Tea1 MGG_02875T0 [1459–1438] SDSLDTTSEYNSGGGERNSAALDNLANELETLR S1419
Tea1 MGG_02875T0 [1459–1438] SDSLDTTSEYNSGGGERNSAALDNLANELETLR S/YT
Tea1 MGG_02875T0 [1459–1438] SDSLDTTSEYNSGGGERNSAALDNLANELETLR S1425
Tea1 MGG_02875T0 [1459–1438] SDSLDTTSEYNSGGGERNSAALDNLANELETLR T1437
Septin4 MGG_06726T0 [300–323] QLLALKDFSAQGHSSRPSPAAER S
Septin5 MGG_03087T0 [364–383] EAASSVGFRIDGSEANGEDGN S12(98.6); S
Septin5 MGG_03087T0 [364–383] EAASSVGFRIDGSEANGEDGN S367; S375
Septin5 MGG_03087T0 [364–383] EAASSVGFRIDGSEANGEDGN S368; S375
Tea1 MGG_02875T0 [628–641] SQGSVDSSTDPGLK S631
Tea1 MGG_02875T0 [628–641] SQGSVDSSTDPGLK S/T
Tea1 MGG_02875T0 [463–480] RPSTSNSNEAGGMQVANR S465
Tea1 MGG_02875T0 [463–480] RPSTSNSNEAGGMQVANR S469
Tea1 MGG_02875T0 [463–480] RPSTSNSNEAGGMQVANR M13;1xPhospho T/S
Tea1 MGG_02875T0 [463–480] RPSTSNSNEAGGMQVANR T466
Tea1 MGG_02875T0 [1131–1144] ASRRGETPDTRAAR T1137
Tea1 MGG_02875T0 [14–35] SLSSRDGPPQSSNSSQSTAAR S17
Tea1 MGG_02875T0 [14–35] SLSSRDGPPQSSNSSQSTAAR S16
Tea1 MGG_02875T0 [14–35] SLSSRDGPPQSSNSSQSTAAR S/T
Mse4 MGG_06572T0 [301–322] DGPDAAYTSGRFSPTNTPTGGR T/S/Y
Mse4 MGG_06572T0 [301–322] DGPDAAYTSGRFSPTNTPTGGR T317
Septin4 MGG_06726T0 [306–323] DPSAQGHSSRPSPAAER S318
Rvs167 MGG_11497T0 [323–347] YGAVHSPAQRAAAENPPPPYSPGVK Y342
Rvs167 MGG_11497T0 [323–347] YGAVHSPAQRAAAENPPPPYSPGVK Y323
Rvs167 MGG_11497T0 [323–347] YGAVHSPAQRAAAENPPPPYSPGVK S328
Rvs167 MGG_11497T0 [323–347] YGAVHSPAQRAAAENPPPPYSPGVK Y/S
Rvs167 MGG_11497T0 [323–347] YGAVHSPAQRAAAENPPPPYSPGVK S343
Tea1 MGG_02875T0 [664–682] SSNRNSQTVVLLKELDMAR S664; T671
Tea1 MGG_02875T0 [664–682] SSNRNSQTVVLLKELDMAR M17;2xPhospho T8(91); S
Tea1 MGG_02875T0 [664–682] SSNRNSQTVVLLKELDMAR S669; T671
Tea1 MGG_02875T0 [664–678] SSNRNSQTVVLLK S665; T671
Tea1 MGG_02875T0 [664–678] SSNRNSQTVVLLK T671; S
Tea1 MGG_02875T0 [1129–1144] SRASRRGETPDTRAAR T1137
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN S368
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN S367
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN M1;1xPhospho S
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN S375
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN S375; S
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN S367; S375
Septin5 MGG_03087T0 [361–383] MQREAASSVGFRIDGSEANGEDGN S368; S375
Tea1 MGG_02875T0 [697–712] KADYTPNASMSPVFER S707
Las17 MGG_02862T0 [263–283] KESPRRQQAAHPPPPPPPSNDR S262
Mse4 MGG_06572T0 [323–330] RASLSLVR S327
Mse4 MGG_06572T0 [323–330] RASLSLVR S325
Mse4 MGG_06572T0 [323–330] RASLSLVR S
Septin5 MGG_03087T0 [364–372] EAASSVGFR S
Chm1 MGG_06320T0 [555–566] MSTMSETEVMAK S556
Chm1 MGG_06320T0 [555–566] MSTMSETEVMAK M1; M4; M10;1xPhospho S/T
Septin4 MGG_06726T0 [300–323] QLLALKDFSAQGHSSRPSPAAER S318
Septin4 MGG_06726T0 [300–323] QLLALKDFSAQGHSSRPSPAAER S308
Septin4 MGG_06726T0 [300–327] QLLALKDFSAQGHSSRPSPAAEREMSR M26;1xPhospho S
Septin4 MGG_06726T0 [300–327] QLLALKDFSAQGHSSRPSPAAEREMSR S308
Septin4 MGG_06726T0 [300–327] QLLALKDFSAQGHSSRPSPAAEREMSR S318
Septin4 MGG_06726T0 [300–327] QLLALKDFSAQGHSSRPSPAAEREMSR S326
Septin4 MGG_06726T0 [300–327] QLLALKDFSAQGHSSRPSPAAEREMSR S313
Tea1 MGG_02875T0 [468–488] QQGTRRPSTSNSNEAGGMQVANR S469
Tea1 MGG_02875T0 [468–488] QQGTRRPSTSNSNEAGGMQVANR M18;1xPhospho S/T
Tea1 MGG_02875T0 [468–488] QQGTRRPSTSNSNEAGGMQVANR S465
Guy11–0
Guy11–1
Guy11–1.5
Guy11–2
Guy11–4
Guy11–6
dpmk1–0
dpmk1–1
dpmk1–1.5
dpmk1–2
dpmk1–4
dpmk1–6