

**Supplementary Table 4: Alignment of 484 lipobox-containing proteins from halophilic archaea**

The N-terminal region of the 484 lipoproteins from 6 halophilic archaea is shown. Proteins are named by their ordered locus tag obtained upon genome annotation (code). Each of the proteins was positive in at least two of the three predictors (lipobox prediction: A: Prosite PS51257; B: lipoP; C: predLipo). The first 400 proteins are TatFind positive ("T"), the last 84 are TatFind negative ("‐‐‐‐‐"). All TatFind positives and the first 50 of the TatFind negatives have a twin-Arg motif ("RR") while the last 34 TatFind negatives lack a twin-Arg motif ("‐‐‐‐‐"). The twin-Arg motif and the lipobox motif were juxtaposed by introducing a gap of variable length between positions 5 and 6 after the twin-Arg motif. Spaces were added on both sides of the twin-Arg motif and of the lipobox to increase readability. If there were three or more consecutive Arg residues in the twin-Arg motif, the positioning is according to TatFind, which is based on the detection of a conserved pair of hydrophobic residues at position +2/+3. Dashes indicate the single gap of variable length introduced between position +5 and +6 after the twin-Arg motif and also the region in front of the initiator-Met.

code	N-terminal sequence	lipobox prediction	TatFind	twin-Arg
pNG2038	-----MTERT RR QLLAG-----IGTSLTLA TAGC GGNNSGGSN	A,B,C	T	RR
pNG3012	-----MK RR NLLSA-----LGTSVA LAGC STSTNSTETN	A,B,C	T	RR
pNG5085	-----MTHKNIPRNSSID RR TVLGL-----LSGSLAA LAGC SGGLSDSGDG	A,B,C	T	RR
pNG6063	-----ME RR QVLKT---AGILATGGVTG LAGC SSSGNGDGGG	A,B,C	T	RR
pNG6067	-----MTTRFGAPGTGLS RR EFLAA-----TGATGIAS LAGC SAGGANEPAA	A,B,C	T	RR
pNG7023	-----MTNSN RR KFLKA-----TGVGLLGG LAGC TRGGDSSGGD	A,B,C	T	RR
pNG7073	-----MD RR TFLAA-----SGLTVAGT LAGC SAPSDNQQRD	A,B,C	T	RR
pNG7088	-----MPSTS RR TFLGM-----VATGTVAS VAGC SSTCPDEGAP	A,B,C	T	RR

pNG7122	-----MNRSSVS RR QFLGA-----SGAALVTG LAGC SAGEGGTSTD	A,B,C	T	RR
pNG7185	-----MHQLH RR QLLGV-----VCSGLTV LAGC GSDQSGNSSE	A,B,C	T	RR
pNG7190	-----MDKADDNSNGLT RR DYLSG-----GSVLLGTSI VAGC SGDGGESPST	A,B,C	T	RR
pNG7250	-----MVNTNRTIS RR EMLGA-----LGATGVTA LAGC SGGSDDTDA	A,B,C	T	RR
pNG7251	-----MVDKKHTRS RR ELLTA-----LGAAGITG LAGC SGGGDGGDGE	A,B,C	T	RR
pNG7384	-----MVPST RR TLLHS-----VCGLATV LAGC SGVFESSGES	A,B,C	T	RR
rrnAC0147	-----MD RR SFLTA-----AGATVSVS FAGC AGLGGSDESS	A,B,C	T	RR
rrnAC0158	-----MN RR QFLAS-----AGLAGTTV LAGC GSLSSQSTRA	A,B,C	T	RR
rrnAC0296	-----MKQLSLT RR AMLGA-----LGTAVAA TAGC QSPGTGSDSG	A,B,C	T	RR
rrnAC0358	-----MFRNDKHSLVSK RR SFLKM-----VGTAGLAG LAGC GGDGSGDGG	A,B,C	T	RR
rrnAC0381	-----MPREARVD RRR YLQA-----IGGTVATLS VAGC QSDSGDSQTL	A,B,C	T	RR
rrnAC0389	-----MET RR TFVRG-----LGLAAGVA LAGC SGGGSDSGD	A,B,C	T	RR
rrnAC0498	-----MPS RR SVLTA-----GTLVA LSGC AALSPASARL	A,B,C	T	RR
rrnAC0508	-----MTNDLD RR TFIRT-----TAAVSGAGL LAGC GGSGGDGGSD	A,B,C	T	RR
rrnAC0566	-----MQNTAT RRR VLGA-----LGAGVVG LSGC VSDDSSMTEQ	A,B,C	T	RR
rrnAC0576	-----MSSDTT RRR ILLGI-----GTGVTVG IAGC SGGGNGGAE	A,B,C	T	RR
rrnAC0830	-----MSSDTN RR AFLKR--TGAVTTVGLLGG LAGC STEQTGGDGG	A,B,C	T	RR
rrnAC0850	-----MTQ-RR-AT RRR FLMG-----CSAIASVG LAGC TGGDESTATD	A,B,C	T	RR
rrnAC0856	-----MTRFT RR GLLAA-----TVGAVGA MAGC AGGSSESESA	A,B,C	T	RR
rrnAC0899	-----MN RR EYIVA-----AGVGLAGT VAGC SGQSEAGSGS	A,B,C	T	RR
rrnAC0998	-----MLRTT RRR LLSA-----LGASLSI LAGC TQFDVESGAE	A,B,C	T	RR
rrnAC1082	-----MD RR TYIGT-----IGSSMAALS LAGC SGSDGGDGNE	A,B,C	T	RR
rrnAC1892	-----MN RRR FLTL-----SGVGVVGA VAGC SSEADSAESV	A,B,C	T	RR
rrnAC1944	-----MEFEST RRR FMQL-----AGTGATVS LAGC NALQGGDSDG	A,B,C	T	RR
rrnAC2421	-----MRNPARDARS RR QYLAA-----LGGIGATA LAGC RGESGTQTEP	A,B,C	T	RR
rrnAC2469	-----MNEHDTRSID RR SVIKA-----AASAGLVG LAGC SG GAPDEGDG	A,B,C	T	RR
rrnAC2580	-----MTNNST RRR FLKT-----AGVAGVAA LAGC GGGGDGGDGG	A,B,C	T	RR
rrnAC2656	-----MSQ RRR TFISS-----VGTATAIG LAGC LGGDGGDGSG	A,B,C	T	RR

rrnAC2761	-----MRNHNIT RR QLLAS-----GGAAATAA LAGC AGGF GGSGQG	A,B,C	T	RR
rrnAC2848	-----M RRRR FIAT-----GAGVGVGL FAGC SGSSD SGGSD	A,B,C	T	RR
rrnAC2863	-----MK RR NYIAS-----LGSLT LAGC ATGE GS GE GE	A,B,C	T	RR
rrnAC3028	-----MRDTSQ-RR-S RR GFLGT-----LALAAVT SAGC SSLGGTQNGQ	A,B,C	T	RR
rrnAC3119	-----MGRPTS RR NVIKI-----AGGALVA LAGC SDTGGSTESD	A,B,C	T	RR
rrnAC3197	-----MQESSNRT RR TFIKS-----AGVIGAAA LAGC SGGSDSSTED	A,B,C	T	RR
rrnAC3198	-----MTERGRT RR AFLGT-----LAATTVGS VAGC QSQFNPLAST	A,B,C	T	RR
rrnAC3204	-----MDRN RR QFLGA-----LTAVVTGT IAGC SGGD SGE SPP	A,B,C	T	RR
rrnAC3220	-----MRDQQQSGTAS RR DYLKA---AGGLAA ASTVG LAGC SGSGSETGTL	A,B,C	T	RR
rrnAC3248	-----M RRR ALLAS-----VPGALAG LAGC SF GTPGSDET	A,B,C	T	RR
rrnAC3267	-----MPIQ RR EFIAA-----IGTGALAT TAGC AQSGEGSGQT	A,B,C	T	RR
rrnAC3299	-----MPRSSSME RR SFLKA-----TGSA AAAA LAGC SSDSDETEGA	A,B,C	T	RR
rrnAC3315	-----M RRR QFLAG-----GTVLLSAA VAGC GHPSVVLMD	A,B,C	T	RR
rrnB0022	-----M RRR QALAL-----VTGSLSL VAGC SSDTTTDLTP	A,B,C	T	RR
rrnB0097	-----MTSSYS RR TLLRT-----AGVAVAGA VGGC GGDSTAQDM	A,B,C	T	RR
rrnB0117	-----MK RR TFLAS-----TGIGLTTA VAGC TTQAGSSDSP	A,B,C	T	RR
rrnB0275	-----M RRR TYIAS-----GTVMASGL LAGC SSGSSESEST	A,B,C	T	RR
rrnB0319	-----MSASTMKQNPVD RR SVLKI-----GATALAAG VAGC SQESSQSTET	A,B,C	T	RR
pNG6086	-----MN RR TFLKQ-----GTALSGGLV LSGC LGRLGFETQS	A,B,-	T	RR
pNG6087	-----MN RR DVLRA-----VGGTSLVG LAGC TGLFETRSAR	A,B,-	T	RR
pNG7039	-----MSRYDMNREGPT RR DYMKY-----SGAVVGGSL LAGC AGQSGSEAAP	A,B,-	T	RR
pNG7040	-----MTDDEKT PHERPT RR DYIKG-----AGIIAGGSL LAGC AGDAVSESTA	A,-,C	T	RR
pNG7041	-----MTDPDRAACEWPT RR DIVRS-----SGALAFGGF LAGC SGESGQESSV	A,-,C	T	RR
pNG7059	-----M RRR TLLKS-----TGAAGTIVG VAGC LGGRLGDSN	A,B,-	T	RR
pNG7234	-----MRPT RR EFLGT-----AAALS LSGC LGGTATPTLT	A,B,-	T	RR
rrnAC0082	-----MDQT RR AVLAG-----VAGL FAGC GTTSQRTTTP	A,B,-	T	RR
rrnAC0299	-----MSSNSNDQSELT RR EYVTC-----GGTVLAGGL LSGC TGSSGSDPSD	-,B,C	T	RR
rrnAC0460	-----MPSS RRR YLRG-----CVATLGLA SAGC LGSNGYIETS	A,B,-	T	RR

