

Integrative meta-analysis of protein interaction data identified multiple GID/MRCTLH protein complexes in plants

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Table S1.- Protein interactors of the Arabidopsis proteins homologues of the proteins in the yeast and human GID/MRCTLH complex based on The Arabidopsis Interactions Viewer (Geisler-Lee et al. 2007; http://bar.utoronto.ca/interactions/cgi-bin/arabidopsis_interactions_viewer.cgi).

Gene 1	Gene 2	Definition gene 2
At1g61150 (Twa1)	At1g16610	ARGININE/SERINE-RICH 45
At1g61150 (Twa1)	At3g55070 (MAEA)	LisH/CRA/RING-U-box domains-containing protein
At1g61150 (Twa1)	At4g09340 (RanBPM)	SPla/Ryanodine receptor (SPRY) domain-containing protein
At1g61150 (Twa1)	At4g33200	MYOSIN XI 1
At1g61150 (Twa1)	At4g37880 (RMD5)	LisH/CRA/RING-U-box domains-containing protein
At1g61150 (Twa1)	At5g08560 (WDR26)	LisH and RanBPM domains / WD-40 repeat containing protein
At1g61150 (Twa1)	At5g43900	MYOSIN 2
At3g55070 (MAEA)	At1g09100	26S PROTEASOME AAA-ATPASE SUBUNIT RPT5B
At3g55070 (MAEA)	At1g59900	PYRUVATE DEHYDROGENASE COMPLEX E1 ALPHA SUBUNIT
At3g55070 (MAEA)	At1g61150 (Twa1)	LisH and RanBPM domains containing protein
At3g55070 (MAEA)	At2g22690 (Rmd5)	LisH/CRA/RING-U-box domains-containing protein
At3g55070 (MAEA)	At2g29540	RNA polymerase 14 kDa subunit
At3g55070 (MAEA)	At2g39630	A putative dolichyl-phosphate β -glucosyltransferase
At3g55070 (MAEA)	At2g40110	Yippee family putative zinc-binding protein
At3g55070 (MAEA)	At3g05530	REGULATORY PARTICLE TRIPLE-A ATPASE 5A, RPT5A
At3g55070 (MAEA)	At3g29800	P-loop containing nucleoside triphosphate hydrolases
At3g55070 (MAEA)	At4g09300	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At3g55070 (MAEA)	At4g27740	Yippee family putative zinc-binding protein
At3g55070 (MAEA)	At4g32410	CELLULOSE SYNTHASE 1
At3g55070 (MAEA)	At5g08560 (WDR26)	LisH/CRA/RING-U-box domains-containing protein ,transducin family protein / WD-40 repeat family protein
At3g55070 (MAEA)	At5g45430	Protein kinase superfamily protein
At4g37880 (RMD5)	At1g10320	Zinc finger C-x8-C-x5-C-x3-H type family protein
At4g37880 (RMD5)	At1g30090	Galactose oxidase/kelch repeat superfamily protein
At4g37880 (RMD5)	At1g61150 (Twa1)	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At4g37880 (RMD5)	At1g70290	Trehalose-6-phosphatase synthase S8
At4g37880 (RMD5)	At1g80830	Natural resistance-associated macrophage protein 1
At4g37880 (RMD5)	At3g14120	Undetermined
At4g37880 (RMD5)	At3g62030	Nuclear-encoded chloroplast stromal cyclophilin CYP20-3
At4g37880 (RMD5)	At4g16830	Hyaluronan / mRNA binding domain-containing protein
At4g37880 (RMD5)	At4g33200	MYOSIN XI 1
