

	10	20	30	40	50	60	70	80	90	100
LOC_Os01g70270.1 11667.m07038_altsplice/1-809										
LOC_Os01g13520.1 11667.m01333/1-700										
LOC_Os02g06910.1 11668.m00642/1-909										
LOC_Os02g35140.1 11668.m03331/1-757										
LOC_Os02g41800.1 11668.m03990/1-784										
LOC_Os04g57610.1 11670.m05695/1-819										
LOC_Os04g43910.1 11670.m04249/1-696										
LOC_Os04g49230.1 11670.m04840/1-294										
LOC_Os04g36060.1 11670.m03490/1-1674										
LOC_Os05g43920.1 11682.m04181/1-700										
LOC_Os06g09660.1 11680.m00929/1-1056										
LOC_Os06g46410.1 11680.m04579/1-918										
LOC_Os06g47150.1 11680.m04656/1-731										
LOC_Os06g48950.1 11680.m04850/1-1116										
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056										
LOC_Os11g32110.1 11687.m02952/1-812										
LOC_Os12g41950.1 11686.m04180/1-900										
LOC_Os12g29520.1 11686.m02861_altsplice/1-842										
LOC_Os04g56850.1 11670.m05610_altsplice/1-956										
LOC_Os05g48870.1 11682.m04683_altsplice/1-696										
LOC_Os01g48060.1 11667.m04657_altsplice/1-723										
LOC_Os01g54990.1 11667.m05421/1-656										
LOC_Os10g33940.1 11676.m02996/1-699										
LOC_Os02g04810.1 11668.m00419/1-1094										
LOC_Os04g59430.1 11670.m05892/1-530										
LOC_Os07g08520.1 11673.m00789/1-729										
LOC_Os07g08530.1 11673.m00790/1-408										
LOC_Os07g08600.1 11673.m00797/1-525										
ARF1/1-665										
ARF3/1-608										
ARF4/1-788										
ARF5/1-902										
ARF6/1-933										
ARF7/1-1165										
ARF8/1-811										
ARF9/1-638										
ARF10/1-693										
ARF11/1-601										
ARF12/1-593										
ARF13/1-623										
ARF14/1-605										
ARF15/1-598										
ARF16/1-670										
ARF17/1-585										
ARF18/1-602										
ARF19/1-1086										
ARF20/1-615										
ARF21/1-606										
ARF22/1-600										
ARF23/1-222										
ARF2/1-859										
PoptrARF8.1/1-827										
PoptrARF1.1/1-660										
PoptrARF7.3/1-1113										
PoptrARF2.3/1-792										
PoptrARF5.1/1-933										
PoptrARF5.2/1-944										
PoptrARF7.4/1-1137										
PoptrARF16.1/1-669										
PoptrARF1.2/1-662										
PoptrARF9.1/1-666										
PoptrARF2.1/1-854										
PoptrARF16.3/1-700										
PoptrARF3.1/1-709										
PoptrARF6.2/1-914										
PoptrARF6.4/1-953										
PoptrARF17.1/1-594										
PoptrARF2.4/1-879										
PoptrARF16.2/1-708										
PoptrARF2.2/1-852										
PoptrARF10.1/1-708										
PoptrARF3.3/1-109										
PoptrARF9.2/1-670										
PoptrARF6.1/1-884										
PoptrARF4/1-713										
PoptrARF6.3/1-163										
PoptrARF3.2/1-714										
PoptrARF9.3/1-579										
PoptrARF6.5/1-907										
PoptrARF9.4/1-632										
PoptrARF7.1/1-1047										
PoptrARF7.2/1-1093										
PoptrARF8.2/1-816										
PoptrARF16.6/1-91										
PoptrARF16.4/1-701										
PoptrARF16.5/1-536										
PoptrARF10.2/1-713										
PoptrARF17.2/1-592										
PoptrARF2.6/1-724										
PoptrARF2.5/1-614										

MP L P S C R R R E Q R P R Q S C F Q N P E C H L R L R A R Q Q R P Q E S A V Q S T L S S S I Y F S D H F V L T P P P F Q I T P T S I Q N S T W L P I P T S P A V A A P P L P R C L G G V D G E G A P C A P P P S P

Consensus

	110	120	130	140	150	160	170	180	190	200	210
LOC_Os01g70270.1 11667.m07038_altsplice/1-809											
LOC_Os01g13520.1 11667.m01333/1-700											
LOC_Os02g06910.1 11668.m00642/1-909											
LOC_Os02g35140.1 11668.m03331/1-757											-MKL
LOC_Os02g41800.1 11668.m03990/1-784											
LOC_Os04g57610.1 11670.m05695/1-819											-MSSS
LOC_Os04g43910.1 11670.m04249/1-696											
LOC_Os04g49230.1 11670.m04840/1-294											
LOC_Os04g36060.1 11670.m03490/1-1674											
LOC_Os05g43920.1 11682.m04181/1-700											
LOC_Os06g09660.1 11680.m00929/1-1056											
LOC_Os06g46410.1 11680.m04579/1-918											-MRL
LOC_Os06g47150.1 11680.m04656/1-731											
LOC_Os06g48950.1 11680.m04850/1-1116											
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056											
LOC_Os11g32110.1 11687.m02952/1-812											-MATAEVGGGGGEGD
LOC_Os12g41950.1 11686.m04180/1-900											-M
LOC_Os12g29520.1 11686.m02861_altsplice/1-842											-MAAAA
LOC_Os04g56850.1 11670.m05610_altsplice/1-956											-MASSQEKAKTGV
LOC_Os05g48870.1 11682.m04683_altsplice/1-696											-MTGI DLNTVEE
LOC_Os01g48060.1 11667.m04657_altsplice/1-723											-MVGIDLNTVEE
LOC_Os01g54990.1 11667.m05421/1-656											
LOC_Os10g33940.1 11676.m02996/1-699											
LOC_Os02g04810.1 11668.m00419/1-1094											-MASMKQQTPASSAVTA
LOC_Os04g59430.1 11670.m05892/1-530											
LOC_Os07g08520.1 11673.m00789/1-729											
LOC_Os07g08530.1 11673.m00790/1-408											
LOC_Os07g08600.1 11673.m00797/1-525											I PAGPASSTVSAASSAPATRDYLGGITGHRLQGEDDSGGHEYTSVSGIHVPSSSGVAFVSFGQCPQEVISTDLCLGLGPAAATAPLQPPPPPPAAAAAGSPSATTTP
ARF1/1-665											
ARF3/1-608											-MGGLIDLNVMTTEEDETQTPTPSSASG
ARF4/1-788											-MEFDLNTIEIAEVEEEENDDVGVGVGGGTRIDKGRLGIS
ARF5/1-902											-MMASLSCVEDKMKTSCLVNGGGITTT
ARF6/1-933											
ARF7/1-1165											
ARF8/1-811											
ARF9/1-638											
ARF10/1-693											
ARF11/1-601											
ARF12/1-593											
ARF13/1-623											
ARF14/1-605											
ARF15/1-598											
ARF16/1-670											
ARF17/1-585											
ARF18/1-602											
ARF19/1-1086											
ARF20/1-615											
ARF21/1-606											
ARF22/1-600											
ARF23/1-222											
ARF2/1-859											-MASSEVSMKGNRGGDNFSSSGFSDPKETRNVSVA
PoptrARF8.1/1-827											
PoptrARF1.1/1-660											
PoptrARF7.3/1-1113											
PoptrARF2.3/1-792											-MINGSYSVD
PoptrARF5.1/1-933											-MGSAAEEKIKTGGINGA
PoptrARF5.2/1-944											-MGSVEEKIKTGGLANGA
PoptrARF7.4/1-1137											-MKSP
PoptrARF16.1/1-669											
PoptrARF1.2/1-662											
PoptrARF9.1/1-666											-M
PoptrARF2.1/1-854											-MASSEISAKANSNGNIKGGESFSSGY
PoptrARF16.3/1-700											
PoptrARF3.1/1-709											-MVGMI DLNTTEEDETTPSSGSL
PoptrARF6.2/1-914											
PoptrARF6.4/1-953											-M
PoptrARF17.1/1-594											
PoptrARF2.4/1-879											-MELRAGAGVFRWLKVKVPMI IHNDFLSFS
PoptrARF16.2/1-708											-MFEAKLRDGYTQVLQIPKLHS
PoptrARF2.2/1-852											-MASSEISAKANSNGNIKGGESFTSGYSE
PoptrARF10.1/1-708											
PoptrARF3.3/1-109											
PoptrARF9.2/1-670											-MAMNR
PoptrARF6.1/1-884											
PoptrARF4/1-713											-MEIDLNPVTEVEK
PoptrARF6.3/1-163											
PoptrARF3.2/1-714											-MVGMI DLNTIEEDETTPSCGSL
PoptrARF9.3/1-579											
PoptrARF6.5/1-907											
PoptrARF9.4/1-632											
PoptrARF7.1/1-1047											
PoptrARF7.2/1-1093											
PoptrARF8.2/1-816											
PoptrARF16.6/1-91											
PoptrARF16.4/1-701											
PoptrARF16.5/1-536											
PoptrARF10.2/1-713											
PoptrARF17.2/1-592											
PoptrARF2.6/1-724											-MGKKRSVAGEAKTSRSHEVG
PoptrARF2.5/1-614											