rrnAC0510	-----MQRPST RR QFLTG-----TGVAALAI TAGC VSSGSSSEPA	A,B,-	T	RR
rrnAC0702	-----MYSDS RR GVLKK-----CIAAGSLGL SSGC LGTLGGSGDT	A,B,-	T	RR
rrnAC0906	-----MSDSNDMTGRNATQS RRR FLAL-----GSATAATA LTGC SGILSGDDGG	-,B,C	T	RR
rrnAC0915	-----MTDDNSNKRLT RR NALRI-----AGAAGAAS LAGC GGSDGGDGSS	-,B,C	T	RR
rrnAC0957	-----MSRS RR TFLTT-----LAAASVTLP LAGC AGGDEQSADE	-,B,C	T	RR
rrnAC1162	-----MRIS RRR FLAT-----VGAGVTLG GAGC VGDEGATAGG	A,B,-	T	RR
rrnAC1183	-----MARQT RR AVLAA-----LGSGIAA TAGC GA FGQERVDV	A,B,-	T	RR
rrnAC1260	-----MN RR DYLLA-----GATLGTAG LAGC SFLAAAEAPP	A,B,-	T	RR
rrnAC1325	-----MTS RR GLLLRL-----LSALTGAAG LSGC SSLLARQAES	A,B,-	T	RR
rrnAC1365	-----MNRNQT RR KTLAL-----LGSALSIP VAGC SSGGLFGGET	-,B,C	T	RR
rrnAC1430	-----MD RR QFLRA-----AGPAAVAG LAGC LGGGSADTDY	A,B,-	T	RR
rrnAC1436	-----MRT RR TVLVL-----SAGTLC LAGC GSPGDGSNGG	-,B,C	T	RR
rrnAC1509	-----MMTDSPSGGIS RRR FLTS-----SGAIGTLA LAGC VQNTNSGDGS	-,B,C	T	RR
rrnAC1654	-----MN RRR YLTV-----STAALSG LAGC LGDPEYTISS	A,B,-	T	RR
rrnAC1910	-----MPQ RR DLLKT-----VGIAATGL LAGC PSQGSTESNT	A,B,-	T	RR
rrnAC2010	-----MNRT RR ETLGL-----VGGLAATV LGGC LAPGAGGDGG	A,B,-	T	RR
rrnAC2149	-----MAHDSNGLSEFVS RR KFIAT-----TGATSIAA IAGC SSGSSDSGSD	-,B,C	T	RR
rrnAC2228	-----MSSGDSID RR SFLTA-----AGSAAAAT LAGC SGDGGDGGGT	-,B,C	T	RR
rrnAC2349	-----MTME RR TVLKR-----IGGVGAATA LAGC SVQEQQNGGS	A,B,-	T	RR
rrnAC2589	-----MTLPT RR GFVVG-----CAAT LAGC GVLPEEREPI	A,B,-	T	RR
rrnAC2853	-----MSTIPTAT RRR VLEA-----LGVGTAA LAGC ASAPGAKEQA	-,B,C	T	RR
rrnAC2868	-----MAMDQSAS RR SFLTG-----VGAAATAG LAGC TGLGFQDQTL	-,B,C	T	RR
rrnAC3111	-----MK RR KFLQT-----MGVGATAV GSGC LGGGEVVVS	A,B,-	T	RR
rrnAC3132	-----MPTNGSSVN RR QLLKS-----TGVAGVAGLTG LAGC SGGDGGDG	-,B,C	T	RR
rrnB0207	-----MGSVAMHDTV S RR TVMKS-----IGAAGAVG LAGC STDGGSSGNT	-,B,C	T	RR
rrnB0311	-----MVQSGGGYDGVS RR SVLKA-----SGTALTVGTVG LAGC SSSSGASVRP	-,B,C	T	RR
OE1391R	-----MQ RRR EFLQA-----TGAALAAVG LAGC SDSGGTTDGD	A,B,C	T	RR
OE1806R	-----MPTEHT RRR FLQA-----TGATSIAA LAGC AGGDDGAGDG	A,B,C	T	RR

OE2157F	-----MH RR AFLAG-----GTTLSVGV LAGC IGPSVSKSDY	A,B,C	T	RR
OE2171F	-----MSRDTRLD RR TMLKS----TGAAVTATL LAGC SGGGDTGDDE	A,B,C	T	RR
OE2317R	-----MADFD RR EFLKL----AGGTVGASL VAGC SSSGGGGDTT	A,B,C	T	RR
OE3320F	-----MSAMGRAPD RR TFLRS----AVAGGLAA IAGC TDRTTTGTTN	A,B,C	T	RR
OE3419F	-----MAGEAT RRR CLGHV----AAGAAL LAGC TSHSDTSTST	A,B,C	T	RR
OE3629R	-----MG RR SFLAA----TGAAASATT LAGC LGGTTGTPTI	A,B,C	T	RR
OE3641F	-----MTS RRR FVAA---VGSATAASLG LAGC VGDRETTTAT	A,B,C	T	RR
OE3859F	-----MH RR SFIGG----VSAAGLGI LGGC SAAVGTVAPP	A,B,C	T	RR
OE4008R	-----MRGQPVH RR SVLAL----VGGGAVSA LAGC TDTTDGDATT	A,B,C	T	RR
OE4030F	-----MH RR ALLGG----VAAAAAAS LTGC MGAQSDDQYE	A,B,C	T	RR
OE4197F	-----MPH RRR VLTA----TAAAASLS LAGC LSNDGGGDSQ	A,B,C	T	RR
OE4225F	-----MRT RR QFLAT----TTSLTTVGL LAGC ARSPDNTDER	A,B,C	T	RR
OE4576F	-----MQ RR AFLKA---GSAATLAGL LAGC SSPDSDATST	A,B,C	T	RR
OE4593R	-----MA RRR QILAG----GASLIAAS LAGC TSSQSPTDDG	A,B,C	T	RR
OE5273R	-----MRTT RR SFIGA----STTALLSG FAGC TSGGDDGANS	A,B,C	T	RR
OE1254R	-----M RRR TYLSL----VGSAAAAG TAGC LGVLGGDDTP	A,B,-	T	RR
OE1361F	-----MPS RR DVRLI----GAGVLAAG TAGC TDTAPNRVAA	A,B,-	T	RR
OE1597F	-----MK RR AFLAT----AAVLT TTGC LGRTDSVSDS	A,B,-	T	RR
OE1679R	-----MPADDAERTTRT RR QVLAG----MGATGAAA LAGC QSTSSEDPAD	-,B,C	T	RR
OE1743R	-----MD RR TLLGR----AVAG LTAC VSLSVGAGCL	A,B,-	T	RR
OE2175F	-----MNSDQRGVP RR EFLKA---AVAIGGASA LSAC LGRTDDPIPG	A,B,-	T	RR
OE2330R	-----MN RR AFLTA---SAGLGSTAA LAGC LGALGFERQS	A,B,-	T	RR
OE2514F	-----MMACT RR KALAA----VGTTLS LSGC ARIARPSSER	A,B,-	T	RR
OE3612R	-----MHSTT RR EWLGA---IGATAATG LAGC AGVGGAGQPV	A,-,C	T	RR
OE3910R	-----MD RR NFLKT---AGAAGTIGISG LSGC LGVLGGGED	A,B,-	T	RR
OE3933F	-----MSRART RRR LLSS----VALAVVAG LTGC STHRQSDTAP	A,B,-	T	RR
OE4093F	-----M RRR TVLVG----VTGALFGV GGGC LTMGADETAR	A,B,-	T	RR
OE4305R	-----MLLSPMTMTGDGMSNTS RRR FLKA----TGAAALTAT VAGC SDSTDADGS	-,B,C	T	RR

