Ocar_g9308 | Pkinase.HG49.3:NA | NA |
Tetwil_g20556.t1 | Pkinase.HG49.0:NA | NA |
Ocar_g3531 | Pkinase.HG49.0:NA | NA |
Plajan_Pjan_scaffold19586_01 | Pkinase.HG49.0:NA | NA |
Plajan_Pjan_scaffold19587_01 | Pkinase.HG49.0:NA | NA |
Cgig_XM_034467594.1 | Pkinase.HG49.0:NA | NA |
Drer_ENSDART00000050934 | Pkinase.HG49.0:NA | NA |
Spis_XP_022794966_1 | Pkinase.HG49.0:NA | NA |
Bralan_BLAG08000295_1 | Pkinase.HG49.0:NA | NA |
Spur_XM_011684200.2 | Pkinase.HG49.0:NA | NA |
Tetwil_g13390.t1 | Pkinase.HG49.1:NA | NA |
Tetwil_g21541.t1 | Pkinase.HG49.1:NA | NA |
Tetwil_g13389.t3 | Pkinase.HG49.1:NA | NA |
Exapal_XM_028662242.1 | Pkinase.HG49.1:NA | NA |
Exapal_XM_021045182.2 | Pkinase.HG49.1:NA | NA |
Exapal_XM_028660416.1 | Pkinase.HG49.1:NA | NA |
Bralan_BLAG18000684_4 | Pkinase.HG49.2:NA | NA |
Spur_XM_030993653.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030993633.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030986522.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030987641.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030987522.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030987264.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030987263.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030986808.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030986526.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_011668633.2 | Pkinase.HG49.2:NA | NA |
Spur_XM_030987633.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030986524.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030986523.1 | Pkinase.HG49.2:NA | NA |
Spur_XM_030987640.1 | Pkinase.HG49.2:NA | NA |
Bralan_BLAG18000708_1 | Pkinase.HG49.2:NA | NA |
Skow_XM_006813871.1 | Pkinase.HG49.2:NA | NA |
Skow_XM_002734065.1 | Pkinase.HG49.2:NA | NA |
Skow_XM_002735303.1 | Pkinase.HG49.2:NA | NA |
Cgig_XM_034449738.1 | Pkinase.HG49.2:NA | NA |
Skow_XM_002730968.1 | Pkinase.HG49.2:NA | NA |
Skow_XM_002738296.1 | Pkinase.HG49.2:NA | NA |
Dgig_XM_028556253.1 | Pkinase.HG49.2:NA | NA |
Nvec_v1g139065 | Pkinase.HG49.2:NA | NA |
Nvec_v1g131691 | Pkinase.HG49.2:NA | NA |
Nvec_NVE17967 | Pkinase.HG49.2:NA | NA |
Exapal_XM_021062017.2 | Pkinase.HG49.2:NA | NA |
Exapal_XM_021050880.2 | Pkinase.HG49.2:NA | NA |
Adig_XM_015908184.1 | Pkinase.HG49.2:NA | NA |
Spis_XP_022807184_1 | Pkinase.HG49.2:NA | NA |
Emue_Em0017g941a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0002g420a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0001g188a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0006g1003a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0021g721a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0632g3a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0021g187a.t1 | Pkinase.HG49.2:NA | NA |
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Emue_Em0032g20a.t1 | Pkinase.HG49.2:NA | NA |
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Emue_Em0010g953a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0022g942a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0022g689a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0022g771a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0685g1a.t1 | Pkinase.HG49.2:NA | NA |
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Emue_Em0001g709a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0040g30a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0017g323a.t1 | Pkinase.HG49.2:NA | NA |
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Emue_Em0001g1968a.t1 | Pkinase.HG49.2:NA | NA |
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Emue_Em0001g2551a.t1 | Pkinase.HG49.2:NA | NA |
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Emue_Em0011g997a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0019g674a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0001g3303a.t1 | Pkinase.HG49.2:NA | NA |
Emue_Em0006g1364a.t1 | Pkinase.HG49.2:NA | NA |
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