At4g37880 (RMD5)	At4g37880 (RMD5)	LisH/CRA/RING-U-box domains-containing protein ,LisH/CRA/RING-U-box domains-containing protein
At4g37880 (RMD5)	At5g48010	Thalianol synthase 1
At1g35470 (RanBPM)	At4g10570	UBIQUITIN-SPECIFIC PROTEASE 9, UBP9
At1g35470 (RanBPM)	At4g18040	EUKARYOTIC TRANSLATION INITIATION FACTOR 4E1
At1g35470 (RanBPM)	At5g14080	Tetratricopeptide repeat (TPR)-like superfamily protein
At1g35470 (RanBPM)	At5g20020	RAN2, RAS-RELATED GTP-BINDING NUCLEAR PROTEIN 2
At1g35470 (RanBPM)	At5g55190	RAN GTPASE 3, RAN3
At1g35470 (RanBPM)	At5g64320	Pentatricopeptide repeat (PPR) superfamily protein
At1g35470 (RanBPM)	At1g61150 (Twa1)	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At1g35470 (RanBPM)	At3g55070 (MAEA)	LisH/CRA/RING-U-box domains-containing protein
At1g35470 (RanBPM)	At4g37880 (RMD5)	LisH/CRA/RING-U-box domains-containing protein
At1g35470 (RanBPM)	At5g08560 (WDR26)	LisH and RanBPM domains / WD-40 repeat containing protein

At1g35470 (RanBPM)	At5g43920 (WDR26)	LisH and RanBPM domains / WD-40 repeat containing protein
At5g08560 (WDR26)	At1g61150 (Twa1)	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At5g08560 (WDR26)	At1g78900	VACUOLAR ATP SYNTHASE SUBUNIT A, VHA-A
At5g08560 (WDR26)	At2g22690 (Rmd5)	LisH/CRA/RING-U-box domains-containing protein
At5g08560 (WDR26)	At2g39760	MATH-BTB domain proteins (BPMs)
At5g08560 (WDR26)	At2g40110	Yippee family putative zinc-binding protein
At5g08560 (WDR26)	At3g06720	IMPA1, IMPORTIN ALPHA
At5g08560 (WDR26)	At3g55070 (MAEA)	LisH/CRA/RING-U-box domains-containing protein
At5g08560 (WDR26)	At4g02150	IMPORTIN ALPHA 3
At5g08560 (WDR26)	At4g09300	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At5g08560 (WDR26)	At4g16143	IMPORTIN ALPHA ISOFORM 2
At5g08560 (WDR26)	At4g24900	TITAN-LIKE, TTL
At5g08560 (WDR26)	At5g19000	nucleus,transducin family protein / WD-40 repeat family protein, BTB-POZ AND MATH DOMAIN 1, BPM1
At5g08560 (WDR26)	AT1G49450	Transducin/WD40 repeat-like superfamily protein
At5g08560 (WDR26)	At1g35470 (RanBPM)	RAN-BINDING PROTEIN IN THE MICROTUBULE-ORGANIZING CENTRE, RANBPM, SPRY, LisH, CTLH and CRA domain-containing proteins
At5g43920 (WDR26)	AT1G78070	Transducin/WD40 repeat-like superfamily protein
At5g43920 (WDR26)	AT5G50120	Transducin/WD40 repeat-like superfamily protein
At5g43920 (WDR26)	At1g35470 (RanBPM)	RAN-BINDING PROTEIN IN THE MICROTUBULE-ORGANIZING CENTRE, RANBPM, SPRY, LisH, CTLH and CRA domain-containing proteins
At4g09340 (RanBPM)	At1g61150 (Twa1)	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At4g09340	At4g26630	DEK-DOMAIN CONTAINING PROTEIN 3, DEK3
At2g22690 (Rmd5)	At1g59900	PYRUVATE DEHYDROGENASE COMPLEX E1 ALPHA SUBUNIT
At2g22690 (Rmd5)	At2g40110	Yippee family putative zinc-binding protein