OE4357F	-----M RRR DYLR-----VGGGATGVA AAGC LQMGGNSAET	A,B,-	T	RR
OE4485R	-----MHSDPDDGASGPVS RR AFVAA-----TGTAGVAA LAGC ANSTDGSGGD	-,B,C	T	RR
OE4551F	-----MSDDTVS RR GFLKA---AGAATVVATS TAGC TDSGGGGDGG	-,B,C	T	RR
OE4700F	-----MH RR PFLSR-----LCLGAVAA TAGC LSRRDDRHS	A,B,-	T	RR
OE5147R	-----MDEQTHTHLS RRR TLTA-----SAGVLSAG LAGC ITSGDTDGDP	-,B,C	T	RR
OE5188F	-----MVDSICT RRR LLAA-----VGATSVSA VAGC SSTLFGGDVE	-,B,C	T	RR
OE5404F	-----MTRNNNPHIS RR TALRT-----VAGTALAS MAGC MDDNGSAGPN	A,B,-	T	RR
HVO_0022	-----M RRR SFLRA-----AGAGGV SAL LAGC TGTGGEQTTT	A,B,C	T	RR
HVO_0062	-----MPDTNKLS RRR FLKA-----TGGAATAAA LAGC TGGDGEETT	A,B,C	T	RR
HVO_0447	-----MAS RR EFIRT---AGITGVAGLTA LSGC TGGDDSSSEAT	A,B,C	T	RR
HVO_0518	-----MN RR ALLLG-----TAGLCAS LAGC ASGGDAAEGS	A,B,C	T	RR
HVO_0530	-----MSRDSQSNDNRLS RR QYVAG-----AGALATMG LAGC SGGGGSNDGG	A,B,C	T	RR
HVO_0553	-----MTVRYD RR TFLRT-----AAGAVGLGS LSGC VGTFGTESGE	A,B,C	T	RR
HVO_0899	-----MARDSKPLN RR DVLKA--TGAVGTAGLAG LAGC TGGGDGGGGD	A,B,C	T	RR
HVO_1007	-----MNNGS RR DFLAA-----TGATLLGG LAGC ASAPTDDGAG	A,B,C	T	RR
HVO_1119	-----MTQRVS RR SFLHA-----TAASTGIAL LAGC SSSSGDGSSGG	A,B,C	T	RR
HVO_1228	-----MEYS RRR LLQT-----TGLAVAAAG LAGC NGRSSDETTT	A,B,C	T	RR
HVO_1401	-----MD RR SFVKA-----AGVAGIAG LAGC TGGPSEGSED	A,B,C	T	RR
HVO_1464	-----MEEKSTGQEGLT RR NYIRY-----GGTALGAGM LAGC ASDSSGSDEA	A,B,C	T	RR
HVO_1624	-----MH RR TLLAA-----AGTGLAAA LTGC IASAGDGADD	A,B,C	T	RR
HVO_1705	-----MHERDREPRGHS RRR FIAT-----TAALGVGA LAGC TGGSDEETET	A,B,C	T	RR
HVO_2038	-----MVNKYKRLT RR DYVKY-----GAAAATAA LAGC GGQSASDPPT	A,B,C	T	RR
HVO_2141	-----MLSTT RRR TLQW-----LGLGGVAS LAGC ATKSPPTAAQS	A,B,C	T	RR
HVO_2149	-----M RRR AFLAS-----ALTAATTL TAGC SSLGPQTEHT	A,B,C	T	RR
HVO_2286	-----MK RR KLLL-----TTGLLTA LAGC NTITGNNKDT	A,B,C	T	RR
HVO_2375	-----MTRNSDSGLS RR KFLIA-----SGAAGLAG LAGC TENNTDGGS	A,B,C	T	RR
HVO_2432	-----ME RR TFLKG-----SAGAAAFT LAGC LGGGEGESES	A,B,C	T	RR
HVO_2607	-----MTNPNDARLS RR AFLGT-----LGTGAAVG LAGC VGGDEGSTGD	A,B,C	T	RR

HVO_2651	MAGRPS-RR-QLL RR GAVAV-----GVGLTGA LAGC SSLRGCSSAE	A,B,C	T	RR
HVO_2695	MVDADSRKGKRNGVS RR SFVKA----AGASGVAVG LAGC ISTGGGDGDN	A,B,C	T	RR
HVO_A0181	MSLT RR TVLGS-----TALLSFGL LAGC VGNNGSDDGA	A,B,C	T	RR
HVO_A0190	MK RR TYLRT----LGAVGVGAA LAGC NTPDEETPTE	A,B,C	T	RR
HVO_A0192	MNLS RR SYLGL-----GAAV LSGC VARGGEPAGT	A,B,C	T	RR
HVO_A0299	MN RR NFVKF----ASGAAAGAA LAGC AGNGGDGGSG	A,B,C	T	RR
HVO_A0362	MPSIS RR EYLAV----SATTVLAA TAGC QTSSCTPTDP	A,B,C	T	RR
HVO_A0380	MVKRDKGGRID RR SFLKY----SGLAGTGIT VAGC TGNQSEETTS	A,B,C	T	RR
HVO_A0428	MSNKSMPDTD RR KLLKT--IAAGATVGPATL LSGC TGGDAKTDVG	A,B,C	T	RR
HVO_A0477	MMAQDSKKLAGSVS RR QFLVA----SSAAGVAG LAGC SGQSNPEADA	A,B,C	T	RR
HVO_A0576	MGKTLKSSTSN RRR FLKA---TGASVAALT LAGC ASDDAGNGGG	A,B,C	T	RR
HVO_A0611	MRTKT RRR FLAT----VAGSAAVG LAGC LGGDDSSDSA	A,B,C	T	RR
HVO_A0620	MSLPT RR QVLRA----GGASLVAA LAGC SGEGSSSSSD	A,B,C	T	RR
HVO_B0093	MAGRSTNWS RR DLIKY----GAVAGTAG LAGC FDSSGQTTED	A,B,C	T	RR
HVO_B0132	MHT RR AVLAA----GGATALAS LSGC VGRALGALDL	A,B,C	T	RR
HVO_B0139	MLS RR EVLAA----GGAALVSG VAGC GGSSPAPDAT	A,B,C	T	RR
HVO_B0187	MVST RR SFIGG----SAAAAG LAGC LTTSSSEGDA	A,B,C	T	RR
HVO_B0198	MV RRR QILAG----SAGLFASA LAGC TAGGSADSES	A,B,C	T	RR
HVO_B0228	MVREAITSRT RR KFIKS----AGVAGTV A LAGC AGNSDNGDS	A,B,C	T	RR
HVO_B0358	MK RR TFLQL----TGLGAAGG LAGC LGSNSPPR	A,B,C	T	RR
HVO_0066	MPRLRS RRR FLQL----LGSASF VAGC LGRDGSSGP	A,B,-	T	RR
HVO_0154	MQ RR QVLAS----LGSLAL LSGC LGGPAESPTE	A,B,-	T	RR
HVO_0564	MN RR TILKQ----LAGTTAVGA LAGC VGVSETDSTQ	A,B,-	T	RR
HVO_0628	MSQDGNDVS RRR FLTA----AGAAAATAG LAGC SGGGGEDTTT	-,B,C	T	RR
HVO_1144	MRDHGSSNGGRT RR QFLAA----TGAAATAG LAGC ATLTASGDDF	A,-,C	T	RR
HVO_1240	MTS RRR FLAA----SGAVAAAG LSGC LAELGRLYTS	A,B,-	T	RR
HVO_1241	MK RR TYLRA----AGAGAAVG LAGC LGGGDANPNV	A,B,-	T	RR
HVO_1244	M RRR ALLSL----LAGAGVAG VAGC LGDGSAPAEA	A,B,-	T	RR