Consensus

-----DL-----MMA+I++++++S++E++KT+GSASSA

LOC_Os01g70270.1|11667.m07038_altsplice/1-809
LOC_Os01g13520.1|11667.m01333/1-700
LOC_Os02g06910.1|11668.m00642/1-909
LOC_Os02g35140.1|11668.m03331/1-757
LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-819
LOC_Os04g43910.1|11670.m04249/1-696
LOC_Os04g49230.1|11670.m04840/1-294
LOC_Os04g36060.1|11670.m03490/1-1674
LOC_Os05g43920.1|11682.m04181/1-700
LOC_Os06g09660.1|11680.m00929/1-1056
LOC_Os06g46410.1|11680.m04579/1-918
LOC_Os06g47150.1|11680.m04656/1-731
LOC_Os06g48950.1|11680.m04850/1-1116
LOC_Os08g40900.1|11674.m04106_altsplice/1-1056
LOC_Os11g32110.1|11687.m02952/1-812
LOC_Os12g41950.1|11686.m04180/1-900
LOC_Os12g29520.1|11686.m02861_altsplice/1-842
LOC_Os04g56850.1|11670.m05610_altsplice/1-956
LOC_Os05g48870.1|11682.m04683_altsplice/1-696
LOC_Os01g48060.1|11667.m04657_altsplice/1-723
LOC_Os01g54990.1|11667.m05421/1-656
LOC_Os10g33940.1|11676.m02996/1-699
LOC_Os02g04810.1|11668.m00419/1-1094
LOC_Os04g59430.1|11670.m05892/1-530
LOC_Os07g08530.1|11673.m00789/1-729
LOC_Os07g08530.1|11673.m00790/1-408
LOC_Os07g08600.1|11673.m00797/1-525
ARF1/1-665
ARF3/1-608
ARF4/1-788
ARF5/1-902
ARF6/1-933
ARF7/1-1165
ARF8/1-811
ARF9/1-638
ARF10/1-693
ARF11/1-601
ARF12/1-593
ARF13/1-623
ARF14/1-605
ARF15/1-598
ARF16/1-670
ARF17/1-585
ARF18/1-602
ARF19/1-1086
ARF20/1-615
ARF21/1-606
ARF22/1-600
ARF23/1-222
ARF2/1-859
PoptrARF8.1/1-827
PoptrARF1.1/1-660
PoptrARF7.3/1-1113
PoptrARF2.3/1-792
PoptrARF5.1/1-933
PoptrARF5.2/1-944
PoptrARF7.4/1-1137
PoptrARF16.1/1-669
PoptrARF1.2/1-662
PoptrARF9.1/1-666
PoptrARF2.1/1-854
PoptrARF16.3/1-700
PoptrARF3.1/1-709
PoptrARF6.2/1-914
PoptrARF6.4/1-953
PoptrARF17.1/1-594
PoptrARF2.4/1-879
PoptrARF16.2/1-708
PoptrARF16.2/1-852
PoptrARF10.1/1-708
PoptrARF3.3/1-109
PoptrARF9.2/1-670
PoptrARF6.1/1-884
PoptrARF4/1-713
PoptrARF6.3/1-163
PoptrARF3.2/1-714
PoptrARF9.3/1-579
PoptrARF6.5/1-907
PoptrARF9.4/1-632
PoptrARF7.1/1-1047
PoptrARF7.2/1-1093
PoptrARF8.2/1-816
PoptrARF16.6/1-91
PoptrARF16.4/1-701
PoptrARF16.5/1-536
PoptrARF10.2/1-713
PoptrARF17.2/1-592
PoptrARF2.6/1-724
PoptrARF2.5/1-614

Consensus

MESGSSVSSAGALPGG+EGEKKALYSELWHACAGPLVGSPL+VGSRV+YFPQGHSEQ

LOC_Os01g70270.1|11667.m07038_altsplice/1-809
LOC_Os01g13520.1|11667.m01333/1-700
LOC_Os02g06910.1|11668.m00642/1-909
LOC_Os02g35140.1|11668.m03331/1-757
LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-819
LOC_Os04g43910.1|11670.m04249/1-696
LOC_Os04g49230.1|11670.m04840/1-294
LOC_Os04g36060.1|11670.m03490/1-1674
LOC_Os05g43920.1|11682.m04181/1-700
LOC_Os06g09660.1|11680.m00929/1-1056
LOC_Os06g46410.1|11680.m04579/1-918
LOC_Os06g47150.1|11680.m04656/1-731
LOC_Os06g48950.1|11680.m04850/1-1116
LOC_Os08g40900.1|11674.m04106_altsplice/1-1056
LOC_Os11g32110.1|11687.m02952/1-812
LOC_Os12g41950.1|11686.m04180/1-900
LOC_Os12g29520.1|11686.m02861_altsplice/1-842
LOC_Os04g56850.1|11670.m05610_altsplice/1-956
LOC_Os05g48870.1|11682.m04683_altsplice/1-696
LOC_Os01g48060.1|11667.m04657_altsplice/1-723
LOC_Os01g54990.1|11667.m05421/1-656
LOC_Os10g33940.1|11676.m02996/1-699
LOC_Os02g04810.1|11668.m00419/1-1094
LOC_Os04g59430.1|11670.m05892/1-530
LOC_Os07g08520.1|11673.m00789/1-729
LOC_Os07g08530.1|11673.m00790/1-408
LOC_Os07g08600.1|11673.m00797/1-525
ARF1/1-665
ARF3/1-608
ARF4/1-788
ARF5/1-902
ARF6/1-933
ARF7/1-1165
ARF8/1-811
ARF9/1-638
ARF10/1-693
ARF11/1-601
ARF12/1-593
ARF13/1-623
ARF14/1-605
ARF15/1-598
ARF16/1-670
ARF17/1-585
ARF18/1-602
ARF19/1-1086
ARF20/1-615
ARF21/1-606
ARF22/1-600
ARF23/1-222
ARF2/1-859
PoptrARF8.1/1-827
PoptrARF1.1/1-660
PoptrARF7.3/1-1113
PoptrARF2.3/1-792
PoptrARF5.1/1-933
PoptrARF5.2/1-944
PoptrARF7.4/1-1137
PoptrARF16.1/1-669
PoptrARF1.2/1-662
PoptrARF9.1/1-666
PoptrARF2.1/1-854
PoptrARF16.3/1-700
PoptrARF3.1/1-709
PoptrARF6.2/1-914
PoptrARF6.4/1-953
PoptrARF17.1/1-594
PoptrARF2.4/1-879
PoptrARF16.2/1-708
PoptrARF2.2/1-852
PoptrARF10.1/1-708
PoptrARF3.3/1-109
PoptrARF9.2/1-670
PoptrARF6.1/1-884
PoptrARF4/1-713
PoptrARF6.3/1-163
PoptrARF3.2/1-714
PoptrARF9.3/1-579
PoptrARF6.5/1-907
PoptrARF9.4/1-632
PoptrARF7.1/1-1047
PoptrARF7.2/1-1093
PoptrARF8.2/1-816
PoptrARF16.6/1-91
PoptrARF16.4/1-701
PoptrARF16.5/1-536
PoptrARF10.2/1-713
PoptrARF17.2/1-592
PoptrARF2.6/1-724
PoptrARF2.5/1-614