At2g22690 (Rmd5)	At3g55070 (MAEA)	LisH/CRA/RING-U-box domains-containing protein
At2g22690 (Rmd5)	At4g27740	Yippee family putative zinc-binding protein
At2g22690 (Rmd5)	At5g08560 (WDR26)	LisH and RanBPM domains / WD-40 repeat containing protein
At2g22690 (Rmd5)	At2g38880	nucleus,zinc ion binding ,ATHAP3
At2g22690 (Rmd5)	At5g27970	zinc ion binding ,ARM repeat superfamily protein
At2g22690 (Rmd5)	At5g65180	zinc ion binding ,ENTH/VHS family protein
At2g22690 (Rmd5)	At1g06790	zinc ion binding ,RNA polymerase Rpb7 N-terminal domain-containing protein
At2g22690 (Rmd5)	At4g33100	Mitochondrial distribution/morphology family 35
At2g22690 (Rmd5)	At5g52640	HEAT SHOCK PROTEIN 83
At2g22690 (Rmd5)	AT5G67320 (WDR26)	HIGH EXPRESSION OF OSMOTICALLY RESPONSIVE GENES 15, HOS15
At2g22690 (Rmd5)	At2g13370	CHROMATIN REMODELING 5, DEAD-like helicase
At4g09300 (TWA1)	At3g55070 (MAEA)	LisH/CRA/RING-U-box domains-containing protein
At4g09300 (TWA1)	At5g08560 (WDR26)	LisH and RanBPM domains / WD-40 repeat containing protein
At4g09300 (TWA1)	At4g09300	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At4g09300 (TWA1)	At5g20490	MYOSIN XI K
At4g09300 (TWA1)	At5g09630 (Rmd5)	LisH/CRA/RING-U-box domains-containing protein
AT5G67320 (WDR26)	At1g02690	IMPA-6, IMPORTIN ALPHA ISOFORM 6
AT5G67320 (WDR26)	At1g03190	ULTRAVIOLET HYPERSENSITIVE 6, UVH6
AT5G67320 (WDR26)	At1g08780	ABI3-INTERACTING PROTEIN 3, AIP3, PFD4, PREFOLDIN 4
AT5G67320 (WDR26)	At1g09020	ATSNF4, HOMOLOG OF YEAST SUCROSE NONFERMENTING 4, KIN β , SNF4
AT5G67320 (WDR26)	At1g11890	ATSEC22, SEC22, SECRETION 22
AT5G67320 (WDR26)	At1g17760	ARABIDOPSIS THALIANA CLEAVAGE STIMULATION FACTOR 77, ATCSTF77, CSTF77
AT5G67320 (WDR26)	At1g18940	Nodulin-like
AT5G67320 (WDR26)	At1g29150	ATS9, NON-ATPASE SUBUNIT 9, REGULATORY PARTICLE NON-ATPASE 6, RPN6
AT5G67320 (WDR26)	At1g29990	PFD6, PREFOLDIN 6
AT5G67320 (WDR26)	At1g30070	SGS domain-containing protein
AT5G67320 (WDR26)	At1g52740	HISTONE H2A PROTEIN 9, HTA9
AT5G67320 (WDR26)	At1g59890	SIN3-LIKE 5, SNL5
AT5G67320 (WDR26)	At1g64790	ILA, ILITHYIA
AT5G67320 (WDR26)	At1g77460	CELLULOSE SYNTHASE INTERACTIVE 3, CSI3
AT5G67320 (WDR26)	At1g78900	VACUOLAR ATP SYNTHASE SUBUNIT A, VHA-A
AT5G67320 (WDR26)	At1g79350	EMB1135, EMBRYO DEFECTIVE 1135
AT5G67320 (WDR26)	At1g80510	Amino acid transporter
AT5G67320 (WDR26)	At2g13370	CHROMATIN REMODELING 5, DEAD-like helicase
AT5G67320 (WDR26)	At2g13440	Glucose-inhibited division family A protein
AT5G67320 (WDR26)	At2g15430	Non-catalytic subunit of nuclear DNA-dependent RNA polymerases II, IV and V
AT5G67320 (WDR26)	At2g16600	CYCLOPHILIN 19
AT5G67320 (WDR26)	At2g18000	TBP-ASSOCIATED FACTOR 14
AT5G67320 (WDR26)	At2g18510	EMB2444, EMBRYO DEFECTIVE 2444
