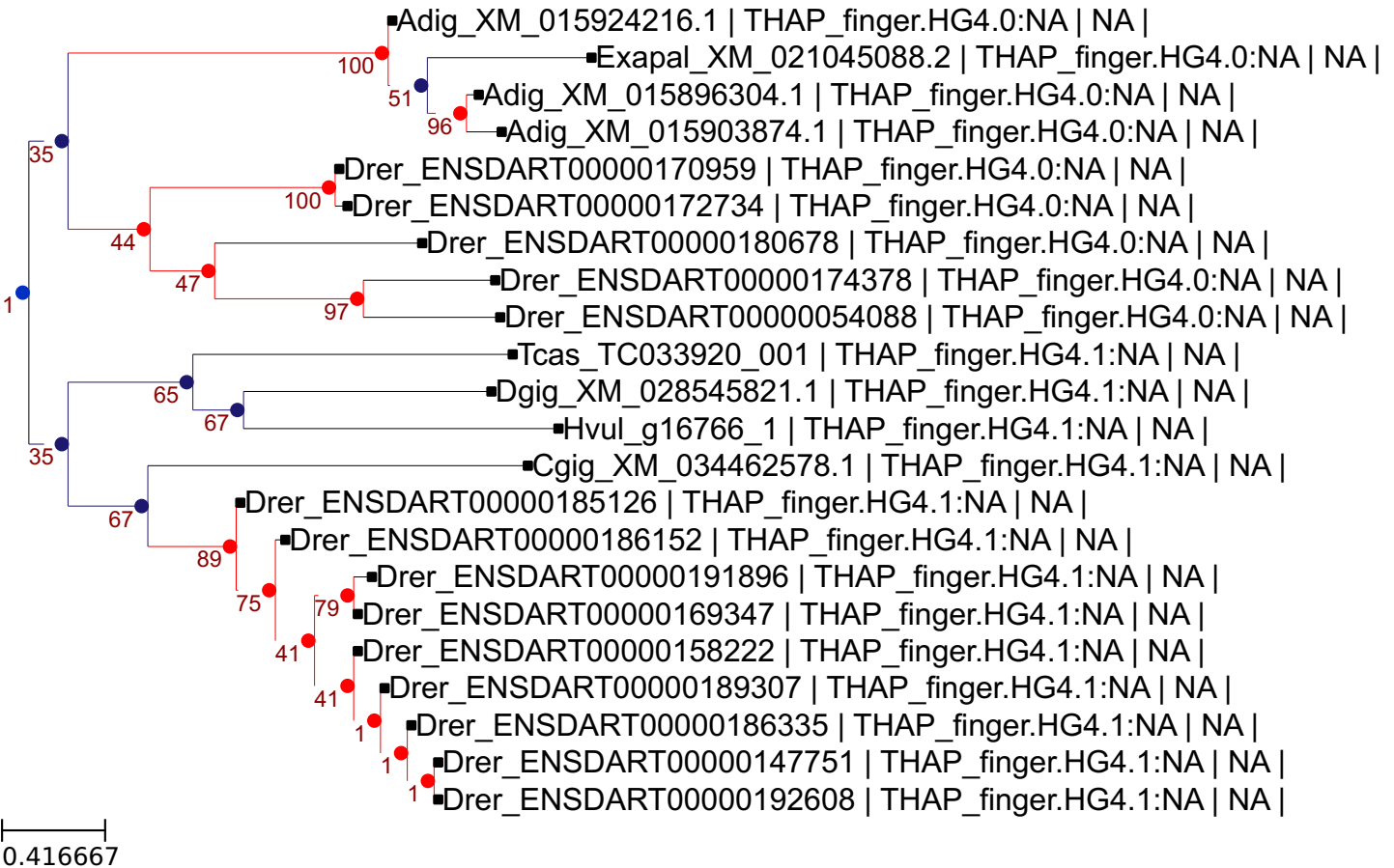
Adig_XM_015924216.1 | THAP_finger.HG4.0:NA | NA |
Exapal_XM_021045088.2 | THAP_finger.HG4.0:NA | NA |
Adig_XM_015896304.1 | THAP_finger.HG4.0:NA | NA |
Adig_XM_015903874.1 | THAP_finger.HG4.0:NA | NA |
Drer_ENSDART00000170959 | THAP_finger.HG4.0:NA | NA |
Drer_ENSDART00000172734 | THAP_finger.HG4.0:NA | NA |
Drer_ENSDART00000180678 | THAP_finger.HG4.0:NA | NA |
Drer_ENSDART00000174378 | THAP_finger.HG4.0:NA | NA |
Drer_ENSDART00000054088 | THAP_finger.HG4.0:NA | NA |
Tcas_TC033920_001 | THAP_finger.HG4.1:NA | NA |
Dgig_XM_028545821.1 | THAP_finger.HG4.1:NA | NA |
Hvul_g16766_1 | THAP_finger.HG4.1:NA | NA |
Cgig_XM_034462578.1 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000185126 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000186152 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000191896 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000169347 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000158222 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000189307 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000186335 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000147751 | THAP_finger.HG4.1:NA | NA |
Drer_ENSDART00000192608 | THAP_finger.HG4.1:NA | NA |
100
51
96
35
100
44
47
97
1
65
67
35
67
89
75
79
41
41
1
1
1
0.416667