320 330 340 350 360 370 380 390 400 410 420
- - - - - VEASMN - QVADSQMRL YDL PSKL - - - - - LCRVL NVLEL KAEQDTDEVYA QVML MP - EPEQN - - - - - EM
- - - - - LQEPD DP - ALL AEQ I KMFQVPYK I - - - - - LCKVVNVLEL KAEETDEVF AQI TLLP - DPDQENL - - - - - PT
- - - - - VAASTNK - EMESQ I PNY PNL PPQL - - - - - I CQL HNV T MHADAETDEVYA QMT LQP - LSPQELK - - - - - LT
- - - - - LEASTD - QQL DQHL PL FNL PSK I - - - - - LCKVVNVLEL RAETDSDEVYA QI MLQP - EADQNE - - - - - LT
- - - - - ALGLAAPEL SAARVPALV - - - - - PCRVASVRYMADPDTDEVFA R IRLVP - LRA - - - - - AE
- - - - - VAASTNK - EVEGH I PNY PNL PAQL - - - - - I CQL HDVT MHADVETDEVYA QMT LQP - LNPQE QN - - - - - VG
- - - - - ASAAVDLS SARVPPLV - - - - - PCRVAVVRF MADAEDEVFAK IRLVPLRPGDAVVD - - - - - VG
- - - - - ATASSSSPT SVSPSAT - - - - - ASAAAST SASGSPFRSSDGA - - - - - GA
- - - - - LEASTN - QQL DQY LPMFNL PSK I - - - - - LCSVNVNLEL RAEADSDEVYA QI MLQP - EADQSE - - - - - LT
- - - - - DGGGGEVPPAGAAAVPPHV - - - - - ACRVVGVLEL RADAATDEVYA RALALVAEGE MLQRNFREGG - - - - - GE
- - - - - VAASTMH - KELDN I PGYPSL PSKL - - - - - ICKLLSLTLHADSETDEVYA QMT LQP - VNKYDRDA - - - - - ML
- - - - - VAASTNK - EMESQ I PNY PNL PPQL - - - - - I CQL HNV T MHADAETDEVYA QMT LQP - LSPQELK - - - - - LT
- - - - - AQGHGPVEFPGGRVPALV - - - - - LCRVAGVRF MADPDTDEVFAK IRLVPRANEQGYA - - - - - GD
- - - - - ADPETDEVYA QMT LQP - VTSYGKEA - - - - - LT
- - - - - DADAQ I PSYPNL PSKL - - - - - I CILHSVTMLADPDTDEVYA RMT LQP VNSCDKET - - - - - LL
- - - - - VEASTN - QVGEQRMQL YNL PWM I - - - - - LCEVMNVLEL KAEPTDEVYA QLTLLP - ESKQQEDN - - - - - GS
- - - - - VAASTNK - EVDAQ I PNY PNL PPQL - - - - - I CQL HNV T MHADAETDEVYA QMT LQP - LSPQEQK - - - - - LT
- - - - - VEASTNQ - VAEQGGAPL YNL PWM I - - - - - PCKVMNVLEL KAEPTDEVYA QLTLLP - EKQQDGNSSGNGNVVS - KD
- - - - - VAATTNRK - IPNSR I PNY PNL PSQL - - - - - LCQVHN I TLHADKDTDEVYA QMT LQP VNSD DVF - - - - - LT
- - - - - LGDAPAAAAAAA VPPHV - - - - - FCRVVDVTL LADAAETDEVYA QLSLYPEKEEVARR - - - - - AD
- - - - - LGAAPGSGPGA VPPHV - - - - - FCRVVDVSL HADAATDEVYA QVSLVADNEEVERRMREG - - - - - DG
- - - - - AGAGGIGRGEVAVAL PPHV - - - - - ACRVVDVLELC - - - - - VFERNLHGG - - - - - GI
- - - - - ADGGGAAAAAAEL GPRAL PPLV - - - - - LCRVEGVQFLADRRSDEVYA K IRLAPVAPGEAEFREP - - - - - DE
MRKKVYKSN I I L RNAFSML L ETAEYLL NLSFLF QVAASMQL - - - - - ICLLHGVLHADPDTDEVYA QMT LQP - VNTYGKEA - - - - - LQ
- - - - - CPAPL PDPLPSAHRFF - - - - - LCTIATVDLSADTTTGEPYAT I SLLPLRHDAPA - - - - - PA
- - - - - CAEDPALL SRL PDP IHPVPCTVADLVLDVAESGEAYAT I SLLP - - - - - GS
- - - - - FCQFPAPLLEQLAVPG - PRVF - - - - - LCTVAAVRLRADAL TNEAYAEITLDP - VADHDV - - - - - PR
- - - - - CLSRPQEP L PG - RIF - - - - - LCKVTDVRLGA - AATNEALAT I SLVP - IADDHAFQ - - - - - LQ
- - - - - LEASMH - QQL EQMPSFNL PSK I - - - - - LCKV I NIQRRAEPTDEVYA QI TLLP - LSPQEQK - - - - - PT
- - - - - APDFSAA I YGL PPHV - - - - - FCRI L DVKLHAETT LDEVYA QVSL LPESED I ERKVREG - - - - - I I
- - - - - DAMVSYSPL E I PKFDL NPQ I - - - - - VCRVVNVQL LANKDTEDEVY QTLTLLP - LQEF SMLNGEGKE - - - - - VK
- - - - - VAVSTRR - SAT TQVPNY PNL PSQL - - - - - MCQVHNVT LHADKDSDE IYAQMSLQPVHSEKDV - - - - - FP
- - - - - VAASTNK - EVDAH I PNYPSL HPQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LNAQE QK - - - - - LT
- - - - - VAASTMQ - KQTF I PSYPNL PSKL - - - - - I CML HNV TL NADPETDEVYA QMT LQP - VNKYDRDA - - - - - LL
- - - - - VAATTNK - EVDGH I PNYPSL PPQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LTPQEQK - - - - - LT
- - - - - LEASTQQ - VDLNTMKPL FVL PPK I - - - - - LCNVMNVSLQAEKDTDEVYA QI TL I PVGT EYDE - - - - - PM
- - - - - AHAPPDFHAPRVPL I - - - - - LCRVVSVKFLADAETDEVFAK I TLLP - LPGA ND LDELND - - - - - AV
- - - - - LVASTNQ - GVVDQE I PVFNL PPK I - - - - - LCRVL SVTLKAEHETDEVYA QI TLLP - EEDQSE - - - - - PT
- - - - - VETSTR - EELNEL QP I CDL PSKL - - - - - QCRV I A IHLKVENNSDETYAEITLMP - - - - - D
- - - - - IENSTR - DELDH I RP I FDL PSKL - - - - - RCRVVA I DRKV DKTDEVYA QI SLM P - - - - - D
- - - - - VEASTR - EELNEL QP I CDF PSKL - - - - - QCRV I A IQLKVENNSDETYAEITLMP - - - - - D
- - - - - VEASTR - EELNEL QP I CDL PSKL - - - - - QCRV I A IHLKVENNSDETYAKITLMP - DTVSYENL - - - - - D
- - - - - AYDCVD FGNLP I PPMV - - - - - LCRVLA I KYMADAESDEVFAK LRLP - LQDESY - - - - - DH
- - - - - CCPLLSTLPSSTSPV - - - - - PCIITS IQLLADPVTDEVFAHL I LQP - MTQQFT - - - - - PT
- - - - - LVASTNQ - GINSEE I PVFDL PPK I - - - - - LCRVL DVT LKAEHETDEVYA QI TLLP - EEDQSE - - - - - PT
- - - - - VAASTMQ - KQTF I PNY PNL PSKL - - - - - ICLLHSVTLHADTETDEVYA QMT LQP - VNKYDREA - - - - - LL
- - - - - VDASTR - EELNEL QP I CDL PSKL - - - - - QCRV I A IHLKVENNSDETYAEITLMP - - - - - D
- - - - - VQASTR - EELNEL QP I CDL PSKL - - - - - QCRV I A IHLKVENNSDE IYAEITLMP - - - - - D
- - - - - VEASTR - EELNEL KP I CDL PSKL - - - - - QCRV I A IQLKVENNSDETYAEITLMP - - - - - D
- - - - - VEASTR - EELNEL QPNCOL PSKL - - - - - QCRV I A IHLKVENNSDETYAEITLMP - - - - - D
- - - - - VEASTN - QAEEQQMPL YDL PSKL - - - - - LCRV I NVDLKAEADTDEVYA QI TLLP - EANQDEN - - - - - AI
- - - - - VAATTNK - EVDAH I PNYPSL PPQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LTPQEQK - - - - - LT
- - - - - LEASMH - EQGEEQQMPSFNL PSK I - - - - - LCKVVNVQRAEPETDEVYA QI TLLP - EPDQSE - - - - - VT
- - - - - VAASLKK - DVNAQ I PNY PNL PSKL - - - - - LCLLHNVTLHADPETDEVYVQMT LQP - VSSFDDKA - - - - - LL
- - - - - VEASTNQ - VADDQOMPAYNL PPK I - - - - - LCRVVNVQLKAE LDTDEVFAQV ILLP - LAEDQVD - - - - - LV
- - - - - VAVSTRK - SATSQ I PNY PNL PSQL - - - - - LCQVHNVT LHADKDTDE IYAQMSLQPVNSEKDV - - - - - FP
- - - - - VAVSTRR - SATSQ I PNY PNL PSQL - - - - - LCQVHNVT LHADKDTDE IYAQMSLQPVNTEKDV - - - - - FP
- - - - - VAA SMK K - DADAQ I PNY PNL PSKL - - - - - LCLLHNVTLHADPETDEVYA QMT LQP - VSSFDKA - - - - - LL
- - - - - AQGSVEFGHFQ I PAL I - - - - - PCKVSA I KYMADPETDEVYAK IRLIP - LNNSDLMLGH - - - - - GC
- - - - - LEASMH - QGMEPQML FNL PSK I - - - - - LCKVVNVQRAEPETDEVYA QI TLLP - EPDQSE - - - - - VT
- - - - - LEASTN - QELNQRVPL FNL PSK I - - - - - LCRV I HTQLLAEQDTDEVYA QI TLLP - ESDQIE - - - - - TT
- - - - - VEASTN - QVADQOMPL YDL PPK I - - - - - LCRVVNVQLKAEPTDEVFAQV TLLP - LHNQDEN - - - - - AS
- - - - - ACEPVDFRNLP GASHT - - - - - LCRVSA I KFMADPETDEVFAK IRLVP - INSNEIDL - - - - - DD
- - - - - LPDLPLGI YDL PPHV - - - - - FCRVVDVKLHAAEASDDVYAQVSLYPESEE I EQKLREG - - - - - VF
- - - - - VAASTNK - EVDAR I PNYPSL PPQL - - - - - I CQL HNV T MHADVETDEVYA QLTLLP - LSPQEQK - - - - - LT
- - - - - VAASTNK - EVDAH I PNYPSL PAQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LSPQEQK - - - - - LT
- - - - - SSSSTAPHP PFLTNALSKPS I - - - - - PCQI SAVDFLADPVTDEVFTLLP I LPLDNFNSNPLSFL - - - - - EP
- - - - - VRFHLLSPVEASTNQ - VADDQOMPAYNL SPK I - - - - - LCRVVNVQLKAE LDTDEVFAQV ILLP - ETQQDVE - - - - - LV
- - - - - AQGSVDFGHFQ I PAL I - - - - - PCKVSA I KYMAEPETDEVYAK IRLTP - SSNSDL MFGD - - - - - GC
- - - - - VEASTN - QVADQOMPL YNL LPK I - - - - - LCRVVNVQLKAEPTDEVFAQV TLLP - EHNQDES - - - - - VL
- - - - - SQSPVDFDQ - R I PSLV - - - - - LCRVASVKFLADPGTDEVFAK I SLPV - LPDADLD - - - - - IS
- - - - - LEASTN - QELNQRVPL FNL PSK I - - - - - LCRV I HTQLLAEQDTDEVYA QI TLLP - ESDQIE - - - - - PT
- - - - - VAASTNK - EVNAQ I PSYPSL PAQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LSPQEQK - - - - - LT
- - - - - LASSS - - PFSHRDMPNFDLHPQ I - - - - - FCKVVNVQLLANRENDEVYTR L TLLPQPEVVGQDL EGKE - - - - - LQ
- - - - - LPDLPLAVYDLPSHV - - - - - FCRVVDVKLHAAEASDEVYAQVSLYPESEE I EQKLREG - - - - - IF
- - - - - LEASTN - QELNQQ I PRFNL PPK I - - - - - LCRVVNIQLLAEQDTDEVYA QI TLLP - EVDQTE - - - - - PT
- - - - - VAASTNK - EVDAH I PNYPSL PPQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LSPQEQK - - - - - LT
- - - - - LEASTN - QELNQQ I PRFNL PPK I - - - - - LCRVVNIQLLAEQDTDEVYA QI TLLP - EVDQTE - - - - - LT
- - - - - VAASTMQ - KETDFVPSYPNL TSKL - - - - - I CML HNV TLHADVETDEVYA QMT LQP - VSKYDKEA - - - - - LL
- - - - - VAASTMQ - KETDF I PSYPNL PSKL - - - - - I CML HNV TLHADVETDEVYA QMT LQP - VSKYDKEA - - - - - LL
- - - - - VAATTNK - EVDAH I PNYPSL PPQL - - - - - I CQL HNV T MHADVETDEVYA QMT LQP - LTPQEQK - - - - - LT
- - - - - ACEPVDFRNLP RVSHN - - - - - LCRVSD I KFMADPETDEVFAK IRLVP - INSNEIDL - - - - - DD
- - - - - SQSPVNFQP - R I PSL I - - - - - LCRVATVKFLADPDTDEVYAK I GFVP - LPNTDLDF - - - - - AH
- - - - - SSSSTAPHP PFLSNALSKPL I - - - - - SCQI SAVDFLADPVTDEVF IRL LLLP - LNNHSDNPLSFL - - - - - EP
- - - - - VAAARMN - EEGKEMEP I YDLPYK I - - - - - LCKVVHVELKAEAGTDEVFA R I TLLP - VAEDEL - - - - - LT
- - - - - VAAFLN - EDSKTAMP I YDLPYK I - - - - - LCKVVHVQLKAEAKTDEVFAH I TLLP - - - - - VAE

Consensus



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LOC_Os02g35140.1|11668.m03331/1-757
LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-189
LOC_Os04g46410.1|11670.m04249/1-696
LOC_Os04g49230.1|11670.m04840/1-294
LOC_Os04g36060.1|11670.m03490/1-1674
LOC_Os05g43920.1|11682.m04181/1-700
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LOC_Os06g46410.1|11680.m04579/1-918
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LOC_Os11g32110.1|11687.m02952/1-812
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LOC_Os05g48870.1|11682.m04683. *altsplce*/1-696
LOC_Os01g48060.1|11667.m04657. *altsplce*/1-723
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LOC_Os10g33940.1|11676.m02996/1-699
LOC_Os02g04810.1|11668.m00419/1-1094
LOC_Os04g59430.1|11670.m05892/1-530
LOC_Os07g08520.1|11673.m00789/1-729
LOC_Os07g08530.1|11673.m00790/1-408
LOC_Os07g08600.1|11673.m00797/1-525
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ARF4/1-708
ARF5/1-902
ARF6/1-933
ARF7/1-1165
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ARF9/1-638
ARF10/1-693
ARF11/1-601
ARF12/1-593
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ARF14/1-605
ARF15/1-598
ARF16/1-670
ARF17/1-585
ARF18/1-602
ARF19/1-1086
ARF20/1-615
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ARF22/1-600
ARF23/1-222
ARF2/1-859
PoptrARF8.1/1-827
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PoptrARF7.3/1-1113
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PoptrARF9.4/1-632
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PoptrARF16.5/1-536
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PoptrARF17.2/1-592
PoptrARF2.6/1-724
PoptrARF2.5/1-614

