

YLR008C	153	KINEAKDFLEKREISK*----	168
gi 85079621 ref XP_956385.1	95	NEAKDFLEKSV-----	105
gi 66810570 ref XP_638992.1	101	NEAKDFLSSKNSN-----	113
gi 17509089 ref NP_491662.1 Ce	101	NEAKDFLSSKNS-----	112
Ba 0159_0032	195	AKINLAKELLLMGRK*----	209
gi 21667103 ref NP_660304.1	104	NEAKDFLSSKNS-----	116
CMP334C CMP334CT	96	NEAKDFLLGKTKGSRPTAGV-	113
g9834 g9834.t.1	97	NEAKDFLLGKTKGSRPTAGV-	114
Vocar20006154m.g Vocar20006154m	97	NEAKDFLLGKTKGSRPTAGV-	114
GRMZM2G029385 GRMZM2G029385.T01	115	NEAKDFLLGKTKGSRPTAGV-	132
LOC_Os03g56540 LOC_Os03g56540.3	80	NEAKDFLLGKTKGSRPTAGV-	96
Potri.012G038800 Potri.012G038800.1	95	NEAKDFLLGKTKGSRPTAGV-	112
PGSC0003DMG400027048 PGSC0003DMT40006957	92	NEAKDFLLGKTKGSRPTAGV-	109
Glyma1g23760 Glyma1g23760.1	90	NEAKDFLLGKTKGSRPTAGV-	107
Cucsa.308220 Cucsa.308220.1	95	NEAKDFLLGKTKGSRPTAGV-	112
Eucgr.G02806 Eucgr.G02806.1	93	NEAKDFLLGKTKGSRPTAGV-	110
Bra023030 Bra023030	98	NEAKDFLLGKTKGSRPTAGV-	115
Carubv10024357m.g Carubv10024357m	95	NEAKDFLLGKTKGSRPTAGV-	112
AT2G35795	95	NEAKDFLLGKTKGSRPTAGV-	112
GSVIVG01025634001 GSVIVT01025634001	112	NEAKDFLLGKTKGSRPTAGV-	129
29740.t000014 29740.m000482	95	NEAKDFLLGKTKGSRPTAGV-	112
Potri.008G043500 Potri.008G043500.1	95	NEAKDFLLGKTKGSRPTAGV-	112
Glyma05g16850 Glyma05g16850.1	95	NEAKDFLLGKTKGSRPTAGV-	112
GSVIVG01016322001 GSVIVT01016322001	48	EAKDFLLGKTKGSRPTAGV-	64
GSVIVG01029410001 GSVIVT01029410001	147	KINEAKDFVLMGRKCTGSAF*	166
Eucgr.H00599 Eucgr.H00599.1	98	NEAKDFLLGKTKGSRPTAGV-	115
420475 420475	97	NEAKDFLLGKTKGSRPTAGV-	114
LOC_Os01g06454 LOC_Os01g06454.1	95	NEAKDFLLGKTKGSRPTAGV-	112
GRMZM2G040587 GRMZM2G040587.T01	95	NEAKDFLLGKTKGSRPTAGV-	112
GRMZM2G038950 GRMZM2G038950.T02	95	NEAKDFLLGKTKGSRPTAGV-	112
LOC_Os07g09450 LOC_Os07g09450.1	95	NEAKDFLLGKTKGSRPTAGV-	112
Phpat.003G112600 Phpat.003G112600.1	95	NEAKDFLLGKTKGSRPTAGV-	112
Bra034080 Bra034080	95	NEAKDFLLGKTKGSRPTAGV-	112
Carubv10015707m.g Carubv10015707m	95	NEAKDFLLGKTKGSRPTAGV-	112
AT3G09700	95	NEAKDFLLGKTKGSRPTAGV-	112
Bra005744 Bra005744	124	NEAKDFLLGKTKGSRPTAGV-	142
Bra009555 Bra009555	91	NEAKDFLLGKTKGSRPTAGV-	108
Bra028855 Bra028855	95	NEAKDFLLGKTKGSRPTAGV-	112
Carubv10002291m.g Carubv10002291m	95	NEAKDFLLGKTKGSRPTAGV-	112
AT5G03030	95	NEAKDFLLGKTKGSRPTAGV-	112
Eucgr.A01173 Eucgr.A01173.1	96	NEAKDFLLGKTKGSRPTAGV-	112
tr 137234 137234_MEDTR	93	NEAKDFLLGKTKGSRPTAGV-	110
Glyma1g05710 Glyma1g05710.1	93	NEAKDFLLGKTKGSRPTAGV-	110
Glyma1g20060 Glyma1g20060.1	93	NEAKDFLLGKTKGSRPTAGV-	110
PGSC0003DMG400001767 PGSC0003DMT40000443	95	NEAKDFLLGKTKGSRPTAGV-	112
Cucsa.065620 Cucsa.065620.1	95	NEAKDFLLGKTKGSRPTAGV-	112
Potri.006G131200 Potri.006G131200.1	95	NEAKDFLLGKTKGSRPTAGV-	112
PGSC0003DMG400017591 PGSC0003DMT40004533	95	NEAKDFLLGKTKGSRPTAGV-	112