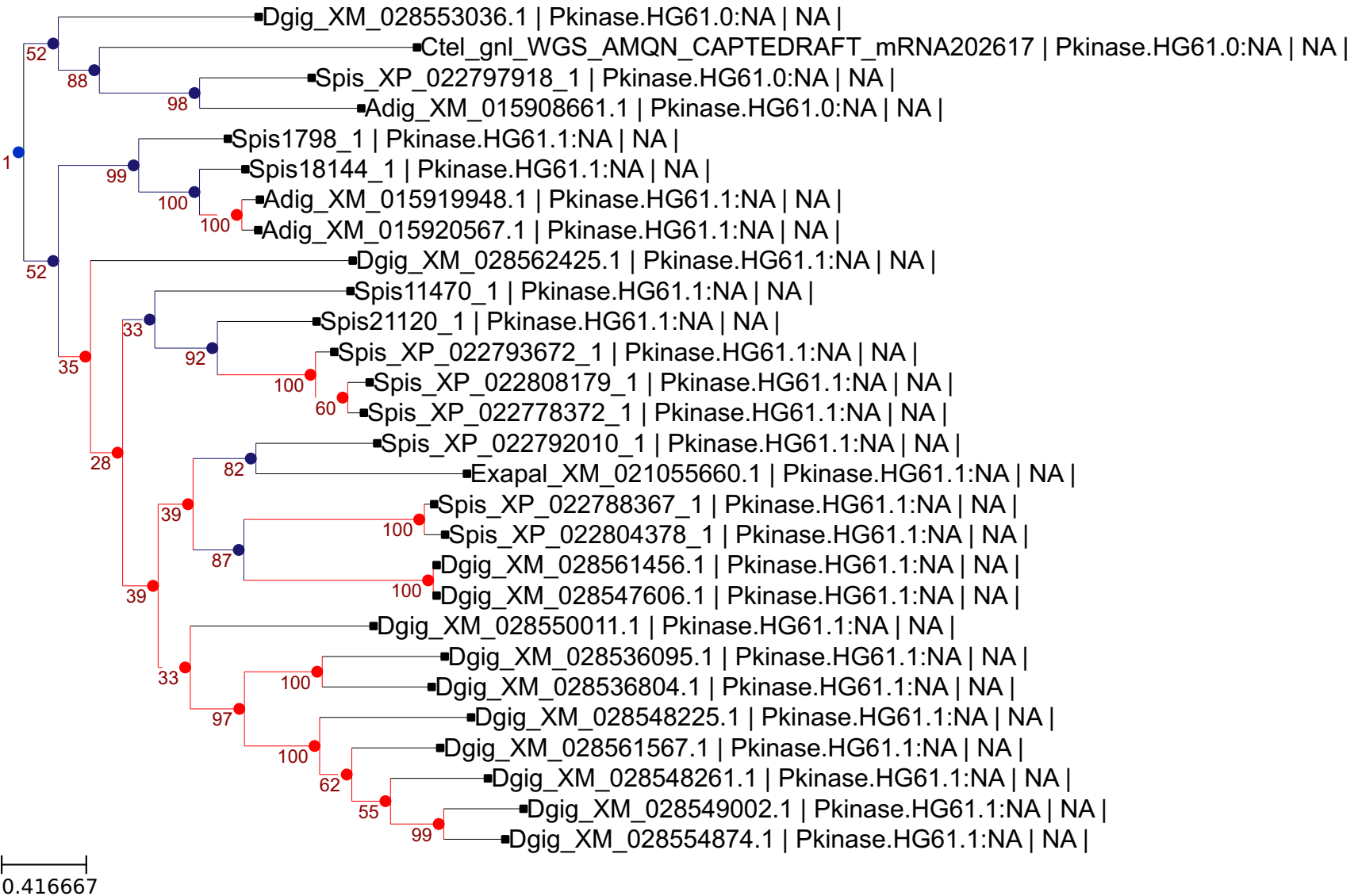
Dgig_XM_028553036.1 | Pkinase.HG61.0:NA | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA202617 | Pkinase.HG61.0:NA | NA |
Spis_XP_022797918_1 | Pkinase.HG61.0:NA | NA |
Adig_XM_015908661.1 | Pkinase.HG61.0:NA | NA |
Spis1798_1 | Pkinase.HG61.1:NA | NA |
Spis18144_1 | Pkinase.HG61.1:NA | NA |
Adig_XM_015919948.1 | Pkinase.HG61.1:NA | NA |
Adig_XM_015920567.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028562425.1 | Pkinase.HG61.1:NA | NA |
Spis11470_1 | Pkinase.HG61.1:NA | NA |
Spis21120_1 | Pkinase.HG61.1:NA | NA |
Spis_XP_022793672_1 | Pkinase.HG61.1:NA | NA |
Spis_XP_022808179_1 | Pkinase.HG61.1:NA | NA |
Spis_XP_022778372_1 | Pkinase.HG61.1:NA | NA |
Spis_XP_022792010_1 | Pkinase.HG61.1:NA | NA |
Exapal_XM_021055660.1 | Pkinase.HG61.1:NA | NA |
Spis_XP_022788367_1 | Pkinase.HG61.1:NA | NA |
Spis_XP_022804378_1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028561456.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028547606.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028550011.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028536095.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028536804.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028548225.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028561567.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028548261.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028549002.1 | Pkinase.HG61.1:NA | NA |
Dgig_XM_028554874.1 | Pkinase.HG61.1:NA | NA |
1
52
88
98
99
100
100
52
33
92
100
60
35
28
82
39
100
87
100
39
33
100
97
100
62
55
99
0.416667