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	Scil_scict010238.1 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Scil_scict010238.1 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Mlei_ML0077.g2.i1 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Cgig_XM_011434135.3 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA131028 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK1/SGK3/SGK3/SGK1/SGK3/SGK3/SGK1/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3	■Nvec_v1g27751 Pkinase.HG12.3:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA GK1/SGK2/SGK3 NA GK2/SGK3 NA
	■Dmel_FBtr0072464 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 ■Tcas_TC002973_001 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 ■Hhon_g10523.t1 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA ■HoiH23_PIH23_008482-RA Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA ■TrH2_TrispH2_008866-RA Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA 99 ■Tadh_TriadT31577 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA	
70	Drer_ENSDART00000003789 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Skow_XM_002735355.2 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Adig_XM_015922557.1 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Dgig_XM_028539625.1 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Ocar_g205 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Plajan_Pjan_scaffold21438_01 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK1/SGK1/SGK1/SGK1/SGK1/SGK1/SGK1	2/SGK3 NA
	Plajan_Pjan_scaffold21439_01 Pkinase.HG12.4:like:C8orf44-SGK3/SGK1/SGK1/SGK2/SGK3 NA -Aque_Aqu2.1.30465_001 Pkinase.HG12.5:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Emue_Em0015g1018a.t1 Pkinase.HG12.5:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Tetwil_g23492.t2 Pkinase.HG12.5:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Plajan_Pjan_C571910_01 Pkinase.HG12.5:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA -Scil_scict004171.1 Pkinase.HG12.5:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA	
91	Gracom_Gcom_C521738_01 Pkinase.HG12.5:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA	(3 NA
73	**Ocar_g574 Pkinase.HG12.8:like:C8orf44-SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/	/SGK1/SGK2/SGK3 NA 1/SGK2/SGK3 NA 1/SGK2/SGK3 NA NA GK3 NA
	Plajan_Pjan_scaffold20635_01 Pkinase.HG12.8:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Plajan_Pjan_scaffold20636_01 Pkinase.HG12.8:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Cgig_XM_011448993.3 Pkinase.HG12.8:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Bralan_BLAG18000059_1 Pkinase.HG12.8:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Dgig_XM_028540584.1 Pkinase.HG12.8:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA Spis_XP_022804682_1 Pkinase.HG12.8:like:C8orf44-SGK3/SGK1/SGK2/SGK3 NA	
	**Scil_scict007450.1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 NA **Scil_scict007450.1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 NA **Gracom_Gcom_scaffold11908_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 NA **Gracom_Gcom_scaffold11908_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3/SGK3/SGK1/SGK2/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3/SGK3	
100	Gracom_Gcom_scaffold2132_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Scil_scict023087.2 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Gracom_Gcom_scaffold17222_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Car_g8784 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Gracom_g8784 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3	
98	Plajan_Pjan_scaffold16256_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Aque_Aqu2.1.32726_001 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Emue_Em0005g22a.t1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Emue_Em0005g45a.t1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Coemet_Cmet_scaffold9786_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Coemet_Cmet_scaffold9787_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3	
67 [•]	Lampan_Lpan_13828_01 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Mlei_ML0622.g1.i1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Horcal_Hcv1.av93.c5.g923.i3 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK4 Horcal_Hcv1.av93.c5.g923.i4 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK4 Horcal_Hcv1.av93.c5.g923.i5 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK4 Horcal_Hcv1.av93.c5.g923.i1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK4/SGK4/SGK4/SGK4/SGK4/SGK4/SGK4/SGK4	3 3 3K3
	99 Horcal_Hcv1.av93.c5.g923.i2 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Drer_ENSDART00000090368 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3 Mmus_ENSMUST00000097826 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3 Hsap_ENST00000345714 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3 Drer_ENST00000519289 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3 Drer_ENSDART00000135027 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 SGK2 Jehsap_ENST00000341458 Jehsap_ENST000000341458 Jehsap_ENST000000341458 Jehsap_ENST000000000000000000000000000000000000	ΣK3
	**Mmus_ENSMUST0000018012 Pkinase.HG12.0:C80rf44-SGK3/SGK1/SGK2/SGK3 SGK2 **Drer_ENSDART00000010580 Pkinase.HG12.0:C80rf44-SGK3/SGK1/SGK2/SGK3 NA **Drer_ENSDART00000008152 Pkinase.HG12.0:C80rf44-SGK3/SGK1/SGK2/SGK3 NA **Drer_ENSDART00000035031 Pkinase.HG12.0:C80rf44-SGK3/SGK1/SGK2/SGK3 SGK1 **Hsap_ENST00000367858 Pkinase.HG12.0:C80rf44-SGK3/SGK1/SGK2/SGK3 SGK1 **Mmus_ENSMUST00000120509 Pkinase.HG12.0:C80rf44-SGK3/SGK1/SGK2/SGK3 SGK1	
	Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA172316 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Spur_XM_030983204.1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Spur_XM_030983207.1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Bralan_BLAG02001634_1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Skow_XM_002730567.2 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK3/SGK1/SGK2/SGK3/SGK3/SGK1/SGK2/SGK3/SGK3/SGK3/SGK1/SGK2/SGK3/SGK3/S	GK1/SGK2/SGK3
	TrH2_TrispH2_004479-RA Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Tadh_TriadT27989 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 HoiH23_PIH23_005586-RA Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Hhon_g07449.t1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Hhon_g09747.t1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Dgig_XM_028546712.1 Pkinase.HG12.0:C8orf44-SGK3/SGK1/SGK2/SGK3 C8orf44-SGK3/SGK1/SGK2/SGK3 Table 100	
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100	Mlei_ML0922.g2.i1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 "Mlei_ML0922.g2.i1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 "Bolinf_Binf_sb1257183 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 "Bolinf_Binf_sb1257186 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 "Bolinf_Binf_sb1257184 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 "Hvul_g33421_1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3	
96 96	Nemnom_g3118.t1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Morvir_scaffold188.g18.t1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Exapal_XM_021050331.2 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3	
94	*Adig_XM_015907309.1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 *Adig_XM_015907337.1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 *Ocar_g8834 Pkinase.HG12.1:AKT1/AKT2/AKT3 NA *Plajan_Pjan_scaffold22974_01 Pkinase.HG12.1:AKT1/AKT2/AKT3 NA *HoiH23_PIH23_001833-RA Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 *Hhon_g06571.t1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3	
85	TrH2_TrispH2_004957-RA Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Tadh_TriadT51134 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Plajan_Pjan_scaffold6021_01 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Plajan_Pjan_scaffold6022_01 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Scil_scict008994.1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Gracom_Gcom_scaffold13787_01 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3 Scil_scict004815.1 Pkinase.HG12.1:AKT1/AKT2/AKT3 AKT1/AKT2/AKT3	
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73	**Gracom_Gcom_scaffold12145_01 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Scil_scict022118.2 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Aque_Aqu2.1.34192_001 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Tetwil_g5214.t2 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Ocar_g3208 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Ocar_g3208 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Tetwil_g5214.t2 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Tetwil_g5214.t2 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Tetwil_g5214.t2 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 **Tetwil_g5214.t2 Pkinase.HG12.3:PKN1/PKN2/PKN3 **Tetwil_g5214.t2 Pkinas	
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	Beraby_Baby_sb334012 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Bolinf_Binf_sb1224445 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Bolinf_Binf_sb1224446 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Horcal_Hcv1.av93.c7.g954.i6 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Horcal_Hcv1.av93.c7.g954.i11 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Horcal_Hcv1.av93.c7.g954.i3 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3	
	Horcal_Hcv1.av93.c7.g954.i5 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Horcal_Hcv1.av93.c7.g954.i2 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Horcal_Hcv1.av93.c7.g954.i1 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Horcal_Hcv1.av93.c7.g954.i4 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 Cint_ENSCINT00000018412 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1 Prer_ENSDART00000091017 Pkinase.HG12.2:PKN1/PKN2/PKN3 NA	
	Drer_ENSDART00000132555 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1 "Mmus_ENSMUST00000144258 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1 "Hsap_ENST00000342216 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1 "Drer_ENSDART00000112546 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN3 "Hsap_ENST00000291906 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN3 "Mmus_ENSMUST00000045246 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN3	
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	*HoiH23_PIH23_006565-RA Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 *Tadh_TriadT20881 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 *TrH2_TrispH2_009334-RA Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 *Cgig_XM_034447138.1 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 *Bralan_BLAG08000530_4 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3 *Skow XM 006816415.1 Pkinase.HG12.2:PKN1/PKN2/PKN3 PKN1/PKN2/PKN3	
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73	100	
	Tetwil_g25223.t2 Pkinase.HG12.6:PRKCI/PRKCZ PRKCI/PRKCZ PRKCI/PRKCZ —Emue_Em0008g100a.t1 Pkinase.HG12.6:PRKCI/PRKCZ PRKCI/PRKCZ —Aque_Aqu2.1.41475_001 Pkinase.HG12.6:PRKCI/PRKCZ PRKCI/PRKCZ —Aque_Aqu2.1.02349_001 Pkinase.HG12.6:PRKCI/PRKCZ PRKCI/PRKCZ —Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA213310 Pkinase.HG12.6:PRKCI/PRKCZ —Cgig_XM_011437951.3 Pkinase.HG12.6:PRKCI/PRKCZ PRKCI/PRKCZ	
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