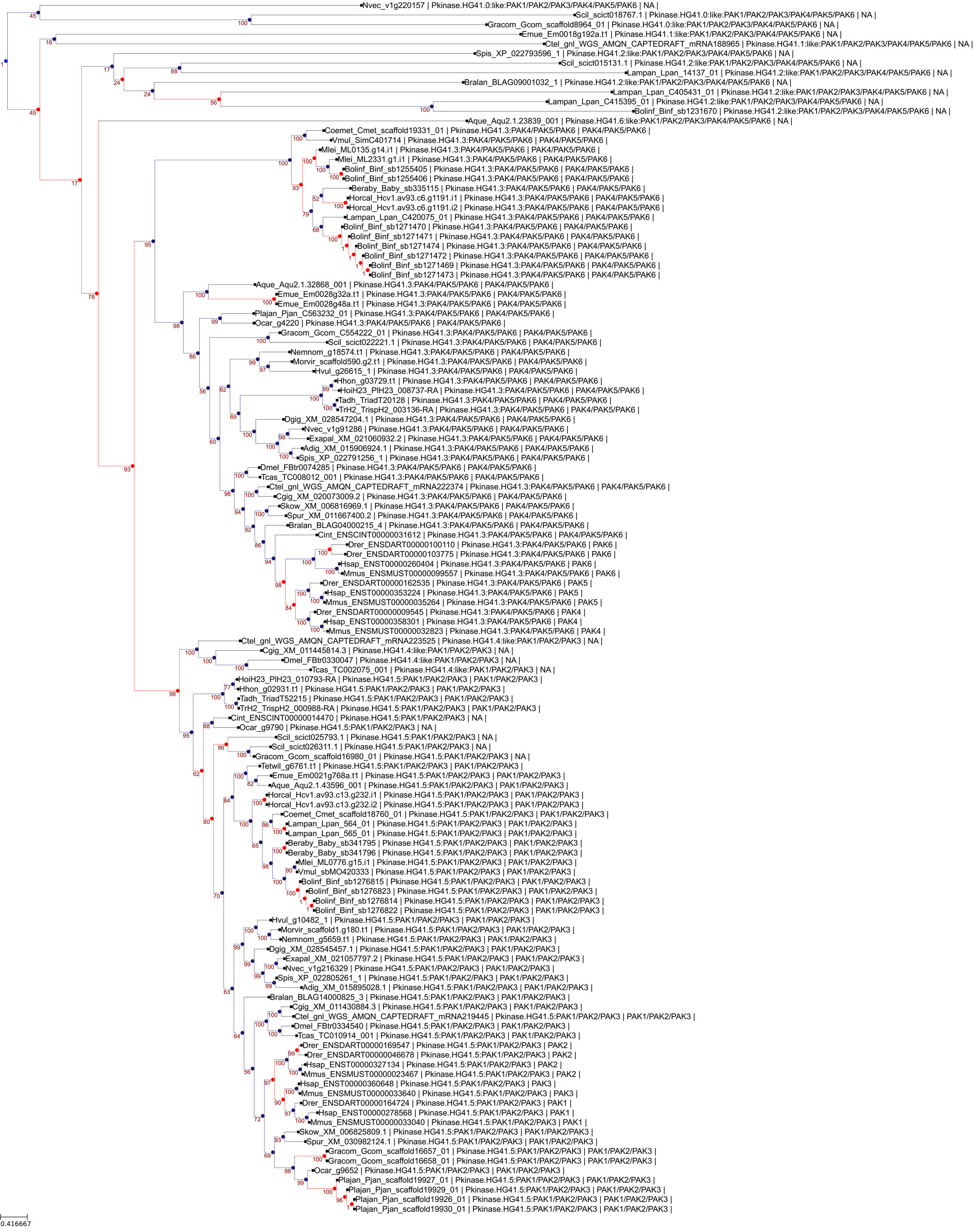
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Gracom_Gcom_scaffold8964_01 | Pkinase.HG41.0:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Emue_Em0018g192a.t1 | Pkinase.HG41.1:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA188965 | Pkinase.HG41.1:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Spis_XP_022793596_1 | Pkinase.HG41.2:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
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Lampan_Lpan_14137_01 | Pkinase.HG41.2:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Bralan_BLAG09001032_1 | Pkinase.HG41.2:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Lampan_Lpan_C405431_01 | Pkinase.HG41.2:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Lampan_Lpan_C415395_01 | Pkinase.HG41.2:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Bolinf_Binf_sb1231670 | Pkinase.HG41.2:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Aque_Aqu2.1.23839_001 | Pkinase.HG41.6:like:PAK1/PAK2/PAK3/PAK4/PAK5/PAK6 | NA |
Coemet_Cmet_scaffold19331_01 | Pkinase.HG41.3:PAK4/PAK5/PAK6 | PAK4/PAK5/PAK6 |
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Mlei_ML2331.g1.i1 | Pkinase.HG41.3:PAK4/PAK5/PAK6 | PAK4/PAK5/PAK6 |
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