AT5G67320 (WDR26)	At2g22690 (Rmd5)	LisH/CRA/RING-U-box domains-containing protein
AT5G67320 (WDR26)	At2g36740	ATSWC2, sequence-specific DNA binding transcription factor
AT5G67320 (WDR26)	At2g43810	SM-LIKE 6B, Small nuclear ribonucleoprotein
AT5G67320 (WDR26)	At2g44150	ASHH3, HISTONE-LYSINE N-METHYLTRANSFERASE
AT5G67320 (WDR26)	At3g01090	SNF1-RELATED PROTEIN KINASE 1.1, SNRK1.1
AT5G67320 (WDR26)	At3g03600	RIBOSOMAL PROTEIN S2, RPS2
AT5G67320 (WDR26)	At3g04710	TETRATRICOPEPTIDE REPEAT 10, TPR10
AT5G67320 (WDR26)	At3g06720	IMPA1, IMPORTIN ALPHA
AT5G67320 (WDR26)	At3g12810	PHOTOPERIOD-INDEPENDENT EARLY FLOWERING 1, PIE1
AT5G67320 (WDR26)	At3g21060	RBBP5 LIKE, RBL
AT5G67320 (WDR26)	At3g22480	PDF2, PREFOLDIN 2
AT5G67320 (WDR26)	At3g22590	PLANT HOMOLOGOUS TO PARAFIBROMIN
AT5G67320 (WDR26)	At3g52250	Encodes a protein with a putative role in mRNA splicing

AT5G67320 (WDR26)	At3g58040	SEVEN IN ABSENTIA OF ARABIDOPSIS 2, SINAT2
AT5G67320 (WDR26)	At4g01370	TMPK4, MAP KINASE 4
AT5G67320 (WDR26)	At4g04940	Transducin
AT5G67320 (WDR26)	At4g05190	KINESIN 5
AT5G67320 (WDR26)	At4g22140	EARLY BOLTING IN SHORT DAYS, EBS
AT5G67320 (WDR26)	At4g27180	KINESIN 2
AT5G67320 (WDR26)	At4g38740	ROC1, ROTAMASE CYP 1
AT5G67320 (WDR26)	At5g02470	DPA
AT5G67320 (WDR26)	At5g02560	HISTONE H2A 12, HTA12
AT5G67320 (WDR26)	At5g02570	Histone
AT5G67320 (WDR26)	At5g03415	DPB
AT5G67320 (WDR26)	At5g09230	SIRTUIN 2
AT5G67320 (WDR26)	At5g09740	HISTONE ACETYLTRANSFERASE OF THE CBP FAMILY 11
AT5G67320 (WDR26)	At5g10260	RAB GTPASE HOMOLOG H1E
AT5G67320 (WDR26)	At5g10400	HISTONE 3.1
AT5G67320 (WDR26)	At5g10980	HISTONE 3.3
AT5G67320 (WDR26)	At5g14120	Major facilitator superfamily protein
AT5G67320 (WDR26)	At5g15770	GLUCOSE-6-PHOSPHATE ACETYLTRANSFERASE 1, GNA1
AT5G67320 (WDR26)	At5g19330	ARM REPEAT PROTEIN INTERACTING WITH ABF2
AT5G67320 (WDR26)	At5g22220	E2F TRANSCRIPTION FACTOR 1, E2F1, E2FB
AT5G67320 (WDR26)	At5g22880	HISTONE B2
AT5G67320 (WDR26)	At5g23290	PREFOLDIN 5
AT5G67320 (WDR26)	At5g41880	POLA3, DNA primase, small subunit
AT5G67320 (WDR26)	At5g42190	SKP-LIKE 2
AT5G67320 (WDR26)	At5g45600	TBP-ASSOCIATED FACTOR 14B
AT5G67320 (WDR26)	At5g48120	HOMOLOG OF YEAST MET18, ARM repeat superfamily protein
AT5G67320 (WDR26)	At5g49510	PREFOLDIN 3
AT5G67320 (WDR26)	At5g53360	TRAF-like superfamily protein
AT5G67320 (WDR26)	At5g53480	HOMOLOG OF HUMAN KPNB1, ARM repeat superfamily protein
AT5G67320 (WDR26)	At5g54670	KINESIN 3
AT5G67320 (WDR26)	At5g61070	HISTONE DEACETYLASE OF THE RPD3/HDA1 SUPERFAMILY 18
AT5G67320 (WDR26)	At5g63110	HISTONE DEACETYLASE 6, RNA-MEDIATED TRANSCRIPTIONAL SILENCING 1
AT5G67320 (WDR26)	At5g66020	IMPAIRED IN BABA-INDUCED STERILITY 2, SACIB, SUPPRESSOR OF ACTIN 1B
AT5G67320 (WDR26)	At5g59970	Histone