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EGDGEEDGEATVKMTTPHMFCKLTASDTSTHGGSVPR-RAAEDCFPPL-----DYTEQQRPSQELVAKDLHGSEWFRHIFR-----GQPRRH
SPDPCPPEP----AKPTIHSCFKILTASDTSTHGGSVSLR-KHATECLPPL-----DMSQATPTQELAARDLHGFEWFRKHIFR-----GQPRRH
--DAYLLPAELGTASKQPTNYFCKLTASDTSTHGGSVPR-RAAEKVFPPL-----DYSQTPPAQELIARDLDHNEWHFRHIFR-----GQPKRH
-----AKQTVHSFCKILTASDTSTHGGSVSLR-KHATECLPPL-----DMSQATPTQELAARDLHGVEWFRKHIFR-----GQPRRH
ASDLGQKQS----RQPTTEFFCKLTASDTSTHGGSVPR-RAAEKIFPPL-----DFSMPQPAQELVARDLDHNTWTFRHIYR-----GQPKRH
ASDMGLKQN----RQPTTEFFCKLTASDTSTHGGSVPR-RAAEKIFPPL-----DFSMPQPAQELVARDLDHNTWTFRHIYR-----GQPKRH
--ETFLPLDLGMPSKQPTNYFCKLTASDTSTHGGSVPR-RAAEKVFPPL-----DFTQPPAQELIARDLDHVEWFRHIFR-----GQPKRH
-----ILESTFSRL-----DYTEAPPEETILAKDVHGETWFRHIFR-----GAPRRH
QEVAVNGGMEAAQDNKPKVSFCKLTQSDANNNGGFSVPR-YCAEMIFPPL-----DYTEADPPVQTLAKDVHGETWFRHIFR-----GTPRRH
-----XVSFAFCKLTASDTSTHGGSVPR-YCAEMIFPPL-----DYTEADPPVQTLAKDVHGETWFRHIFR-----GTPRRH
DRGLCGNGNDGDCSPDKPAASFCKLTQSDANNNGGFSVPR-YCAETIFPPL-----DYSDDPPLQTVIARDVHGEVWFRHIFR-----GTPRRH
SRSEGGGVNDVDDENKILAFAKILTPSDANNNGGFSVPR-FCADSIFPPL-----NYQAEPPVQTLTVTDIGHISWDFRHIYR-----GTPRRH
--SSNKDGKSLPLHRKTCARSFTKKLTPTSDTKHGGFSVPR-RHADQCLPPL-----DKSQQPPQELIARDLHGFEWFRKHIFR-----GQPKRH
LSSNKDGESLLLHRKTRVLSFTKLTPTSDTSQGGFSVPK-RHAEESLPPL-----DKSQQPPAQELIARDLHGSEWFRHIFR-----GQPKRH

Consensus



LOC_Os01g70270.1|11667.m07038_altsplice/1-809
LOC_Os01g13520.1|11667.m01333/1-700
LOC_Os02g06910.1|11668.m00642/1-909
LOC_Os02g35140.1|11668.m03331/1-757
LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-819
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LOC_Os06g09660.1|11680.m00929/1-1056
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LL - - - - - QS - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRA MRQLSNVPSSV I S - - - - -
LL - - - - - TT - - GWSFTVTSKRL I SGDAFYVFL R - S - - - - ETGEQRVGVRR L VQKQSTMPASV I S - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSVLF I W - N - - - - DSNQL L L G I RRANRPQTVMPPSSV L S - - - - -
LL - - - - - TT - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRL MRQLNNMPSSV I S - - - - -
LL - - - - - TT - - GWSFTVFNQKLLVAGDS I VFL R - G - - - - DGGDL HVG I RRAKRGF CGGGGAEASL - - - - PGWDQY GGL MRGNASPCA AAKG - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSVLF I W - N - - - - EKNQL L L G I RRASRPQTVMPPSSV L S - - - - -
LL - - - - - TT - - GWSPFVFNKQLTAGDS I VFM R - D - - - - EGGN I HVGL RRAKRGFCS I GGDDES L SS I PGWDQY RGL MRRNATATATGGRTPP - - - - -
VM - - - - - TK - - GWSRFVFEKRLDAGDTVSFCRGA - - - - -
LL - - - - - TT - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ESSEL RVGVRRL MRQVNNMPSSV I S - - - - -
LL - - - - - TT - - GWSSTFVNKRLLVSGDAVLF L R - G - - - - DDGQL RLGVRRAVQL RNEALFEPVN - - - - -
LL - - - - - TT - - GWSVFVTSKRLLAGDSVLF I R - D - - - - EKSQ L L L G I RRATRPQPALSSSV L S - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSVLF I W - N - - - - DNNQL L L G I RRANRPQTVMPPSSV L S - - - - -
LL - - - - - TT - - GWSFTVFNQKLLVAGDS I VFM R - T - - - - ENGDL CVG I RRAKKG VGGVGFEL PPPPPPPPTPAAGGNYGGFSMFLRGDDDGKNMAAARGKV - - - - -
LL - - - - - TT - - GWSL FVSGKRLFAGDSV I FVR - D - - - - EKQQL L L G I RRANRQPTN I SSSV L S - - - - -
LL - - - - - TT - - GWSL FVSGKRLLAGDSVLF I R - D - - - - AKQQL L L G I RRANRQPTN L SSSV L S - - - - -
LL - - - - - QS - - GWSVFVSAKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRA MRQQTNPVSSV I S - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSV I F I W - N - - - - DNNQL L L G I RRANRQQTVMPPSSV L S - - - - -
LL - - - - - QS - - GWSVFVSAKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRA MRQQAN I PSSV I S - - - - -
LL - - - - - TT - - GWSL FVSAKRLKAGDSVLF I R - D - - - - EKSQ L L L GVRRA TRQQTMLSSSV L S - - - - -
LL - - - - - TT - - GWSAFVFNKKLLVSGDAVLF L R - G - - - - DDGEL RLGVRRAAQLKNGSAFPALY - - - - -
LL - - - - - TT - - GWSGF I NKKLLVSGDAVLF L R - G - - - - EDGEL RLGVRRAAQLKNASFPALH - - - - -
LL - - - - - TT - - GWSSTFVNKRLLVSGDAVLF L R - G - - - - DDGEL RLGVRRA TQLKNEA I FKAFS - - - - -
LL - - - - - TT - - GWSFTVFNQKLLVAGDS I VFL R - T - - - - RHGEL CVG I RRAKRMACGGMCEMSGWNAPYGGGGSAFLKEEESKL MKHG HGGGGY MKG - - - - -
LL - - - - - TT - - GWSL FVSGKRLFAGDSV I FVR - D - - - -
LL - - - - - TT - - GWSKFVNAQQLVAGDTVVFMMCG - - - - - APAPERKLLVGVRAARYSGESACNA - - - - -
MLGDLGVNDNDGWRGFGVKA RRLAT RDTVVFMRGGGDDGGDEL L VGVRRAPRARGHPR - - - - -
TL - - - - - AK - - VWKFRDDMD FVDGDSV I FMR - RRDDDDDDGEL YVGVRRQRTL ERPLRNTMRRYRPPTP - - - - -
MF - - - - - RN - - GWMEFSNANGLVTGDNAVFM R - R - - - - GNGEMFMAVRRTRNR - - - - -
LL - - - - - TT - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ENEEL RVGVRRA MRQQTNPSSV I S - - - - -
LL - - - - - TT - - GWSAFVFNKKLLVSGDAVLF L R - G - - - - DDGKL RLGVRRA S Q I EGTAA LSAQY - - - - -
LL - - - - - TT - - GWS I FVSQKLLVSGDAVLF L R - D - - - - EGGEL RLG I RRAARPRNGLPDS I I - - - - -
LL - - - - - TT - - GWSL FVSGKRLRAGDSVLF I R - D - - - - EKSQ L MVGVRRA NRQQTALPSSV L S - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSVLF I W - N - - - - DKNQL L L G I RRANRPQTVMPPSSV L S - - - - -
LL - - - - - TT - - GWSVFVSTKRLFAGDSVLF I R - D - - - - GKAQL L L G I RRANRQPALSSV I S - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSV I F I R - N - - - - EKNQL F L G I RHATRPQT I VPSSV L S - - - - -
LL - - - - - TT - - GWSFTVTSKRLVAGDTFVFL R - G - - - - ENGEL RVGVRRA NLQSSMPSSV I S - - - - -
LL - - - - - TT - - GWSFTVFNQKLL I AGDS I VFL R - S - - - - ESGL CVG I RRAKRGL GSNAGSDNPYPGSGFLR DDESTTTTSKLMMMKRNGNNDGNAAA - - - - -
LL - - - - - TT - - GWSFTVTSKRLVAGDSVFL R - G - - - - ETGDL RVGVRRLAKQSTMPASV I S - - - - -
LL - - - - - TT - - GWNAFTTSKLLVAGDV I FVR - G - - - - ETGEL RVG I RRA RHQQGN I PSS I VS - - - - -
MF - - - - - TSGGGWSVFATT KRL I VGD I FVLL R - G - - - - ENGEL RFG I RRAKHQQGHI PSS I VS - - - - -
LL - - - - - TT - - GWNAFTTSKLLVEGDV I FVR - G - - - - ETGEL RVG I RRAHQGN I PSS I VS - - - - -
LL - - - - - TT - - GWNEFTTSKLLVKGDV I FVR - G - - - - ETGEL RVG I RRA RHQQGN I PSS I VS - - - - -
LL - - - - - TT - - GWSNFFVNKLLVAGDS I VFM R - A - - - - ENGDL CVG I RRAKRGG I GNGPEYSAGWNP I GSGCGYSSLLREDESNSLRRSNCSLADR - - - - -
LL - - - - - TT - - GWSKFVNSKLL I AGDSVVFMR - K - - - - SADEMF I GVRRTPISSSDGGSSYYGGDEYNGYSSQSSVAKEDDGSPPKKTFRRS G - - - - -
LL - - - - - TT - - GWSFTVSSKRLVAGDAFVFL R - G - - - - ENGDL RVGVRRL ARHQSTMPTSV I S - - - - -
LL - - - - - TT - - GWSVFVSTKRLFAGDSVLFVR - D - - - - EKSQ LMLG I RRANRQPTLSSV I S - - - - -
SL - - - - - TT - - GWNEFTTSKLLVKGDV I FVR - G - - - - ETGEL RVG I RRA RHQQGN I PSS I VS - - - - -
SL - - - - - TT - - GWNEF I TSKLLVKGDV I FVR - G - - - - ETGEL RVG I RRA RHQQGN I PSS I VS - - - - -
LL - - - - - TT - - GWNAFTTSKLLVAGDV I FVR - G - - - - ETGEL RVG I RRAHQGN I PSS I S - - - - -
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LL - - - - - QS - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRA MRQQGNVPSSV I S - - - - -
LL - - - - - TT - - GWSVFVSAKRLVAGDSVLF I W - N - - - - EKNQL L L G I RRATRPQTVMPPSSV L S - - - - -
LL - - - - - TT - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRL MRQQTNPMPSSV I S - - - - -
LL - - - - - TT - - GWSL FVSGKRLFAGDSVLFMR - D - - - - EKQQL L L G I RRANRQPTN L SSSV L S - - - - -
LL - - - - - QS - - GWSL FVSAKLLVAGDAF I FL R - G - - - - ETEEL RVGVRRA LSQPSNVPPSSVMS - - - - -
LL - - - - - TT - - GWSL FVSGKRLKAGDSVLF I R - N - - - - EKS L MVGVRRA NRQQT L P SSV L S - - - - -
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LL - - - - - TT - - GWSNFFVNQKLLVAGDS I VFL R - A - - - - ENGDL CVG I RRAKRGI GGNCESSGWSNFAGYSGFFREDESKL MRRNGNGDM - - - - -
LL - - - - - TT - - GWSVFVSSKRLVAGDAF I FL R - G - - - - ENGEL RVGVRRL MRQQTNPMPSSV I S - - - - -
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LL - - - - - TT - - GWSAFVFNKKLLVSGDAVLF L R - G - - - - EDGEL RLGVRRAAQV KCGPTFPALW - - - - -
LL - - - - - TT - - GWSVFVSAKRL I AGDSVLF I W - N - - - - EKNQL L L G I KRATRPQTVMPPSSV L S - - - - -
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- - - - - G - - - - -
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LL - - - - - NT - - GWSNFFVNNKNSMLGTRLCF - - - - -
LL - - - - - TT - - GWSPFVNHKKLL I AGDSVVFMR - A - - - - ENGDL CVGVRRAKRTSGGPESLWNPAGGSSAVPSGGGAF LREDEHKL MRSASGNGNGSKSNESL M - - - - -
LL - - - - - TT - - GWSPFVNHKKLL I AGDSVVFFR - A - - - - ENGDL CVGVRRAKRTSGGPESLWNPAGGSSAVPSGGGAF LREDEHKL MRSASGNGNGSKSNESL M - - - - -
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LL - - - - - TT - - GWSKFVNNKLL I AGDSVVFMR - N - - - - LKGE MF I GVRRAVRLNNSARWREQ I AGGGGEGKVKVEEGFSMSG - - - - -
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Consensus



