

Locus_tag	C	Locus_tag	C	Product	Gene	BH O. sativa	E-value
MbP032N20cg010	N-f	-		Putative D-3-phosphoglycerate dehydrogenase precursor		LOC_Os06g44460	4.5E-58
MbP032N20cg020		MbP036B13g010	C-f	Putative aspartate carbamoyltransferase		LOC_Os08g15030	4.3E-154
MbP032N20cg025		MbP036B13g020		Histone H4	H4	LOC_Os01g61920	5.0E-56
MbP032N20cg030		MbP036B13g030		Conserved hypothetical protein		LOC_Os08g14620	2.1E-230
MbP032N20cg040	N-f	MbP036B13g040	N-f	Putative fasciclin-like arabinogalactan protein precursor		LOC_Os02g26320	5.5E-55
MbP032N20cg050		MbP036B13g050		Chlorophyll synthase	VTE2	LOC_Os06g44840	1.9E-145
MbP032N20cg070		MbP036B13g060		Hypothetical protein		No similarity found	
MbP032N20cg080		MbP036B13g070		Plasma membrane ATPase	PMA	LOC_Os04g56160	0
MbP032N20cg090		MbP036B13g080		Conserved hypothetical protein		LOC_Os04g56309	9.1E-14
MbP032N20cg100	N-r	MbP036B13g090	N-r	Villin N-terminal remnant		LOC_Os06g44890	2.9E-18
MbP032N20cg105		MbP036B13g100		Conserved hypothetical protein		LOC_Os06g44900	5.4E-168
MbP032N20cg110		MbP036B13g110		Conserved hypothetical protein		LOC_Os08g40500	1.1E-13
MbP032N20cg120		MbP036B13g120		Putative WRKY transcription factor		LOC_Os08g13840.2	7.5E-60
MbP032N20cg140		MbP036B13g140		Conserved hypothetical protein		LOC_Os02g07920	4.1E-52
MbP032N20cg150		MbP036B13g150		Conserved hypothetical protein		LOC_Os08g13350	1.4E-221
MbP032N20cg160		-		Disease resistance protein (CC-NBS-LRR)	RGA08A	LOC_Os11g45050	2.3E-163
MbP032N20cg165	r	-		RGA08 remnant	RGA08B	LOC_Os11g45180	2.1E-55
MbP032N20cg170	r	-		RGA08 remnant	RGA08C	LOC_Os11g45050	1.5E-69
MbP032N20cg180		-		Disease resistance protein (CC-NBS-LRR)	RGA08D	LOC_Os11g44960	3.0E-166
-		MbP036B13g160	ψ	Disease resistance protein (CC-NBS-LRR)	RGA08S	LOC_Os11g45050	2.0E-146
-		MbP036B13g165	C-r	MTERF C-terminal remnant	MTERF5	LOC_Os09g38720	2.8E-41
-		MbP036B13g170		Disease resistance protein (CC-NBS-LRR)	RGA08T	LOC_Os11g45180	3.4E-154
-		MbP036B13g180	N-r	MTERF N-terminal remnant	MTERF6	LOC_Os09g38720	1.2E-52
-		MbP036B13g190		Disease resistance protein (CC-NBS-LRR)	RGA08U	LOC_Os11g44960	1.1E-163
MbP032N20cg190	ψ	MbP036B13g260	ψ	MTERF domain-containing protein, mitochondrial	MTERF1	LOC_Os09g38720	2.9E-99
MbP032N20cg200	ψ	MbP036B13g270	ψ	Disease resistance protein (CC-NBS-LRR)	RGA08E	LOC_Os01g05620	1.4E-118
MbP032N20cg205	r	MbP036B13g275	r	Gag-Pol remnant			
MbP032N20cg210	ψ	-		Disease resistance protein (CC-NBS-LRR)	RGA08F	LOC_Os11g45090	7.5E-155
MbP032N20cg220	N-f	-		Putative Gag-Pol polyprotein			
MbP032N20cg225	C-r	MbP036B13g195	C-r	Gag-Pol C-terminal remnant			
MbP032N20cg240		MbP036B13g200		Disease resistance protein (CC-NBS-LRR)	RGA08G	LOC_Os11g45180	6.4E-169
MbP032N20cg250	ψ	MbP036B13g210	ψ	Disease resistance protein (CC-NBS-LRR)	RGA08H	LOC_Os11g45180	1.1E-139
MbP032N20cg260	C-r	MbP036B13g230	C-r	RPA1 C-terminal remnant	RPA1	LOC_Os06g40950	1.3E-68
MbP032N20cg280		MbP036B13g250		Disease resistance protein (CC-NBS-LRR)	RGA08I	LOC_Os11g45090	7.5E-163
MbP032N20cg285	ψ	-		Putative Gag-Pol polyprotein			
MbP032N20cg290		-		Disease resistance protein (CC-NBS-LRR)	RGA08J	LOC_Os11g45180	1.2E-164
-		MbP036B13g280	N-r	MTERF N-terminal remnant	MTERF7	LOC_Os09g38720	1.2E-47
-		MbP036B13g290	ψ	Disease resistance protein (CC-NBS-LRR)	RGA08V	LOC_Os11g45180	4.3E-165
MbP032N20cg300		MbP036B13g300		Disease resistance protein (CC-NBS-LRR)	RGA08K	LOC_Os11g45050	1.6E-167
MbP032N20cg310		-		Disease resistance protein (CC-NBS-LRR)	RGA08L	LOC_Os11g47780	2.2E-158
MbP032N20cg320		-		Disease resistance protein (CC-NBS-LRR)	RGA08M	LOC_Os11g45180	4.0E-163
MbP032N20cg330	ψ	-		MTERF domain-containing protein, mitochondrial	MTERF2	LOC_Os09g38720	1.8E-96
MbP032N20cg340	ψ	-		Disease resistance protein (CC-NBS-LRR)	RGA08N	LOC_Os11g13940	5.4E-141
MbP032N20cg360		-		Disease resistance protein (CC-NBS-LRR)	RGA08O	LOC_Os11g45180	1.0E-166
MbP032N20cg380		MbP036B13g310	ψ	Disease resistance protein (CC-NBS-LRR)	RGA08P	LOC_Os09g13820	3.4E-159
-		MbP036B13g315	C-r	MTERF C-terminal remnant	MTERF8	LOC_Os09g38720	1.8E-20
-		MbP036B13g320	N-f	Putative Gag-Pol polyprotein			
MbP032N20cg390	f	MbP036B13g330	N-f	Putative Gag-Pol polyprotein			
MbP032N20cg395	C-r	MbP036B13g325	C-r	MTERF C-terminal remnant	MTERF3	LOC_Os09g38720	9.5E-32
MbP032N20cg400	r	MbP036B13g340	r	RGA08 remnant	RGA08Q	LOC_Os11g45180	1.3E-60
MbP032N20cg410	N-r	-		RGA08 N-terminal remnant	RGA08R	LOC_Os11g45060	2.6E-108
-		MbP036B13g345	r	RGA08 remnant	RGA08W	LOC_Os09g13820	5.5E-77
-		MbP036B13g350		Disease resistance protein (CC-NBS-LRR)	RGA08X	LOC_Os11g45180	7.8E-167
MbP032N20cg420		MbP036B13g360		MTERF domain-containing protein, mitochondrial	MTERF4	LOC_Os09g38720	2.1E-145
MbP032N20cg440	N-f	MbP036B13g380		Putative serine/threonine-protein kinase		LOC_Os06g45280	5.5E-173
-		MbP036B13g390	N-f	Pseudouridylate synthase	PUS	LOC_Os06g45250	1.8E-63