AT5G67320 (WDR26)	At1g07820	Histone
AT5G67320 (WDR26)	At5g52310	Low-temperature-induced protein 78
AT5G67320 (WDR26)	At3g45930	Histone
AT5G67320 (WDR26)	At3g46320	Histone
AT5G67320 (WDR26)	At1g07660	Histone
AT5G67320 (WDR26)	At5g59690	Histone
AT5G67320 (WDR26)	At2G28740	HISTONE H4
AT5G67320 (WDR26)	At3g53730	Histone
At5g09630 (Rmd5)	At1g07930	GTP binding Elongation factor Tu family protein
At5g09630 (Rmd5)	At1g09640	Translation elongation factor EF1B, gamma chain
At5g09630 (Rmd5)	At1g10320	Zinc finger C-x8-C-x5-C-x3-H type family protein
At5g09630 (Rmd5)	At1g11530	C-TERMINAL CYSTEINE RESIDUE IS CHANGED TO A SERINE 1
At5g09630 (Rmd5)	At1g13320	PROTEIN PHOSPHATASE 2A SUBUNIT A3
At5g09630 (Rmd5)	At1g15210	ATP-BINDING CASSETTE G35
At5g09630 (Rmd5)	At1g22780	40S RIBOSOMAL PROTEIN S18
At5g09630 (Rmd5)	At1g27970	NUCLEAR TRANSPORT FACTOR 2B
At5g09630 (Rmd5)	At1g33090	MATE efflux family protein
At5g09630 (Rmd5)	At1g33110	MATE efflux family protein
At5g09630 (Rmd5)	At1g50310	SUGAR TRANSPORTER 9
At5g09630 (Rmd5)	At1g51770	Core-2/I-branching beta-1,6-N-acetylglucosaminyltransferase family protein
At5g09630 (Rmd5)	At1g53900	Eukaryotic translation initiation factor 2B (eIF-2B)
At5g09630 (Rmd5)	At1g62880	Cornichon family protein
At5g09630 (Rmd5)	At1g64480	CALCINEURIN B-LIKE PROTEIN 8, CBL8
At5g09630 (Rmd5)	At1g71280	DEA(D/H)-box RNA helicase family protein
At5g09630 (Rmd5)	At1g72480	Lung seven transmembrane receptor family protein
At5g09630 (Rmd5)	At1g76720	Eukaryotic translation initiation factor 2 (eIF-2) family protein
At5g09630 (Rmd5)	At1g78970	LUPEOL SYNTHASE 1
At5g09630 (Rmd5)	At2g03667	Asparagine synthase family protein
At5g09630 (Rmd5)	At2g13650	GOLGI NUCLEOTIDE SUGAR TRANSPORTER 1
At5g09630 (Rmd5)	At2g14120	DRP3B, DYNAMIN RELATED PROTEIN
At5g09630 (Rmd5)	At2g19860	HEXOKINASE 2
At5g09630 (Rmd5)	At2g20280	Zinc finger C-x8-C-x5-C-x3-H type family protein
At5g09630 (Rmd5)	At2g20450	Ribosomal protein L14
At5g09630 (Rmd5)	At2g24040	Low temperature and salt responsive protein family
At5g09630 (Rmd5)	At2g29690	ANTHRANILATE SYNTHASE 2
At5g09630 (Rmd5)	At2g31170	CYSTEINYL T-RNA SYNTHETASE, FIONA
At5g09630 (Rmd5)	At2g37470	Histone
At5g09630 (Rmd5)	At2g39290	PHOSPHATIDYLGLYCEROLPHOSPHATE SYNTHASE 1
At5g09630 (Rmd5)	At2g46860	PYROPHOSPHORYLASE 3
At5g09630 (Rmd5)	At2g47640	Small nuclear ribonucleoprotein family protein
At5g09630 (Rmd5)	At3g01280	VDAC1, VOLTAGE DEPENDENT ANION CHANNEL 1
At5g09630 (Rmd5)	At3g05960	SUGAR TRANSPORTER 6
At5g09630 (Rmd5)	At3g06470	GNS1/SUR4 membrane protein family
At5g09630 (Rmd5)	At3g08650	ZIP metal ion transporter family
At5g09630 (Rmd5)	At3g08950	HOMOLOGUE OF THE COPPER CHAPERONE SCO1
At5g09630 (Rmd5)	At3g11630	2-Cys peroxiredoxin
At5g09630 (Rmd5)	At3g12690	Serine/threonine kinase