HVO_1245	-----MRNT RR AYLAA-----TAGALTIG TAGC LGGGSGGSGN	A,B,-	T	RR
HVO_1548	-----MH RR ALLSS-----AAALSLAG LSGC LADARRTTSG	A,B,-	T	RR
HVO_1597	-----ME RRR LLGL-----VAVGL SSGC LGSLPGATGP	A,B,-	T	RR
HVO_1612	-----MASTTN RRR FLAL-----GVGAA IAGC LGSEPPVADD	A,B,-	T	RR
HVO_1888	-----MAIE RRR FLQA-----AGVGAVLG LSGC TGNTSPPQAN	A,B,-	T	RR
HVO_1981	-----ME RR TFLKS-----VATTGALLT TAGC MGGGSESQPT	A,B,-	T	RR
HVO_2126	-----MD RR QFIAA-----ASAVGATA FAGC TGGDAGGSGG	A,B,-	T	RR
HVO_2153	-----MRNYVGAPGSTVS RR EFLAA-----TGSGVGALG LAGC AAPTNDGNA	A,B,-	T	RR
HVO_2316	-----MN RR EMLAL-----SGALAAAT IAGC TGDGQSPAET	A,B,-	T	RR
HVO_2397	-----MNTNADSSDPKLS RR SVVAA-----GSGLLTAG LAGC LGSGGGAGSG	-,B,C	T	RR
HVO_2799	-----MK RR ALLST-----VAGSALGL AAGC LSDLDEPVET	A,B,-	T	RR
HVO_A0109	-----MRA RR NVLLG-----ATATLFG FSGC IAPFTSDGSL	A,B,-	T	RR
HVO_A0242	-----MN RR KLLLS-----LGATSLA FPGC LGKQTNLNGTD	-,B,C	T	RR
HVO_A0324	-----MK RR TFVTA-----STLSVAG LSGC LGDTEYRIAD	A,B,-	T	RR
HVO_A0541	-----MADNDDCRSGPT RR DYVKF-----GGTVLGGGL LAGC AGGGSDATTT	-,B,C	T	RR
HVO_A0549	-----MCHDCDRRTVA RR TVLSA-----VGVAGVAA TAGC LGGLGTDGGG	A,B,-	T	RR
HVO_A0557	-----MDQDLTDRATPT RR ACLKG-----GGALLLAGGL LAGC AGQNDAESST	-,B,C	T	RR
HVO_A0558	-----MTDSDGPT RR KFLTY-----GGSVAAGGL LAGC TGSSESDATA	-,B,C	T	RR
HVO_A0623	-----MRSRDARDAGDSDRT RRR FIWL-----AGTAAMAG LAGC SGGGGAEPSE	A,-,C	T	RR
HVO_A0627	-----MK RR DALKT-----LGIGALAA TAGC LGGFEQQSAW	A,B,-	T	RR
HVO_B0082	-----MGSASESVRARFGDAAS RR NFLKA-----LGTAGIAG LAGC SGGESEYDGQ	-,B,C	T	RR
HVO_B0140	-----MPSYS RR DALKT-----IPALAAAG LAGC ASLTGRDDSL	A,B,-	T	RR
HVO_B0144	-----MVDDSENSKTLT RR GCLKY-----GGTIAGSGL LAGC SGTSEPSSSS	-,B,C	T	RR
HVO_B0184	-----MNEARPL RRR TVLSV-----LTAASLGG LAGC LGGADDASPD	A,B,-	T	RR
HVO_B0217	-----MTDQPALTDES RRR YLKA-----IGTAGLTAG LAGC SGGGGGNGGG	-,B,C	T	RR
HVO_B0222	-----MKRTN RRR FLAA-----FSSSAFAF TAGC QRPVASASER	A,B,-	T	RR
HVO_B0318	-----MLDEESSIQ RR DVLSA-----LGAAGVTT LAGC TGGDTGDTDD	-,B,C	T	RR
HVO_C0075	-----MDSDTALD RRR YLGL-----LGGSAVTG IAGC TGGSNQEGL	-,B,C	T	RR

HQ1199A	-----MN RR ELLRV--LACTSISGGTFL TAGC SENTASFNPV	A,B,C	T	RR
HQ1310A	-----MSDESRERDID RR SFVKL----AGTATATAA LAGC GSSGGGGGAE	A,B,C	T	RR
HQ1383A	-----MSS RRR FLQG----IGAVSTVGL VAGC TGSTNNNSNNN	A,B,C	T	RR
HQ1442A	-----MTSDQT RRR FISV----VGAAGITG LAGC SGGGDAGDEG	A,B,C	T	RR
HQ1471A	-----MSKANAGDTQVS RR KFLLT----SGAIGAAG LAGC SSSEGNSSEE	A,B,C	T	RR
HQ1486A	-----MFNKFDKSDLEGPKID RR TAIKL----FAAGGVSS LAGC AGESGDPGEG	A,B,C	T	RR
HQ1619A	-----MVRDNDSLH RR DVLKA---AGASTVGIAG LAGC AGGGGGGDSG	A,B,C	T	RR
HQ1806A	-----MSMQST RR KYIAM----IGAVGGGA MTGC LGGNTNAGGT	A,B,C	T	RR
HQ1816A	-----MK RR SYLRQSVLPLIASAGIGAV FAGC TRTESDDSRI	A,B,C	T	RR
HQ1974A	-----MEIS RRR ALES----ASVTAAVA LTGC IGESTSVSDN	A,B,C	T	RR
HQ1992A	-----MSRNVNS RR SFIKA----ASIAGIAG LAGC SGQSTSNSGG	A,B,C	T	RR
HQ2058A	-----MQ RR TLLST----LSTVGAVT IVGC IGGGDEDPQF	A,B,C	T	RR
HQ2104A	-----MN RR ELLQL----TGVSLGFI ATGC VGIDPPQTD	A,B,C	T	RR
HQ2186A	-----MFDNSGRIN RR NYIRS--VGAASVAGLAS LAGC SGDSSTDDTS	A,B,C	T	RR
HQ2192A	-----MVDEKRTS RR KFLAA----SGSLSAAA LAGC SGGGGGGGGG	A,B,C	T	RR
HQ2247A	-----MNPT RRR LLSSG----LGTAAAVG LAGC SGGGDGSSTS	A,B,C	T	RR
HQ2251A	-----MKQIDKPHN RR TLLSM----IIIAVSVG LAGC TGPGESEDSG	A,B,C	T	RR
HQ2450A	-----MD RR TFVKA----TGIAGIAG LAGC SGGPSGGAGN	A,B,C	T	RR
HQ2477A	-----MSSNID RR TFLTT----AGSAAAAAT LAGC SGSDDGSAASD	A,B,C	T	RR
HQ2598A	-----MLTG RR GFIAA----GASAVLAG LAGC SGTLGGSASG	A,B,C	T	RR
HQ2732A	-----MD RR SYIKS----GAGVIAGGL IAGC TGSDSESEST	A,B,C	T	RR
HQ2754A	-----MYSMIQ RR KFVAG----TGAATVAA LAGC SGSSDTGDES	A,B,C	T	RR
HQ2809A	-----MSERS RR DILKG--LGAAGTAGVVS LAGC SRQSGGGSTE	A,B,C	T	RR
HQ2849A	-----MNNWS RR EFVRI----ATVGAGAG IAGC NALSPSGSIK	A,B,C	T	RR
HQ2969A	-----MARDIE RR AFLKR--AGAAGVLGTVG LSGC VGGGGEGGEG	A,B,C	T	RR
HQ3009A	-----MTLT RR QALLA----TGTTLTAL LSGC GATQQEFTFS	A,B,C	T	RR
HQ3060A	-----MSWT RR GVLAS-----MAGVCG VSGC LASPDRSLPT	A,B,C	T	RR
HQ3135A	-----MPT RR TVIAA----LTTVGTAT LSGC GSFDNTADNN	A,B,C	T	RR