LOC_Os01g70270.1|11667.m07038_altsplice/1-809
LOC_Os01g13520.1|11667.m01333/1-700
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PoptrARF8.2/1-816
PoptrARF16.6/1-91
PoptrARF16.4/1-701
PoptrARF16.5/1-536
PoptrARF10.2/1-713
PoptrARF17.2/1-592
PoptrARF2.6/1-724
PoptrARF2.5/1-614

Consensus

-----FTGT I I GSENLDPV---WPESSWRS L KVRWDEPS-T I P-RPDR-VSPWK I---EPASSPPVN-----
-----FSGT I VGEGDLSL---QWGSSEWKS L KVQWDEVT-NVN-GPER-VSPWE I---ETCDGTAPA I NVP-----
-----YMG T I T G I S D L D P V---RWMNSHWRSVKVGWDEST-AGE-RQPR-VSLWE I---EPLTTFPMYP-----
-----FSGT I I GVGSMSTS---PWANSDWRS L KVQWDEPS-VVP-RPDR-VSPWEL---EPLAVSNSQPS-----
-----FMGTVASVQVADP I---RWPQSPWRL L QVTWDEPD-LLQ-NVKR-VSPWL V---ELVSSMPA I N L S S F-----
-----YMG T I T E V S D A D P V---RWPSSYWRSVKVGWDEST-AGE-RPPR-VSLWE I---EPLTTFPMYP-----
-----FMGT V A G Q A S D P V---RWPQSPWRL L QVTWDEPE-LLQ-NVKR-VCPLV V---ELVSSMPN L H L P S F-----
-----AGASSCRPR-----RPPR-----
-----FSGT I I G I G S V P A M S K S P W A D S D W K S L K V Q W D E P S - A I V - R P D R - V S P W E L ---EPLDASNPPQQ-----
-----SAGL I S G I S E V D P I---RWP G S R W K C L L V R W D D S T - D S S - H Q N R - V S P W E I ---E R V G S S V S V T H S L-----
-----YMG T I T G I G D L D P V---RWKNSHWRLN L QVWDEST-ASE-RRTR-VS I W E I---EPVATPFY I C P P P F-----FRPKL P K Q P G M P-----
-----YMG T I T G I S D L D A A---RWPNSHWRSVKVGWDEST-AGE-RQPR-VSLWE I---EPLTTFPMYP-----
-----FMGTVASVQVADP I---RWPNSPWRL L QVSWDEPD-LLQ-NVKR-VSPWL V---ELVSNM P A I H L A P F-----
-----YMG T I T G I S D L D P V---RWKNSQWRN L QVWDESA-AGE-RRNR-VS I W E I---EPVAAFF I C P P P F-----FGAKRPRQLD-----
-----YMG T I T G I S D L D P V---RWK T S H W R N I Q V A W D E A A - P T E - R R T R - V S L W E I ---E P I I A P F F I Y P S P L-----F T A K R P R L P G M T-----
-----F T G T I V G M G S D P A---G W P E S K W R S L K V R W D E A S - S I P - R P E R - V S P W Q I ---E P A V S P P P V N-----
-----YMG T I T S I S D L S D V---RWPNSHWRSVKVGWDEST-TGD-KQPR-VSLWE I---EPLTTFPMYP-----
-----F T G T I V G V G S D P S---G W A D S K W R S L K V R W D E A A - S V P - R P D R - V S P W Q I ---E P A N S P S P V N-----
-----Y T G T V V G I S D Y D M---R W P N S K W R N L Q V E W D E H G - Y G E - R P E R - V S I W D I ---E T P E N T L V F P S S-----
-----Y T G I I T G S G D T D P M---W H G S K W K C L L V R W D D D A - E F R - R P N R - V S P W E I ---E L T S S V S G S H S L-----
-----R T G I I I G S R E A D P M---W H G S K W K C L V V K W D D D V - E C R - R P N G - V S P W E I ---E L S G S V S G S H S L-----
-----S G M I A G S V E D P I---R W P G S K W R S L L V R W E D A T - D C N - S Q N R - V S P W E I ---E I V G G S I S V A H S L S-----
-----F M G T I S S V Q V A D P N---R W P N S P W R L L Q V T W D E P D - L L Q - N V K C - V S P W L V ---E L V S S I P P I H L G P F-----
-----YMG T I T G I S D L D P V---G W D E S A - A G E - R R N R - V S I W E I ---E P V A A P F F I C P P P F-----F G V K R P R Q L-----
-----L N G T L T L N R L H---Q Q I W R T L E V E W D A S A A S S S - M K N R E V N P W Q V---Q P V D F-----
-----I S G T V R T F D H L R P-----W R M L E V D W Q A A S P I S Y R I H R Q V N S Q V L R Q P Q A A T T S A V R I R D A I-----V A-----
-----P R G K V T A I A T-----G Q L W R N L E I V W D G N S-----E M D M S A N F W Q V---R P V E E V D I S P S-----
-----W A V-----
-----FSGT I V G V Q E N K S S---V W H D S E W R S L K V Q W D E P S - S V F - R P E R - V S P W E L ---E P L V A N S T P S S Q-----
-----S P G I I S G I S D L D P I---R W P G S K W R C L L V R W D D I V - A N G - H Q Q R - V S P W E I ---E P S G S I S N S G-----
-----C A G V V T G W C D L D Y---R W P N S K W R C L L V R W D E S F - V S D - H Q E R - V S P W E I ---D P S V S L P H L S-----
-----YMG T I V G I S D L D P L---R W P G S K W R N L Q V E W D E P G - C N D - K P T R - V S P W D I ---E T P E S L F I F S L-----
-----YMG T I T G I C D L D P T---R W A N S H W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T T F P M Y P-----
-----YMG T V T G I S D L D P V---R W K N S Q W R N L Q I G W D E S A - A G D - R P S R - V S V W D I ---E P V L T P F Y I C P P P F-----F R P R F S G Q P G M P-----
-----YMG T I T G I S D L D S V---R W P N S H W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T T F P M Y P-----
-----Y S G T V I G V K D C S P---H W K D S K W R C L E V H W D E P A - S I S - R P N K - V S P W E I ---E P F V N S E N V P-----
-----F M G T V S A V Q V A D P I---R W P N S P W R L L Q V A W D E P D - L L Q - N V K R - V S P W L V ---E L V S N M P T I H L S P-----
-----F T G T I I G S G D L S S---Q W P A S K W R S L Q I Q W D E P S - S I Q - R P N K - V S P W E I ---E P F S S P A T T P L-----
-----C F G T I I G V S D F S P---H W K C S E W R S L E V Q W D E F T - S F P - G P K K - V S P W D I ---E H L M P A I N V P-----
-----Y D G T I I G V N D M S P---H W K D S E W R S L K V Q W D E L S - P F L - R P N Q - V S P W D I ---E H L I P S S D I S-----
-----S F G T I I G V S D F S P---H W K C S E W R S L E V Q W D E F A - S F P - R P N Q - V S P W D I ---E H L T P M N V N S-----
-----Y F G T I I G V S N F S P---H W K C S D W R S L E V Q W D E F A - S F L - R P N K - V S P W E I ---E H L M P A L N V P-----
-----F M G T V S A V N S D P I---R W P N S P W R L L Q V A W D E P D - L L Q - N V K R - V N P W L V ---E L V S N V N H P I P L T S F-----
-----F Q G I V S S T Y Q E T G---P W R G S P W K Q L Q I T W D E P E - I L Q - N V K R - V N P W Q V---E I A A H A T Q L H T P-----
-----YMG T I V G S G D L S S---Q W P A S K W R S L Q V Q W D E P T - T V Q - R P D K - V S P W E I ---E P F L A T S P I S T P A-----
-----YMG T V T G I S D L D P V---R W K G S Q W R N L Q V G W D E S T - A G D - R P S R - V S I W E I ---E P V I T P F Y I C P P P F-----F R P K Y P R Q P G M P-----
-----Y F G T I I G V N D F S P---H W K C S E W R S L E V Q W D E F A - S F S - R P N K - V S P W E I ---E H L M S A L N V P-----
-----Y F G T I I G V S D F S P---H W K C S E W R S L E V Q W D E F A - S F S - R P N K - V S P W E I ---E H L V P A L N V P-----
-----Y F G T I I G V S D F S P---H W K C S E W R N L E V Q W D E F A - S F S - R P N K - V S P W E I ---E H L M P A L N V P-----

