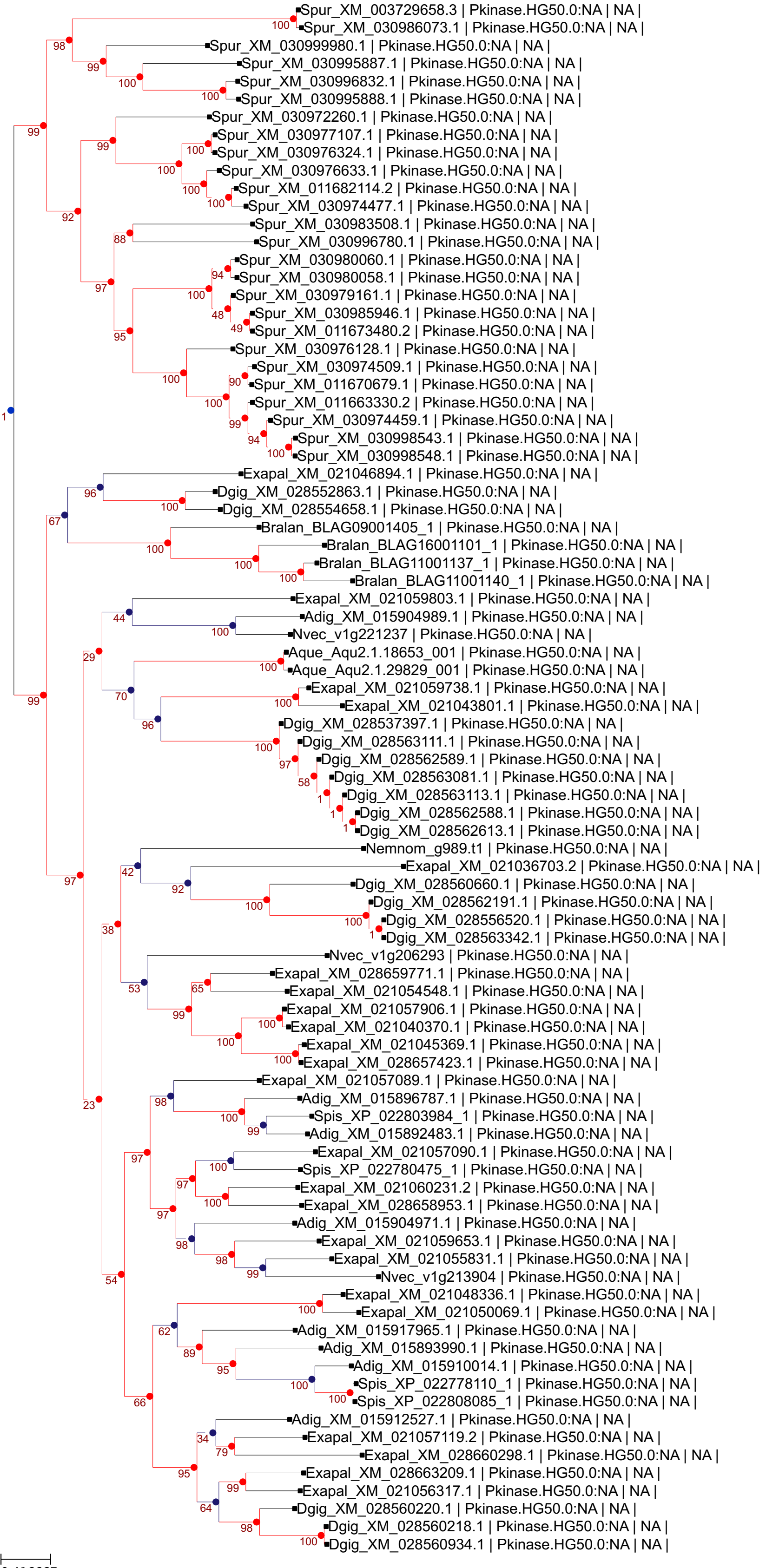
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Spur_XM_030995887.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030996832.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030995888.1 | Pkinase.HG50.0:NA | NA |
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Spur_XM_030977107.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030976324.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030976633.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_011682114.2 | Pkinase.HG50.0:NA | NA |
Spur_XM_030974477.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030983508.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030996780.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030980060.1 | Pkinase.HG50.0:NA | NA |
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Spur_XM_030979161.1 | Pkinase.HG50.0:NA | NA |
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Spur_XM_011673480.2 | Pkinase.HG50.0:NA | NA |
Spur_XM_030976128.1 | Pkinase.HG50.0:NA | NA |
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Spur_XM_030974459.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030998543.1 | Pkinase.HG50.0:NA | NA |
Spur_XM_030998548.1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021046894.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028552863.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028554658.1 | Pkinase.HG50.0:NA | NA |
Bralan_BLAG09001405_1 | Pkinase.HG50.0:NA | NA |
Bralan_BLAG16001101_1 | Pkinase.HG50.0:NA | NA |
Bralan_BLAG11001137_1 | Pkinase.HG50.0:NA | NA |
Bralan_BLAG11001140_1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021059803.1 | Pkinase.HG50.0:NA | NA |
Adig_XM_015904989.1 | Pkinase.HG50.0:NA | NA |
Nvec_v1g221237 | Pkinase.HG50.0:NA | NA |
Aque_Aqu2.1.18653_001 | Pkinase.HG50.0:NA | NA |
Aque_Aqu2.1.29829_001 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021059738.1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021043801.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028537397.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028563111.1 | Pkinase.HG50.0:NA | NA |
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Nemnom_g989.t1 | Pkinase.HG50.0:NA | NA |
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Nvec_v1g206293 | Pkinase.HG50.0:NA | NA |
Exapal_XM_028659771.1 | Pkinase.HG50.0:NA | NA |
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Exapal_XM_021057906.1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021040370.1 | Pkinase.HG50.0:NA | NA |
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Nvec_v1g213904 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021048336.1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021050069.1 | Pkinase.HG50.0:NA | NA |
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Adig_XM_015893990.1 | Pkinase.HG50.0:NA | NA |
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Spis_XP_022808085_1 | Pkinase.HG50.0:NA | NA |
Adig_XM_015912527.1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021057119.2 | Pkinase.HG50.0:NA | NA |
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Exapal_XM_028663209.1 | Pkinase.HG50.0:NA | NA |
Exapal_XM_021056317.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028560220.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028560218.1 | Pkinase.HG50.0:NA | NA |
Dgig_XM_028560934.1 | Pkinase.HG50.0:NA | NA |
0.416667