HQ3182A	-----MTENTN RR TFLKI---AGSTGAVGLAG LAGC AGGGGGGDGT	A,B,C	T	RR
HQ3283A	-----MSDGET RR EFLEA----FGVASISI LGGC QTSGDSDTQT	A,B,C	T	RR
HQ3383A	-----MIDKS RR GFLAG----VGAAATTS LAGC AGIGSGSQTL	A,B,C	T	RR
HQ3619A	-----MSDNRMS RR AVLAT----GAVGLGAS LAGC SSSGGSGGD	A,B,C	T	RR
HQ4034A	-----MDMSMN RR GVIEA----IGVGGLTV LAGC SSNSPNEQSE	A,B,C	T	RR
HQ4036A	-----MD RRR YIGI----VGTIVAGA LAGC NSNTENNAPS	A,B,C	T	RR
HQ1481A	-----MVDDSEVN RRR FLRA----TGAVAATVG VAGC SQGSDGETET	-,B,C	T	RR
HQ1855A	-----MTDSKIG RRR LLGL----TGAAVLGG LAGC NTQTQSQSNT	A,B,-	T	RR
HQ1912A	-----MLDDSLT RR ALLAS----GVAVTLG LTGC AETDPSSTPV	-,B,C	T	RR
HQ2373A	-----MD RR NYLHA----IAGAGAFA TAGC LGSTLSFRDG	A,B,-	T	RR
HQ3303A	-----MSSEGADKKQRIAN RR TYLKG----LGAAGTVL LAGC SGSGGGSGGD	-,B,C	T	RR
Nmag_0005	-----MKRVAPSVT RR SLLAG----ASGVGLSA LAGC SERFWSRAEN	A,B,C	T	RR
Nmag_0106	-----MVH RR NILKG----AAGVGLAS LAGC LGGDNGGVDT	A,B,C	T	RR
Nmag_0499	-----MAGRDLDSA RRR FLGA----AGAAATAG IAGC VSLPTALDDR	A,B,C	T	RR
Nmag_0622	-----MQTDPT RR TFLAG----SATAGIVA VAGC ASDGDGNNGE	A,B,C	T	RR
Nmag_0655	-----MN RR TALRS----AGVLATVG LAGC VDAVEEHFQG	A,B,C	T	RR
Nmag_0697	-----MNGLCD RR AFLAS----AATGGMLG LAGC LSGDAEQVST	A,B,C	T	RR
Nmag_0704	-----MDDRLT RRR LIAI----SGAVSAAA LAGC NDDPEDPDDP	A,B,C	T	RR
Nmag_0795	-----MEGNSRYK RRR VIGL----IGTGTVVVS VAGC LTDDTEDDGV	A,B,C	T	RR
Nmag_0945	-----MDPSTD RRR FVQA----MGGGFAVA LAGC LSDDEENGGE	A,B,C	T	RR
Nmag_1151	-----MPLQ RR QLLAG----IGGAATLT VAGC LGDDDRGTTL	A,B,C	T	RR
Nmag_1294	-----MN RR KLLLG----SGTFVASM AAGC LQAGGTSENK	A,B,C	T	RR
Nmag_1357	-----MHRPSFPS-RR-S RR AFLSG----TAVAGTAA LAGC TTLYHVLPR	A,B,C	T	RR
Nmag_1387	-----MTFRPK RR ELLMG----AGAVSLGG LAGC LSDVPGLESE	A,B,C	T	RR
Nmag_1419	-----ME RR KILLG----SGAALATA LAGC SSSETDEPSG	A,B,C	T	RR
Nmag_1438	-----MGDDSKS RR TVLKG--VGIASAAGLTTs LAGC VSQNGGS DVE	A,B,C	T	RR
Nmag_1742	-----MSDLP RRR LLAL----GGTALTGA IAGC SSSANEDDPN	A,B,C	T	RR
Nmag_1771	-----MQ RR TYLTL----TGSVGLLA VAGC LSEEQSTPKE	A,B,C	T	RR

Nmag_1801	-----MN RR QHLAQ-----LGGVGLVA LSGC LDTISSLTD	A,B,C	T	RR
Nmag_1824	-----MD RR SLIHT-----AVGASFI LSGC LSRSDGNSET	A,B,C	T	RR
Nmag_1899	-----MK RR TYLTT----TSATAVGLA LAGC MGGEESDEDP	A,B,C	T	RR
Nmag_1923	-----MSRVG RR TYLAA---AGGAVAGLAG LAGC LGDGGDSTPI	A,B,C	T	RR
Nmag_1931	-----MQ RR TVLAG----LGSLSGLLA LSGC LSDDHGEIDE	A,B,C	T	RR
Nmag_2002	-----MELT RR KALTT----SFSGLVMGT IAGC TDDTPEDEEE	A,B,C	T	RR
Nmag_2005	-----MDVT RR SVLKR----TLGGAALGA LAGC LNEPDETS GT	A,B,C	T	RR
Nmag_2069	-----MPS RR SVLGA----GATAGFAT VAGC LGSNHSFGAL	A,B,C	T	RR
Nmag_2138	-----ME RRR VLAT----AGGLLPIS VAGC VTVSTESADE	A,B,C	T	RR
Nmag_2173	-----MARHTT RR SVLKG---STAAA AVGTTL LAGC VTDEGN GTGD	A,B,C	T	RR
Nmag_2271	-----MIP RR KFLQC----GVVSFSGSI IAGC SESTETSNPE	A,B,C	T	RR
Nmag_2366	-----MTKRQSRSGGWSS RR TFVLA----TGAIGTSV LAGC LDGDDSDSQA	A,B,C	T	RR
Nmag_2372	-----MAKFTRPV S RR GVLTS----GATAVGVA VAGC ISDDTDGAGQ	A,B,C	T	RR
Nmag_2397	-----MVWNIG RR TLLSV----TGLAL LSGC LDSGNTGETN	A,B,C	T	RR
Nmag_2411	-----MNQT RR AFLGT----TGTVGLGV VAGC LGSEDPPPEPP	A,B,C	T	RR
Nmag_2669	-----MASDRDQT RR QLLGL---AGATVSTML IAGC GDTEPEDENG	A,B,C	T	RR
Nmag_2689	-----M RRR PFLVA----ASATTVG LAGC TSAQERLGDV	A,B,C	T	RR
Nmag_2702	-----MNDSRLD RR QFLAA----AGAGLAGV VAGC SEPNNASSIE	A,B,C	T	RR
Nmag_3059	-----ME RR TYLGS--LGAAGAAGLTG LAGC LDELSGDDQ	A,B,C	T	RR
Nmag_3090	-----MVWELN RRR VLSG----VGAAGIAG LAGC IGDEEEVTEG	A,B,C	T	RR
Nmag_3154	-----MEYN RR TILQS----LLPAVGL LAGC SFRDDVHLFE	A,B,C	T	RR
Nmag_3278	-----MADNRMT RRR LLAG----SSVAAAAG LAGC TDRLLDLRGD	A,B,C	T	RR
Nmag_3316	-----MPS RR EALLT----AGPLLSST LAGC LDTFRSGSL	A,B,C	T	RR
Nmag_3531	-----M RRR KYLTG----ISSIATAG IAGC AGHTPTDDDQ	A,B,C	T	RR
Nmag_3965	-----MAIQ RR GFIAA----VGTGGAVG LAGC TGLGDDTGDS	A,B,C	T	RR
Nmag_4115	-----MN RR TVLTT----AGLVA VAGC LSESSDPVTE	A,B,C	T	RR
Nmag_4271	-----MN RR SLLAM----AVPASFAA LSGC LGSNSALDDW	A,B,C	T	RR
Nmag_0096	-----MVN RR DIIKG----AGAASIAG LAGC LGGDNGGRDI	A,B,-	T	RR