-----F T G T I V G I E E S D P T---R W P K S K W R S L K V R W D E T S - S I P - R P D R - V S P W K V ---E P A L A P P A L S-----
-----YMG T I T G I S D L D P V---R W P N S H W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T T F P M Y P-----
-----FSGT I V G V G D N I S S---G W A D S E W R S L K V H W D E P S - S I L - R P E R - V S P W D L ---E P L V A T T P S N-----
-----H M G T I T G I S D L D A V---R W K N S Q W R N L Q V G W D E S T - A G E - R R N R - V S I W E I ---E P V T A P F F I C P P P F-----F R S K H P R Q P G M P D-----
-----FSGT V I G V E E A D P K---K W P R S K W R C L K V R W D E T S - P V H - R P D R - V S P W K I ---E R A L A P S I P V P G C-----
-----YMG T I V G I S D L D P L---R W P G S K W R N L Q V E W D E P G - C S D - K Q N R - V S S W E I ---E T P E S L F I F-----
-----YMG T I V G I S D L D P L---R W P G S K W R N L Q V E W D E P G - C S D - K Q N R - V S S W E I ---E T P E S L F I F-----
-----H M G T I T G I S D L D A V---R W K N S Q W R N L Q V G W D E S T - A G E - R R N R - V S I W E I ---E P V T A P F F I C P P P F L D L S V Q G N Q E C Q W M A S S G L Y Y Q I D K T V L L W D-----
-----F M G T I S S V Q V A D P I---R W P N S P W R L L Q V A W D E P D - L L H - N V K R - V S P W L V ---E L V S N M P A I H L S P F-----
-----FSGT I V G V G D N I S S---G W A D S D W R S L K V Q W D E P S - S I M - R P E R - V S H W E L ---E P L V A T T P S N-----
-----FSGT I V G V E D F S P---H W N D S K W R S L K V Q W D E P A - S I S - R P D R - V S P W E I ---E P C V A S V A N L S-----
-----F T G T I V G I E D A D P S---R W K D S K W R C L K V R W D E T S - T I P - R P D R - V S P W K I ---E P A L A P P A L N-----
-----F M G T V C S V Q D A D P L---C W P G S P W R L L Q V T W D E P D - L L Q - N V K R - V S P W L V ---E L A S H M S A I H L S P F-----
-----Y T G L I T G I S E L D P T---R W P G S K W C L L V R W D D T E - A N - R H S R - V S P W E V ---E P S G S V S G S G S G S I S S S N-----
-----YMG T I T G I S D L D V A---R W P N S L W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T T F P M Y P-----
I M F D S R Y M G T I T G I S D L D P A---R W P N S H W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T T F P M Y P-----
-----F Q G T V S G T G L P D C G---A W R G S P W R M L Q I T W D E P E - V L Q - N V K R - V S P W Q V---E F V A T T L P Q D A-----
-----FSGT V I G V E E A D P K---K W P R S N W R C L K V H W D E T S - P V H - R P D R - V S P W K V ---E P A L A P S M D P V-----
-----F M G T I S S V Q F A D P I---R W P N S P W R L L Q V A W D E P D - L L Q - N V K R - V S P W L A ---E L V S N M P A I H L S P F-----
-----F T G T I V G I E D A D P G---R W K N S K W R C L K V R W D E T S - T M P - R P E R - V S P W K I ---E P A L A P P A L N-----
-----F M G T V A S V Q V A D P D---R W P N S P W R L L Q V T W D E P D - L L Q - T V K C - V S P W L V ---E L V S N M P V I H L S P F-----
-----Y T G I I T G V S E L D P A---R W P G S K W K C L L V S W-----
-----FSGT I V G V E D F S P---H W N D S K W R S L K V Q W D E P A - P I P - R P D R - V S P W E I ---E P Y L S S-----
-----YMG T I T G I S D L D V V---R W P N S Q W R S V K V G W D E S T - D G E - R Q P R - V S L W E I ---E P L T T F P T Y P-----
-----C S G V V T G T A D L D P Y---K W P N S K W R C L M R W D E D V - I S D - H Q E R - V S P W E I ---D A S V S L P P L I-----
-----N P P S-----
-----Y T G I I T G V S E L D P A---R W P G S K W K C L L V R W D D R E - A N - R L S R - V S P W E V ---E P S G S G S I S S S N-----
-----F T G T I V G V G D I S P---E W S G S I W R S L K I Q W D E P A - T I Q - R P E R - V S P W D I ---E P F A A P A S P N L-----
-----YMG T I T G I S D L D P V---R W P N S H W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T T F P M Y P-----
-----F T G T I V G V G D I S P---E W S G S I W R S L K I Q W D E P A - T I Q - R P E R - V S P W D I ---E P F A A P A S P N L-----
-----YMG T I T G I S D L D P V---R W K N S Q W R N L Q V G W D E S T - A S E - R P N R - V S I W E I ---E P V V T P F Y I C P P P F-----F R P K F P K Q P G M P-----
-----YMG T I T G I S D M D P V---R W K N S Q W R N L Q V G W D E S T - A G E - R P S R - V S I W E I ---E P V V T P F Y I Y P P P F-----F R P K F P K Q P G M P-----
-----YMG T I T G S D L D P V---R W P N S H W R S V K V G W D E S T - A G E - R Q P R - V S L W E I ---E P L T S F P M Y P-----

-----F M G T V C S V Q A A D S L---W M P H S P W R L L Q V T W D E P D - L L Q - N V K R - V S P W L V ---E L A S N M A A I H F P P F-----
-----F M G T V C S V Q A A D S L---W M P H S P W R L L Q V T W D E P D - L L Q - N V K R - V S P W L V ---E L A S N M A A I H F P P F-----
-----F M G T V T S V Q V A D P V---R W P N S P W R L L Q V A W D E P D - L L Q - N V K R - V S P W L V ---E L V S N M P V I H L S P F-----
-----F Q G T V S G T G L P D S G---A W R G S P W R M L Q I T W D E P E - V L Q - N A K R - V S P W Q V---E F V A T T P Q L Q A A-----
-----F E G T V V G T E D V D H I---R W P N S E W R I L K V K W D A A S E P F V - H Q E R - V S P W N I ---E P I E P I R K K H A S R L H-----L H K-----
-----C E G T V V G A E D V D H I---R W P N S E W R I L K V K W D A A S D A F V - R P E R - V S P W N I ---E P I E P I R K K H A S-----