At5g09630 (Rmd5)	At3g14120	Undetermined
At5g09630 (Rmd5)	At3g16050	PYRIDOXINE BIOSYNTHESIS 1.2
At5g09630 (Rmd5)	At3g16950	LIPOAMIDE DEHYDROGENASE 1
At5g09630 (Rmd5)	At3g19940	Major facilitator superfamily protein
At5g09630 (Rmd5)	At3g19980	FLOWER-SPECIFIC, PHYTOCHROME-ASSOCIATED PROTEIN PHOSPHATASE 3
At5g09630 (Rmd5)	At3g20390	REACTIVE INTERMEDIATE DEAMINASE A, RIDA
At5g09630 (Rmd5)	At3g20650	mRNA capping enzyme family protein
At5g09630 (Rmd5)	At3g23570	alpha/beta-Hydrolases superfamily protein
At5g09630 (Rmd5)	At3g28710	ATPase, V0/A0 complex, subunit C/D
At5g09630 (Rmd5)	At3g30842	ATP-BINDING CASSETTE G38
At5g09630 (Rmd5)	At3g49010	40S RIBOSOMAL PROTEIN
At5g09630 (Rmd5)	At3g52390	TatD related DNase
At5g09630 (Rmd5)	At3g52580	Ribosomal protein S11 family protein
At5g09630 (Rmd5)	At3g53510	ATP-BINDING CASSETTE G20
At5g09630 (Rmd5)	At3g56190	ALPHA-SOLUBLE NSF ATTACHMENT PROTEIN 2
At5g09630 (Rmd5)	At3g62770	AUTOPHAGY 18A
At5g09630 (Rmd5)	At4g02120	CTP synthase family protein
At5g09630 (Rmd5)	At4g09300 (TWA1)	LisH/CRA/RING-U-box domains-containing protein ,LisH and RanBPM domains containing protein
At5g09630 (Rmd5)	At4g17770	TREHALOSE -6-PHOSPHATASE SYNTHASE S5
At5g09630 (Rmd5)	At4g24400	CBL-INTERACTING PROTEIN KINASE 8
At5g09630 (Rmd5)	At4g30990	ARM repeat superfamily protein
At5g09630 (Rmd5)	At4g32470	Cytochrome bd ubiquinol oxidase, 14kDa subunit
At5g09630 (Rmd5)	At4g33070	PYRUVATE DECARBOXYLASE 1
At5g09630 (Rmd5)	At4g36860	LIM domain-containing protein
At5g09630 (Rmd5)	At4g39330	CINNAMYL ALCOHOL DEHYDROGENASE 9
At5g09630 (Rmd5)	At5g05420	FKBP-like peptidyl-prolyl cis-trans isomerase family protein
At5g09630 (Rmd5)	At5g09630 (Rmd5)	LisH/CRA/RING-U-box domains-containing protein
At5g09630 (Rmd5)	At5g12350	Regulator of chromosome condensation (RCC1) family with FYVE zinc finger domain
At5g09630 (Rmd5)	At5g13050	5-FCL, 5-FORMYLTETRAHYDROFOLATE CYCLOLIGASE
At5g09630 (Rmd5)	At5g19150	pfkB-like carbohydrate kinase family protein
At5g09630 (Rmd5)	At5g19820	EMBRYO DEFECTIVE 2734
At5g09630 (Rmd5)	At5g23740	RIBOSOMAL PROTEIN S11-BETA
At5g09630 (Rmd5)	At5g41190	NOB1, cytoplasmic protein with RNA endonuclease activity
At5g09630 (Rmd5)	At5g41480	EMBRYO DEFECTIVE 9
At5g09630 (Rmd5)	At5g43370	PHOSPHATE TRANSPORTER 1
At5g09630 (Rmd5)	At5g44500	Small nuclear ribonucleoprotein family protein
At5g09630 (Rmd5)	At5g45420	MEMBRANE ANCHORED MYB
At5g09630 (Rmd5)	At5g47880	EUKARYOTIC RELEASE FACTOR 1-1
At5g09630 (Rmd5)	At5g59870	HISTONE H2A 6
At5g09630 (Rmd5)	At5g60390	GTP binding Elongation factor Tu family protein
At5g09630 (Rmd5)	At5g62300	Ribosomal protein S10p/S20e family protein
At5g09630 (Rmd5)	At1g76720	Eukaryotic translation initiation factor 2