Nmag_0184	-----MQGANYSLD RR TLLQR--TGAATSIAGLAA IAGC IDDSNGDDDD	-,B,C	T	RR
Nmag_0242	-----MN RR GYLGA----IGAVGLTA TAGC VEELQAAADG	A,B,-	T	RR
Nmag_0327	-----MTDFPT RRR VLQL----TGVGATAS LAGC TQFDIPGIGS	A,B,-	T	RR
Nmag_0438	-----MN RR TYLGA----VAGGLTAT SAGC LGDFGDSDID	A,B,-	T	RR
Nmag_0446	-----MAENNPTT RR TILKV----SGAVGAGAF LAGC ADNGNGNDEP	-,B,C	T	RR
Nmag_0460	-----M RRR TFVGT----LGGGAIAG FAGC LSREDTDDEN	A,B,-	T	RR
Nmag_0557	-----MSPS RRR FLLGA----TGLTGA TAGC LSYVRRVSPG	A,B,-	T	RR
Nmag_0563	-----ME RR ALLTS----LCFPLFVS VSGC LAREQSPRLS	A,B,-	T	RR
Nmag_0609	-----MKYD RR TYLSS----AIITIT LSGC TGLINSGRIR	A,B,-	T	RR
Nmag_0793	-----MN RRR ALTV----GVTTGLTA VAGC LTGMLDDESS	A,B,-	T	RR
Nmag_0856	-----MN RR QYLAS----ATGVAIVG LSGC LQILASGEIS	A,B,-	T	RR
Nmag_0963	-----MVDDTTMSNQYALT RR SALAS----ASGLLVG VAGC LSGDDNEVDS	-,B,C	T	RR
Nmag_1091	-----MPST RR SLLAS----ALVGSA A LAGC LSRVRSPSPS	A,B,-	T	RR
Nmag_1209	-----MPS RR AVLAT----CGSAV LAGC SAVRREPSVT	A,B,-	T	RR
Nmag_1789	-----MPAKNTHG RR TFLRS--TAAAGSVAALGG LAGC TGMLDGDDT	A,B,-	T	RR
Nmag_1814	-----MCPPRS RR PVLAS----VASLSAL TAGC STNWFSNPET	A,B,-	T	RR
Nmag_1940	-----MD RR TFVHG----VGGGSVTA LAGC LTRNGENEHH	A,B,-	T	RR
Nmag_1961	-----MK RR TLLRA--TGGAVLGGSVG LAGC LDQFSDSESD	A,B,-	T	RR
Nmag_2110	-----MSQGHTD RR TMLAL----SGAFLAG MAGC TGTAPDSEDE	-,B,C	T	RR
Nmag_2327	-----MDLN RR TMLGG----IGTASLTG LAGC LNLVGLATHE	A,B,-	T	RR
Nmag_2358	-----MAPSQNLRGRLS RR SAIAT----IGSIGALS LAGC LGGDEDGLSG	-,B,C	T	RR
Nmag_2390	-----M RRRR LLST----GVVVA LAGC VSLPADEDET	A,B,-	T	RR
Nmag_2456	-----MH RR SVLAA----ASAVASVG IAGC LEFFEDNGED	A,B,-	T	RR
Nmag_2492	-----MVENNNRYRPGIN RRR VLQG----LTAAGVIG AAGC LDSEETDETG	A,B,-	T	RR
Nmag_2581	-----MPS RRR TLAL----GATLSTTA LAGC TALRSEPEYL	A,B,-	T	RR
Nmag_2647	-----MD RR TYLST----SAATI FAGC TELSGADEPD	A,B,-	T	RR
Nmag_2680	-----MN-RR-NRSANAERGTT RR TILAG----ATTAAIGL AAGC LSEDEESVPD	A,-,C	T	RR
Nmag_2733	-----MD RR QFLAA----STVGVAA TAGC LGNALSTDD	A,B,-	T	RR

Nmag_2739	-----MPSD RR TFLTI-----CGTGATTA LAGC ADARVSARDV	A,B,-	T	RR
Nmag_2761	-----MLS RR TTLQA-----TIPLFASL LSGC VSSMLPRDPP	A,B,-	T	RR
Nmag_2904	-----MAHQPSGQSKLTS RR AVLSA-----AGAGVTAA LAGC SDSQDEDPVR	-,B,C	T	RR
Nmag_2996	-----MTHPS RR TLLGS-----TAGLLAL TAGC IADNDDDD	A,B,-	T	RR
Nmag_3163	-----MD RR TWLTA-----TTVGTLSA FSGC LSELADRNS	A,B,-	T	RR
Nmag_3234	-----MH RRR VLAH-----VGTGTVIG TVGC LSVPLPTGDS	A,B,-	T	RR
Nmag_3274	-----MTDSN RR SFVAA-----TAAVGTLG VAGC LSTIDGWGDG	A,B,-	T	RR
Nmag_3481	-----MATDPRDTT RR SLLAT-----TGSLGVLG LAGC LSDDEEDGG	A,-,C	T	RR
Nmag_3486	-----MRTSFP RRR LLAT-----TGTISALAI GAGC LDDSDSESDP	A,B,-	T	RR
Nmag_3525	-----MTDRHTT RR TVLQA-----AGATSVVAL AAGC LSDDEEENG	A,-,C	T	RR
Nmag_3601	-----MD RR EFVAA-----GAVCGLGL MAGC LDDALADVTS	A,B,-	T	RR
Nmag_3725	-----MLDSDHVS RR TALRL-----SLASGVAI LAGC ADDNDENDVD	-,B,C	T	RR
Nmag_3851	-----MPIGKQS RRR FLTA-----TGGAAALGT VAGC LGGDDDDVVN	A,-,C	T	RR
Nmag_3944	-----MMSFKDGVS RR DALKV-----AGGVGLVG LAGC LGDGEQTDIT	A,B,-	T	RR
Nmag_4024	-----MASRQID RR SFLQV-----AGGAGIAG LAGC LSGDVEAVSI	A,B,-	T	RR
Nmag_4050	-----MEPDGNMQFN RR ELISA-----LSAGGVLA LAGC ADQADGDGND	-,B,C	T	RR
Nmag_4142	-----M RRR EFIAV-----GCCSVCGI SAGC LDEDDSEPVP	A,B,-	T	RR
NP0126A	-----M-RR-PHS RR QFLVA-----AGLTVTAA AAGC FDSASEPTGE	A,B,C	T	RR
NP0278A	-----M RRR ALLTS-----CVTVAAVS ISGC FDDGESPSGG	A,B,C	T	RR
NP0322A	-----MDLS RR GVLAA-----GAVA LAGC ADLEEQADRR	A,B,C	T	RR
NP0660A	-----MPELT RRR LLAA-----AAGGIAG LAGC AAGSRESPPP	A,B,C	T	RR
NP0894A	-----M RRR TFLST-----VAAGTIAA SAGC SSEELGLGLD	A,B,C	T	RR
NP0938A	-----MDST RRR MLGA-----IAGSLSVV AAGC LGGDDAGDAP	A,B,C	T	RR
NP1600A	-----MHTTPT RR TVVRT-----AAAAAGLA VAGC LDSEPDEAPE	A,B,C	T	RR
NP1648A	-----MERTV RRR TVLTV-----LASSGSVL VAGC SDDDGTETPE	A,B,C	T	RR
NP1652A	-----MTRKLS RR QLVSG-----LAAAGAGV AAGC TDGEEGDSED	A,B,C	T	RR
NP1748A	-----M RRR TYLTA-----AGGLFAAG LAGC TDEEPTGNGN	A,B,C	T	RR
NP1978A	-----MS RR SLLAA-----AGAGLAA LSGC TVLGDPPGED	A,B,C	T	RR