	850	860	870	880	890	900	910	920	930	940	950
LOC_Os01g70270.1 11667.m07038_altsplice/1-809				- PLPLSRVKRPR -		- PNAPPASPES -		- P I -			- L TKEAA -
LOC_Os01g13520.1 11667.m01333/1-700				- LQSATKNKRPR -		- EPSE -		- T -			- I DLQSL -
LOC_Os02g06910.1 11668.m00642/1-909				- SPFPLRLKRPW -		- PTGLPS -					- L YGGKE -
LOC_Os02g35140.1 11668.m03331/1-757				- PQQPARNKRAR -		- PPAS -		- N -			- S IAPEL -
LOC_Os02g41800.1 11668.m03990/1-784				- SPPRKKPRILA -		- YPEF -		- P -			- F EQQLL -
LOC_Os04g57610.1 11670.m05695/1-819				- SLFPLRVKHPW -		- YSGVASLHDD -		- SNALMW -			- L RGVAGEGGFQ -
LOC_Os04g43910.1 11670.m04249/1-696				- SPPRKKPRNPP -		- YAEL -		- P -			- L EQQIF -
LOC_Os04g49230.1 11670.m04840/1-294											
LOC_Os04g36060.1 11670.m03490/1-1674				- PPLRNKRARPPASPVVAELPPSFDVDSQI		- SQPSNGNKS		- SDAPGTSSERSPL		- ESQSRQVRSCTKV	- I MQGMAVGRAVD -
LOC_Os05g43920.1 11682.m04181/1-700				- SSGSKRTLKHF -		- PQGSLDTP -					- F L NGNG -
LOC_Os06g09660.1 11680.m00929/1-1056	DDENEVESAFKRAMPWL	ADDFALKDVQ		- SALLFPGLSLVQ -		- VMAMQONPQMLTAASQT		- VQSPYL -			- NSNALA -
LOC_Os06g46410.1 11680.m04579/1-918				- SPFPLRLKRPW -		- PTGL -		- PS -			- L HGGKD -
LOC_Os06g47150.1 11680.m04656/1-731				- SPPRKKL CVPL -		- YPEL -		- P -			- I DGQFP -
LOC_Os06g48950.1 11680.m04850/1-1116	DESSEMENL LKRAMPW	GEEICIKDPQTQNTIM	PGLSLVQ			- VMNMNQSS -		- S -			- FANTAM -
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056	DDETEMDGL LKRAMPW	GEEICKKDLN	I QNSVVPGLNLAQ			- VMNMQHS -		- SS -			- L PGTVV -
LOC_Os11g32110.1 11687.m02952/1-812				- PLPVPRTKRLR -		- PNATALPADS -		- SA -			- I AKEAN -
LOC_Os12g41950.1 11686.m04180/1-900				- SAFPLRLKRPW -		- ASGLPMHGMFNGGGNDDFARYSSL		- MW -			- L RDGNR -
LOC_Os12g29520.1 11686.m02861_altsplice/1-842				- PLPAPRTKRAR -		- PNVLASSPDL -		- SA -			- VNKESA -
LOC_Os04g56850.1 11670.m05610_altsplice/1-956				- TLNSKRQCLPG -		- YGVSVPGMEIGSANMSSF		- PRAQG -			- NPYGVL -
LOC_Os05g48870.1 11682.m04683_altsplice/1-696				- TPHSKRLKPCL -		- PHVNP EYM -					- VPRGGG -
LOC_Os01g48060.1 11667.m04657_altsplice/1-723				- TPHSKRLKSCF -		- PQVNPDI V -					- L PNGSV -
LOC_Os01g54990.1 11667.m05421/1-656				- ASSSKRTLKCP -		- QGNLDVP -					- A L YGNG -
LOC_Os10g33940.1 11676.m02996/1-699				- SPPRKKL RVPP -		- HPDFP -					- FEGHLL -
LOC_Os02g04810.1 11668.m00419/1-1094	DDSEMENL FKRAMPW	GEEVCIKDQTQNTAP	GLSLVQ			- VMNMNRQSS -		- S -			- L ANTAA -
LOC_Os04g59430.1 11670.m05892/1-530				- PPLPMGLKISN -							
LOC_Os07g08520.1 11673.m00789/1-729				- TPQVQIMALPR -				- P -			- P PPTTT -
LOC_Os07g08530.1 11673.m00790/1-408				- T PPPKRLKNCE -							
LOC_Os07g08600.1 11673.m00797/1-525											
ARF1/1-665				- PQQPQRNKPRR -		- P PGL -					- S PATGP -
ARF3/1-608				- SFVT TGPKRSR -		- I GFSSGK PDI -					- V SEGIR -
ARF4/1-788				- IQSSPRPKRPW -		- AGLLDTTPPG -		- NP -			- I TKRGG -
ARF5/1-902				- T SGLKRQLHPS -		- YFAGETEWGSLIKRPLI -		- RVPD -			- SANGIM -
ARF6/1-933				- SPFPLRLKRPW -		- P PGLPSFHGLKEDDMGMS		- SPL -			- M DORGL -
ARF7/1-1165	DDETDMESALKRAMPWL	DNSLEMKDPS		- STIFPGLSLVQ -		- VMNMQQQNG -		- Q -			- L PSAAA -
ARF8/1-811				- SLFPLRLKRPW -		- HAGTSSSLPDG -		- RGDLGSLGTW -			- L RGGGG -
ARF9/1-638				- KSVMLKNKRPR -		- QVSEVSA -					- L DVGIT -
ARF10/1-693				- FSPRKKIRIPQ -		- PFEF -		- P -			- F HGTK -
ARF11/1-601				- TQQQSKRSR -				- P -			- I SEITG -
ARF12/1-593				- RSFL LKNKRLR -				- EV -			- NEIGSS -
ARF13/1-623				- QSSLKKKKH -		- WLQL -					- NEIGAT -
ARF14/1-605				- RSSFLKNKRSR -				- EV -			- NEIGSS -
ARF15/1-598				- RSSFLKNKRLR -				- EV -			- NEFGSS -
ARF16/1-670				- SPPRKKMRLPQ -		- HPDYN -					- L INSIP -
ARF17/1-585				- FPPAKRLKYPQ -		- PGGG -		- F -			- L SGDDG -
ARF18/1-602				- QQQSKCKRSR -				- P -			- I EPSVK -
ARF19/1-1086	DDELDMENAFKRAMPW	MGEDFGMKDAQ		- SSMFPGLSLVQ -		- VMSMQQN -		- NP -			- L SGSAT -
ARF20/1-615				- RSSL LKNKRLR -				- ET -			- L EYLLN -
ARF21/1-606				- RSSL LKNKRLR -				- EV -			- NEFGSS -
ARF22/1-600				- RPSL LKNKRLR -				- EV -			- NEIGSS -
ARF23/1-222											
ARF2/1-859				- PVPMPRPKRPR -		- SNIAPSSPDS -		- SM -			- L TREGT -
PoptrARF8.1/1-827				- SLFPLRLKRPW -		- HPGSPSLLDE -		- ASNGLMW -			- L RGGSG -
PoptrARF1.1/1-660				- SQPMQRNKPRR -		- PSVL P -		- SP -			- T ANLSA -
PoptrARF7.3/1-1113	DDSTDFDSL FKRTMPWL	GDDIYMKD		- PQVL PGLSLAQ -		- RMNMQONPSLANSMQPNYMQSL		- SGSVLQN -			- L PGGLD -
PoptrARF2.3/1-792				- QSKRHRSNMT -				- SS -			- A DSSAP -
PoptrARF5.1/1-933				- PSLT SGLKRPLQSGFL		- GDT EWGGLVKKPLA -					- L L PGSG -
PoptrARF5.2/1-944				- PSLT SGLKRPLQSGFL		- GDSEWGLVKKPLS -		- W -			- L PGSGN -
PoptrARF7.4/1-1137	DDSSDFDSL FKRTMPWL	GDEFCKMD		- PQAL PGLSLVQ -		- VMNMQONPSLANSMQPNYMQSL		- SGSVLQN -			- L PGADL -
PoptrARF16.1/1-669				- SPPRKKLRLPQ -		- PPDF -		- P -			- L L GQIP -
PoptrARF1.2/1-662				- SQPVQRNKRAR -		- PYPV I -		- SP -			- T ADLSA -
PoptrARF9.1/1-666				- QPVQPKNKPRR -				- PP -			- I E I PAF -
PoptrARF2.1/1-854				- PLPMPRPKRPR -		- ANMVPSSPDS -		- SV -			- L TREGS -
PoptrARF16.3/1-700				- SPPRKKLRLPQ -		- HPDF -		- P -			- I DGQFP -
PoptrARF3.1/1-709				- NSMAPGLKRSR -		- SGLPSLKAEF -		- P -			- I PDGIG -
PoptrARF6.2/1-914				- STFPLRLKRPW -		- TPGLHSFHGKLLYTILRAL		- MDGI -			- K DDDL G -
PoptrARF6.4/1-953				- SAFPMRLKRPW -		- PSGLPSFHGLKDDDL		- SINSPMMW -			- L QGGVGD LGVH -
PoptrARF17.1/1-594				- SPPMKKLRYPN -							
PoptrARF2.4/1-879				- SGCR LKRHRPN -		- TVTS -		- SA -			- D SSALT -
PoptrARF16.2/1-708				- SPPRKKLRLPQ -		- PPDF -		- P -			- L L GQIP -
PoptrARF2.2/1-852				- PLPLPRPKRPR -		- ANMVPSSPDS -		- SV -			- L TRDGS -
PoptrARF10.1/1-708				- SPPRKKLRF PQ -		- QLDFP -					- L DGQFQ -
PoptrARF3.3/1-109											
PoptrARF9.2/1-670				- TASTRNSGLTQ -		- SHDL -		- TQ -			- L SVTGE -
PoptrARF6.1/1-884				- SPFPLRLKRPW -		- PPGLPSFHGL -		- GI -			- K DDDL G -
PoptrARF4/1-713				- IQSSPRLKKLR -		- TGLQAAPPDK -		- P -			- I AGGGG -
PoptrARF6.3/1-163											
PoptrARF3.2/1-714				- NFMAPGLKRSR -		- SGLPSSKAEF -		- P -			- I PDGIG -
PoptrARF9.3/1-579				- TQQVMKTKRPR -		- PTDI -					
PoptrARF6.5/1-907				- SAFPMRLKRPW -		- PSGLPSFHGLQDGD		- L N I NSPMMW -			- L QGGVGD LGVQ -
PoptrARF9.4/1-632				- TQQVVKSKRPR -		- SVDI -		- P -			- T SGSIT -
PoptrARF7.1/1-1047	NDESDTENAFKRAVPWL	GDEFGKKDAA		- SSIFPGLSLVQ -		- VMMSQQNNQF -		- QA -			- A QSGFF -
PoptrARF7.2/1-1093	DDSD I ENAFKRAMPWL	GDEF GMDPP		- SSIFPGLSLVQ -		- VMMSQQNNQF -		- PA -			- T QSGLF -
PoptrARF8.2/1-816				- SLFPLRLKRPW -		- HPGSSSLLDS -		- RDEASNGLIW -			- L RGGSG -
PoptrARF16.6/1-91											
PoptrARF16.4/1-701				- SPPRKKLRLPQ -		- HLD F -		- P -			- I DGQFP -
PoptrARF16.5/1-536				- SPPRKKLRLPQ -		- HLD F -		- P -			- I DGQFP -
PoptrARF10.2/1-713				- SPPRKKS RFPQ -		- QLGF -		- P -			- L DLQFQ -
PoptrARF17.2/1-592				- FPPMKKLRYPN -							
PoptrARF2.6/1-724				- MECIADKSLPR -							- F L I SVK -
PoptrARF2.5/1-614				- LPTIAHLLSPP -		- NPEW -		- N -			- F S P I GK -

Consensus



LOC_Os01g70270.1|11667.m07038_altsplice/1-809
LOC_Os01g13520.1|11667.m01333/1-700
LOC_Os02g06910.1|11668.m00642/1-909
LOC_Os02g35140.1|11668.m03331/1-757
LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-819
LOC_Os04g43910.1|11670.m04249/1-696
LOC_Os04g49230.1|11670.m04840/1-294
LOC_Os04g36060.1|11670.m03490/1-1674
LOC_Os05g43920.1|11682.m04181/1-700
LOC_Os06g09660.1|11680.m00929/1-1056
LOC_Os06g46410.1|11680.m04579/1-918
LOC_Os06g47150.1|11680.m04656/1-731
LOC_Os06g48950.1|11680.m04850/1-1116
LOC_Os08g40900.1|11674.m04106_altsplice/1-1056
LOC_Os11g32110.1|11687.m02952/1-812
LOC_Os12g41950.1|11686.m04180/1-900
LOC_Os12g29520.1|11686.m02861_altsplice/1-842
LOC_Os04g56850.1|11670.m05610_altsplice/1-956
LOC_Os05g48870.1|11682.m04683_altsplice/1-696
LOC_Os01g48060.1|11667.m04657_altsplice/1-723
LOC_Os01g54990.1|11667.m05421/1-656
LOC_Os10g33940.1|11676.m02996/1-699
LOC_Os02g04810.1|11668.m00419/1-1094
LOC_Os04g59430.1|11670.m05892/1-530
LOC_Os07g08520.1|11673.m00789/1-729
LOC_Os07g08530.1|11673.m00790/1-408
LOC_Os07g08600.1|11673.m00797/1-525
ARF1/1-665
ARF3/1-608
ARF4/1-788
ARF5/1-902
ARF6/1-933
ARF7/1-1165
ARF8/1-811
ARF9/1-638
ARF10/1-693
ARF11/1-601
ARF12/1-593
ARF13/1-623
ARF14/1-605
ARF15/1-598
ARF16/1-670
ARF17/1-585
ARF18/1-602
ARF19/1-1086
ARF20/1-615
ARF21/1-606
ARF22/1-600
ARF23/1-222
ARF2/1-859
PoptrARF8.1/1-827
PoptrARF1.1/1-660
PoptrARF7.3/1-1113
PoptrARF2.3/1-792
PoptrARF5.1/1-933
PoptrARF5.2/1-944
PoptrARF7.4/1-1137
PoptrARF16.1/1-669
PoptrARF1.2/1-662
PoptrARF9.1/1-666
PoptrARF2.1/1-854
PoptrARF16.3/1-700
PoptrARF3.1/1-709
PoptrARF6.2/1-914
PoptrARF6.4/1-953
PoptrARF17.1/1-594
PoptrARF2.4/1-879
PoptrARF16.2/1-708
PoptrARF2.2/1-852
PoptrARF10.1/1-708
PoptrARF3.3/1-109
PoptrARF9.2/1-670
PoptrARF6.1/1-884
PoptrARF4/1-713
PoptrARF6.3/1-163
PoptrARF3.2/1-714
PoptrARF9.3/1-579
PoptrARF9.4/1-632
PoptrARF7.1/1-1047
PoptrARF7.2/1-1093
PoptrARF8.2/1-816
PoptrARF16.6/1-91
PoptrARF16.4/1-701
PoptrARF16.5/1-536
PoptrARF10.2/1-713
PoptrARF17.2/1-592
PoptrARF2.6/1-724
PoptrARF2.5/1-614