NP2004A	-----MRNPTVS RR KLLAS-----GAAAAGIG LAGC MGGNGGSDGP	A,B,C	T	RR
NP2092A	-----MH RR TFVSG-----VAAGAVAG LAGC LGGDDDAPTT	A,B,C	T	RR
NP2184A	-----MVS RR GVLAA-----LAVGASSS LAGC HTGALVEVQD	A,B,C	T	RR
NP2186A	-----MD-RR-VT RR GVLAA-----AGIA LAGC TDNGDDQPVR	A,B,C	T	RR
NP2288A	-----MQQRILSDS RRRR TVLAA-----IGAGAAG LAGC LDGGSSNDPE	A,B,C	T	RR
NP2536A	-----MSPDLT RR GYLAV-----TAAAGGAA LAGC SDDVTADPDA	A,B,C	T	RR
NP2818A	-----MNRDELVANKTVG RR ALLGS-----VAAFAGV VAGC SQLSPASSKE	A,B,C	T	RR
NP2860A	-----MQPS RR TFVKS-----GIGVVGLGA LAGC AEEDPDGAE	A,B,C	T	RR
NP3028A	-----MTRYT RRR LLGS-----SALAALGA FAGC TAEQPDDPEE	A,B,C	T	RR
NP3032A	-----MKRTT RRR FIAS-----GAAVGAI T IAGC ADEDPEPEPE	A,B,C	T	RR
NP3040A	-----MQ RR EFLAG-----GAGLTAF TAGC ADSDRANVGI	A,B,C	T	RR
NP3142A	-----MSKQTQDTWT RRR MLAV-----AGAGAVVG LAGC ADESSEEPEE	A,B,C	T	RR
NP3232A	-----MD RR AFLAT-----TGAAAAAS VAGC LGRGLSDEDF	A,B,C	T	RR
NP3388A	-----M RRR TYIAL-----SGAALAS LAGC SDDEGENGDG	A,B,C	T	RR
NP3578A	-----MKLT RR QALTA-----FGSAGVVG MAGC LGGGNDDTEL	A,B,C	T	RR
NP3954A	-----MKDIS RRR FVLGT-----GATVAAAT LAGC NGNGNGNGNG	A,B,C	T	RR
NP4140A	-----MTRHIN RR NVLKA-----AGAASMVA LAGC NGNGNGNGNG	A,B,C	T	RR
NP4216A	-----MPS RR ELLRL--GGATLAAASAAG IAGC SEEAFEGPAD	A,B,C	T	RR
NP4616A	-----MNPS RR TVLKS-----GVGVAAAAA LAGC NGNGNGNGEE	A,B,C	T	RR
NP4692A	-----MAPP RRR QILSI-----GGLGLAAA IAGC GDTAPDDTP	A,B,C	T	RR
NP4700A	-----MASDTT RR QFLIA-----ASAAATFTV VAGC SEPEEDPPGY	A,B,C	T	RR
NP4736A	-----MIS RR EFVAS-----VTATGAVV VAGC GTDEDAAPEE	A,B,C	T	RR
NP4744A	-----MGTH T RRR FVRG-----TAAVAATGA LAGC NGNGNGNENG	A,B,C	T	RR
NP4750A	-----MRWT RRR VVAA-----IGAATTAA VAGC SDTDEDAENG	A,B,C	T	RR
NP5018A	-----MKLH RR TALRL-----AGVGVGTA VAGC LDPETDNGET	A,B,C	T	RR
NP5270A	-----M RRR TLLVS-----AAALPA VAGC SNVLPTGTND	A,B,C	T	RR
NP6072A	-----MN RR AALT V-----VTTATASA LAGC GGVDIGTSGA	A,B,C	T	RR
NP6086A	-----M RRR ALLAA-----TGTACGV LAGC LDIDSRFSGS	A,B,C	T	RR

NP6254A	-----ME RR AFLMA-----TGGAAT IAGC TTGGNTEEN	A,B,C	T	RR
NP6272A	-----MN RR AIIVA-----GGVFF LSGC GERATDMSNE	A,B,C	T	RR
NP0250A	-----MELS RR SFLAT-----SAVATAG FAGC TGDSGDPDPP	A,B,-	T	RR
NP0254A	-----MPRFD RR DYLRA-----AGLAGAAA LAGC TDVFDDNGET	A,B,-	T	RR
NP0528A	-----M RRR TVLAL-----AGAAGTAG LAGC LDDVLGSENG	A,B,-	T	RR
NP0544A	-----MVDTSNLD RR SLLKG-----VGVAGIAG LAGC TGNGDGNGNG	-,B,C	T	RR
NP0758A	-----MSDEGVDF RR TFLQY--GGTAAAAASVS VAGC NGNGDGNGNG	-,B,C	T	RR
NP0950A	-----MPT RRR VLAV-----AAAATA TAGC TALGDGENDE	A,B,-	T	RR
NP1070A	-----MRS RR TVLGA-----AAVLASG FAGC FSTEAPDDQE	A,B,-	T	RR
NP1480A	-----M RRR QVLAA-----VPVA LAGC VAAPEPGPPT	A,B,-	T	RR
NP1598A	-----MTQNTNPN RR TFLKA-----AGAGGAFL LAGC VGNEGEPTDE	-,B,C	T	RR
NP1654A	-----MN RR QFLAA-----GALVA AAGC LGPDDAEPPA	A,B,-	T	RR
NP1680A	-----MAT RRR FMCM-----GSTTVGLVA LSGC LGAEGAEVVD	A,B,-	T	RR
NP1858A	-----MVS RR QLLAA-----GGLASAG LAGC LDSLSDGSSP	A,B,-	T	RR
NP2396A	-----METQVS RRR LLQA-----GGVGAAAS FAGC GYMDTEDGGA	A,B,-	T	RR
NP2622A	-----MDPQSSSQSVRMS RR SVLGA-----VGAGATAA LAGC SGGQDGDPPLR	-,B,C	T	RR
NP2694A	-----M RRR ELIAA-----APLVA TAGC TIGGEPPDGS	A,B,-	T	RR
NP2862A	-----MNLS RR GLLTA-----GAGAATAA VAGC VDAAAPGQES	A,-,C	T	RR
NP2890A	-----MK RR DVLAG-----VGATALLG TAGC LGVAGLDEHE	A,B,-	T	RR
NP3030A	-----MKFRERDGLS RR EYVRA-----MVAAGGAAG LAAC LEAFDDGDTG	A,B,-	T	RR
NP3732A	-----MPPE RR DVLRL-----GTVGLGGL VAGC LDNGGADFQS	A,B,-	T	RR
NP3968A	-----MELS RR DALRA-----GGGLLAAAA LSGC VEERVTRRET	A,B,-	T	RR
NP4332A	-----MMPSIS RRR LLVG-----GAAALGSA LAGC PSPTPGGAE	A,B,-	T	RR
NP4786A	-----MN RR SFVAG-----AFGGAGIV TAGC IDIDSEDWEM	A,B,-	T	RR
NP5000A	-----MSQQRQTDGNDGRFTG RR AFIVG-----AAATGVAA FAGC TDNGNGDGNG	A,-,C	T	RR
NP5286A	-----MSSRLVS RRR AVAV-----AATVA VAGC LTEAGRRCRG	A,B,-	T	RR
NP6022A	-----MSLPPTAS RRR MLAA-----AGTALATFS LAGC SDESDDDDPE	-,B,C	T	RR
NP6104A	-----MAHTPSDRVS RR SVLAT-----TGAAAAGF LAGC LGGGGGDGLS	-,B,C	T	RR

rrnAC0160	-MMTERGIS RR EFAKS-----AVAIGGTAA LAAC LDRGSGTVPK	A,B,C	-	RR
rrnAC0236	-MLLE RR HLSLA-----LVVLLIA SAGC SGLFGSETGT	A,B,C	-	RR
rrnAC0863	-MK RR HLLL-----AVLALVA LSGC TGFFGSEEV	A,B,C	-	RR
rrnAC2237	-M RR PFALV-----ILCAVALL LAGC QAPSVDPE	A,B,C	-	RR
pNG7056	-MRCS RR HALRL-----IGASGVITA TAGC LNPGSLDNYA	A,B,-	-	RR
rrnAC0311	-MY RRR VLAL-----CGLAII GAGC QGETADTEV	A,B,-	-	RR
rrnAC1046	-M RR LVLA-----CLVSF LAGC GAFMTDSPPL	A,B,-	-	RR
rrnAC1739	-MGDLS RR GFGAS-----VLAAL SAGC LGSRPDQESE	A,B,-	-	RR
rrnAC2271	-M RRR HFVQT-----LGVSALVG SAGC AEQSDDTTSA	A,B,-	-	RR
rrnAC2940	-M RR GVAIA-----VVCLCLL SAGC TGLFGDNAAA	A,B,-	-	RR
OE3190F	-M RR QLAAL-----LVVALVA VAGC AGGSAPATDD	A,B,C	-	RR
OE4563F	-MA RR LLAVA-----VVCLVV LAGC QGGMSGDATT	A,B,C	-	RR
OE6028R	-MSL RR LLAVF-----AVVGMVV IAGC AGGIDNGEPA	A,B,C	-	RR
OE5030R	-MP RR VLLRS-----IGVCSAVG IAGC LSQGEARSIT	A,B,-	-	RR
HVO_1806	-MN RR LAAVF-----SAFLV VAGC AAPVADPSAA	A,B,C	-	RR
HVO_1926	-MQ RR TIAPI-----FLVGLVL LAGC LSAPLQTTAD	A,B,C	-	RR
HVO_2145	-MTRDTNHT RRR FTAL-----TATAALSTA LAGC SGSGGDAANA	A,B,C	-	RR
HVO_1176	-MN RR LLVGI-----AALALLV TAGC LGGTSSVSND	A,B,-	-	RR
HVO_1477	-MV RR ATATL-----VVVVLAV LAGC LGGGGAGAAT	A,B,-	-	RR
HVO_1673	-MSS RR VVAVL-----ALAALVL TSGC IGFLTGEETL	A,B,-	-	RR
HVO_1802	-MTV RRR SLGM--RGSTLLVAVLVV LAGC AAPVSPGTDG	A,B,-	-	RR
HQ1059A	-MS RR ISRAS-----IALSLLIV LAGC SGGSLSFNPN	A,B,C	-	RR
HQ2489A	-MN RR LLLAV-----CALAFLGV LSGC LGIGTGPIPA	A,B,C	-	RR
HQ3375A	-MNVSQ RR IVMLG-----IALLVV FSGC AEVNDSIAPT	A,B,C	-	RR
HQ1196A	-MI RR QILTK-----CAVALSVG LVGC SGGGGGGSTD	A,B,-	-	RR
HQ2519A	-MI RRR FISA-----IVAGIII IGGC TGPNDSPNTA	A,B,-	-	RR
Nmag_0702	-MNAKEIMN RR HVLSS-----IGTTTGTL LAGC LTDDSGEPMA	A,B,C	-	RR
Nmag_1399	-MY RR LLLSS-----ISLVA VSGC TSNSDKEDGT	A,B,C	-	RR

Nmag_2088	-----MY RRR LLAA-----AGLTTAV VAGC LTSDGTDGDG	A,B,C	-	RR
Nmag_2424	-----MN RR VYLAA-----VGTSLSAS LAGC SSVMSVFDDE	A,B,C	-	RR
Nmag_2835	-----MYVT RR LVCVA-----ILL TAGC VGSDDEGLDD	A,B,C	-	RR
Nmag_3079	-----MK RR HILTT-----IGASLSFA TAGC LQGGSSNNEG	A,B,C	-	RR
Nmag_0572	-----MK RR LVLAT-----AGATL VSGC SSYLRAPSNS	A,B,-	-	RR
Nmag_1805	-----MVS RR HLLSS-----LGCAA LAGC TAVLPYDGTQ	A,B,-	-	RR
Nmag_2536	-----MSVP RR MMGAA-----LLGV VAGC LSIGPGAENN	A,B,-	-	RR
Nmag_2731	-----MID RR TSLVA-----VLGSILA ISGC LERISNGDTE	A,B,-	-	RR
Nmag_3155	-----MN RR LVFGS-----LAVVLLIG LAGC TMFFTGISDD	A,B,-	-	RR
Nmag_3710	-----MN RR VCLRS-----IAASSVVT TAGC LESLSQENDE	A,B,-	-	RR
Nmag_3769	-----MH RR VYLST-----LSAAGIGS LSGC LDILATDSES	A,B,-	-	RR
NP3728A	-----M RRR VLLSL-----VSGTAAA LAGC ADGMEEANDD	A,B,C	-	RR
NP4392A	-----MK RR YYLGT-----IAAAA LAGC SELQPEEAPE	A,B,C	-	RR
NP6066A	-----MPS RR YILAC-----VGAASIA GAGC TDLSDSNGRP	A,B,C	-	RR
NP1172A	-----MD RR LLAAT-----ALAAVLL SAGC TVGYQPAADA	A,B,-	-	RR
NP4102A	-----MN RR LLAAV-----AVFVLFG LAGC TTIFGADVDD	A,B,-	-	RR
NP4728A	-----MSSHPRQNESKGVS RR TFALL-----AGSAVTAG LAGC TDDPMENGDA	A,B,-	-	RR
NP6234A	-----MQ RR KFSTT-----TGSALSIL FAGC LGGTDVPDNE	A,B,-	-	RR
rrnAC0370	-----MDSGRNGGGPSVF RR DLLRS-----IGAAGLAG LAGC GGSEGVTDGD	A,B,C	-	RR
OE3575R	-----MRNDDCCS RR QFGTT-----LGGIGVATM IAGC SAFGLPSKKP	A,B,-	-	RR
HVO_B0280	-----MRETLT RR QFGVG-----VAVATVA LAGC TDTGGGDGPV	A,B,C	-	RR
NP4850A	-----MDE-RR-CT RR QVGAG-----VIGGLSVL VAGC LGNGDGNSSH	A,B,C	-	RR
rrnAC1401	-----M HR IVPAI-----GLAALVV LSGC VTATVDSTVA	A,B,C	-	-
rrnAC2054	-----MT TR LLTVL-----VVSLLV LAGC LGGGPAVGDG	A,B,C	-	-
rrnAC2239	-----M RK ELLIA-----LTVV LAGC TVPSFGGSDH	A,B,C	-	-
rrnAC2345	-----MTA VR PLVLV-----AIAVLVS LAGC GALFGSGGEA	A,B,C	-	-
rrnB0030	-----MYRFARNLCS LR GLARV-----VVVVMLVT ASGC SAIGSLSSEN	A,B,C	-	-
rrnAC0024	-----M RW LSTAA-----VVVVLL TAGC NAFVGADTA	A,B,-	-	-

rrnAC0808	MASR IR VLAVV-----GLLL LAGC TGMGGSGDAG	A,B,-	-	-
rrnAC1293	M RK GILTL-----LGVVLVV SSGC TGLITGETVA	A,B,-	-	-
rrnAC2079	MT KR KTIFS----SVAIALLII SGGC SGFVGGEPEV	A,B,-	-	-
rrnAC2645	MVRS PR VESLA----VAALLV LAGC GGFVANGVDS	A,B,-	-	-
rrnAC2720	M WR VTLAV-----GIAL IAGC SSFFPTSPTA	A,B,-	-	-
rrnAC3140	M RW GVFGV-----VVVLVV ATGC SGFAGSPSQS	A,B,-	-	-
rrnAC3358	MA GR IAVLL-----VLLL FAGC GAFFGAERQT	A,B,-	-	-
rrnAC3416	M TR AVVAI-----VTLALV LAGC SGFVQQDSPE	A,B,-	-	-
OE3264F	MR PR IVLIA----ALVVSSIA LAGC SGTQSGTPA	A,B,C	-	-
OE2121F	M RA LWVAL----AVAALAV TAGC AGFGGPQSAA	A,B,-	-	-
OE2791R	MVRDLHV PR TALLA----VGLAVLLV AAGC TGMPGGSSAD	A,B,-	-	-
OE3655R	M VR SATIP----LVLIALLA AAGC LGSPIMEPVR	A,B,-	-	-
HVO_0494	MYS NR IPVLL----IAVLVL LAGC AGGADVAPAA	A,B,C	-	-
HVO_0807	M RV SVLLA----AALVL STGC LGGGSGTARL	A,B,C	-	-
HVO_1030	M KR TGLAT----VLLALMVV LAGC AGGAGTVTET	A,B,C	-	-
HVO_1580	MSPAA DR GIAPL----ALALVLLV LSGC TSSSPSPVV	A,B,C	-	-
HVO_1807	MAR FR ALPLA----ALAALAV LSGC VGVPTADAPG	A,B,C	-	-
HVO_2535	MQPTA AR VLLAA----LMVL LAGC GGGVGGDATT	A,B,C	-	-
HVO_1609	MD NR LKLLS----VALLVV LAGC TGGAADGGAG	A,B,-	-	-
HVO_1808	MRTTG LR AVFAA----VLLV LAGC AAPTAAPPGD	A,B,-	-	-
HQ1800A	M AR TPIIV----AGVVFLM IAGC SAPTAPGPDD	A,B,C	-	-
Nmag_2608	M RQ LAALL----LVALLVM SAGC SAFDSSPDGD	A,B,C	-	-
Nmag_4102	M KR LLLVV----LVAAMLV LAGC ADGAGTDTDE	A,B,C	-	-
NP5284A	M RI NVAAF----AVLAVVV TAGC AGALLGEDLD	A,B,-	-	-
OE3125R	M PK YSTGG----AGSGGG GQAC ELCGSTADSL	A,B,-	-	-
HQ1883A	MT TK EESTN-KLIEQALTATTIA VIGC STSPGKPAHD	A,-,C	-	-
Nmag_2200	MTKRQGSNSQQVGRGLR TR VRSLP---LGLVAVLFLVV LAGC TLPASPDQFG	-,B,C	-	-
NP4668A	M KL NVVGS-----GYVGTT LAAC LAEMGHNVSA	A,B,-	-	-