Consensus

SLNFQ-FG-APWI+PS+QL+L+DLSP+FLPSL LSSGQSLQSL+NL S+Q+A+SL SQ++QQASL LQGQEQL G+ - - - - - Q - - - - -

Consensus



	1170	1180	1190	1200	1210	1220	1230	1240	1250	1260	1270
LOC_Os01g70270.1 11667.m07038_altsplice/1-809	SQS			- FGDSPGFFMQN							
LOC_Os01g13520.1 11667.m01333/1-700				- FNSNKGVSPT							
LOC_Os02g06910.1 11668.m00642/1-909	GHSQPEY			- LQQPLQHCQS							- NEQK PQL Q
LOC_Os02g35140.1 11668.m03331/1-757				- PSSPNP I FST S							
LOC_Os02g41800.1 11668.m03990/1-784				- PSLSDLHLTHL							
LOC_Os04g57610.1 11670.m05695/1-819	DQK Q			- KIQPDQSYQVP							T
LOC_Os04g43910.1 11670.m04249/1-696				- SPFPEFHI GNL							
LOC_Os04g49230.1 11670.m04840/1-294											
LOC_Os04g36060.1 11670.m03490/1-1674	EAVRRGGDP	SERWRT	GGVERRRRE	PAT RRT	GSSVG	GFF HAS	SSSL I	AGANGG	SDGGG	GARTL	SWG TWRRL
LOC_Os05g43920.1 11682.m04181/1-700				- PPQRNTEFTY Q							
LOC_Os06g09660.1 11680.m00929/1-1056	VEQHKL	KEQTS	SGGQV	ASQAQMLNQI	- FPPSSSQQL	QQL				GLPKSP	THRQGL
LOC_Os06g46410.1 11680.m04579/1-918	IQGHAQ	SEFL	QQQL	QRCQS	- FTEQKP	QLQT Q					QQQESQ
LOC_Os06g47150.1 11680.m04656/1-731				- ISLSDLHLNKL							
LOC_Os06g48950.1 11680.m04850/1-1116	VLVQNM	QQQH	ASST	QGQP	AT	SOP	LLLP	QQQQ	QQQQ	QQQ	
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056	VQAQL	QQP	QV I	LQAQL	QQP	QVVV	QAQL	QQT	QP	SVQ	SHTVL
LOC_Os11g32110.1 11687.m02952/1-812	PPKD			- VQIPQ	QGFCSL	P					EQIT
LOC_Os12g41950.1 11686.m04180/1-900	TQ			- QLLKE	QELQ	QQ					QR
LOC_Os12g29520.1 11686.m02861_altsplice/1-842	PLKD			- TRNPL	SSFP	S					IS
LOC_Os04g56850.1 11670.m05610_altsplice/1-956	RGKTR	RSKGL	PHKT	VSEK	SDL	SSAP	SWI	CDN	QQV	GLES	K
LOC_Os05g48870.1 11682.m04683_altsplice/1-696				- APLGN	P	GFSY	H				
LOC_Os01g48060.1 11667.m04657_altsplice/1-723	LGVK			- T	PSGN	P	GFSY	H			
LOC_Os01g54990.1 11667.m05421/1-656				- LPVQY	SGYT	H	Q				
LOC_Os10g33940.1 11676.m02996/1-699				- LPLT	D	HQ	L	NKL			
LOC_Os02g04810.1 11668.m00419/1-1094	VQNQM	QQK	PSPT	QNPQR	I	NGR	QL	LL	SHQ	K	QNL
LOC_Os04g59430.1 11670.m05892/1-530				- NNIS	A	P	V	C	N	G	D
LOC_Os07g08520.1 11673.m00789/1-729				- FV	I	P	K	P	S	A	A
LOC_Os07g08530.1 11673.m00790/1-408											
LOC_Os07g08600.1 11673.m00797/1-525				- F	S	P	A	A	A	T	F
ARF1/1-665											
ARF3/1-608											
ARF4/1-788				- E	F	V	D	A	T	G	V
ARF5/1-902				- QHGL	E	Q	S	M	E	Q	A
ARF6/1-933	QQQL	S	Q	Q	Q	Q	Q	A	Y	L	G
ARF7/1-1165	QQHS	NNNQ	S	Q	Q	Q	Q	Q	L	L	Q
ARF8/1-811				- E	N	L	P	Q	N	M	R
ARF9/1-638				- W	S	S	P	Q	C	H	R
ARF10/1-693				- F	G	S	P	S	P	L	S
ARF11/1-601				- S	P	V	A	S	S	F	L
ARF12/1-593				- A	S	P	M	N	I	S	L
ARF13/1-623				- S	P	I	S	V	P	E	S
ARF14/1-605				- A	T	P	M	N	I	S	L
ARF15/1-598				- A	S	P	M	N	I	S	L
ARF16/1-670				- N	A	H	Q	Y	Y	G	L
ARF17/1-585				- A	P	D	P	S	P	S	M
ARF18/1-602				- K	L	F	Q	D	P	S	L
ARF19/1-1086	GT	I	A	V	A	N	Q	V	S	C	S
ARF20/1-615				- A	S	P	M	N	T	S	L
ARF21/1-606				- A	S	P	M	N	I	S	L
ARF22/1-600				- A	S	P	M	N	I	S	L
ARF23/1-222											
ARF2/1-859	F	G	T	N	I						
PoptrARF8.1/1-827	Q	I	S	T							
PoptrARF1.1/1-660				- E	S	L	P	R	H	L	Q
PoptrARF7.3/1-1113	Q	N	Q	Q	H	I	M	G	N	Q	Q
PoptrARF2.3/1-792				- S	K	L	N	E	T	G	M
PoptrARF5.1/1-933	V	D	G	K							
PoptrARF5.2/1-944	G	E	R	K	L	K	A	E	T	E	H
PoptrARF7.4/1-1137	Q	Q	N	Q	Q	H	I	M	G	N	Q
PoptrARF16.1/1-669				- L	S	S	S	D	L	H	F
PoptrARF1.2/1-662				- D	L	Y	S	P	N	F	S
PoptrARF9.1/1-666				- L	S	V	T	G	G	W	L
PoptrARF2.1/1-854	F	G	A								
PoptrARF16.3/1-700				- L	P	L	S	D	L	N	L
PoptrARF3.1/1-709				- I	P	T	S	E	N	S	F
PoptrARF6.2/1-914	E	N	K	H	Q	S	Q	P	S	Q	T
PoptrARF6.4/1-953	H	Q	H	Q	H	Q	V	Q			
PoptrARF17.1/1-594				- S	D	L	T	N	S	R	T
PoptrARF2.4/1-879				- T	S	D	E	K	S	M	F
PoptrARF16.2/1-708				- L	S	S	S	D	L	H	F
PoptrARF2.2/1-852	F	G	T								
PoptrARF10.1/1-708				- I	S	L	S	D	F	Q	F
PoptrARF3.3/1-109											
PoptrARF9.2/1-670				- T	Q	T	E	G	G	W	A
PoptrARF6.1/1-884	I	Q	Q	L	L	H	H	L	D	S	P
PoptrARF4/1-713				- L	S	P	V	Y	R	C	E
PoptrARF6.3/1-163											
PoptrARF3.2/1-714				- I	A	T	S	N	N	S	Y
PoptrARF9.3/1-579											
PoptrARF6.5/1-907	Q	P	K	Q	L						
PoptrARF9.4/1-632				- Q	K	E	I	D	S	N	L
PoptrARF7.1/1-1047	Q	Q	Q	Q	Q	Q	L	L	A	S	N
PoptrARF7.2/1-1093	P	P	T	L	N	D	S	V	Q	P	V
PoptrARF8.2/1-816	Q	I	S	S							
PoptrARF16.6/1-91				- L	P	L	S	D	P	H	L
PoptrARF16.4/1-701				- L	P	L	S	D	P	H	L
PoptrARF16.5/1-536				- I	S	L	S	D	I	Q	F
PoptrARF10.2/1-713				- S	D	L	T	N	S	T	M
PoptrARF17.2/1-592				- Y	P	C	P	S	N	T	V
PoptrARF2.6/1-724				- L	G	N	P	S	W	C	S
PoptrARF2.5/1-614											

Consensus



LOC_Os01g70270.1|11667.m07038. altsplice/1-809
LOC_Os01g13520.1|11667.m01333/1-700
LOC_Os02g06910.1|11668.m00642/1-909
LOC_Os02g35140.1|11668.m03331/1-757
LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-189
LOC_Os04g46410.1|11670.m04249/1-696
LOC_Os04g49230.1|11670.m04840/1-294
LOC_Os04g36060.1|11670.m03490/1-1674
LOC_Os05g43920.1|11682.m04181/1-700
LOC_Os06g09660.1|11680.m00929/1-1056
LOC_Os06g46410.1|11680.m04579/1-918
LOC_Os06g47150.1|11680.m04656/1-731
LOC_Os06g48950.1|11680.m04850/1-1116
LOC_Os08g40900.1|11674.m04106. altsplice/1-1056
LOC_Os11g32110.1|11687.m02952/1-812
LOC_Os12g41950.1|11686.m04180/1-900
LOC_Os12g29520.1|11686.m02861. altsplice/1-842
LOC_Os04g56850.1|11670.m05610. altsplice/1-956
LOC_Os05g48870.1|11682.m04683. altsplice/1-696
LOC_Os01g48060.1|11667.m04657. altsplice/1-723
LOC_Os01g54990.1|11667.m05421/1-656
LOC_Os10g33940.1|11676.m02996/1-699
LOC_Os02g04810.1|11668.m00419/1-1094
LOC_Os04g59430.1|11670.m05892/1-530
LOC_Os07g08520.1|11673.m00789/1-729
LOC_Os07g08530.1|11673.m00790/1-408
LOC_Os07g08600.1|11673.m00797/1-525
ARF1/1-665
ARF3/1-608
ARF4/1-788
ARF5/1-902
ARF6/1-933
ARF7/1-1165
ARF8/1-811
ARF9/1-638
ARF10/1-693
ARF11/1-601
ARF12/1-593
ARF13/1-623
ARF14/1-605
ARF15/1-598
ARF16/1-670
ARF17/1-585
ARF18/1-602
ARF19/1-1086
ARF20/1-615
ARF21/1-606
ARF22/1-600
ARF23/1-222
ARF2/1-859
PoptrARF8.1/1-827
PoptrARF1.1/1-660
PoptrARF7.3/1-1113
PoptrARF2.3/1-792
PoptrARF5.1/1-933
PoptrARF5.2/1-944
PoptrARF6.1/1-1137
PoptrARF16.1/1-669
PoptrARF1.2/1-662
PoptrARF9.1/1-666
PoptrARF2.1-854
PoptrARF16.3/1-700
PoptrARF3.1/1-709
PoptrARF6.2/1-914
PoptrARF6.4/1-953
PoptrARF17.1/1-594
PoptrARF2.4/1-879
PoptrARF16.2/1-708
PoptrARF2.2/1-852
PoptrARF10.1/1-708
PoptrARF3.3/1-109
PoptrARF9.2/1-670
PoptrARF16.1/1-884
PoptrARF4/1-713
PoptrARF6.3/1-163
PoptrARF3.2/1-714
PoptrARF9.3/1-579
PoptrARF6.5/1-907
PoptrARF9.4/1-632
PoptrARF7.1/1-1047
PoptrARF7.2/1-1093
PoptrARF8.2/1-816
PoptrARF16.6/1-91
PoptrARF16.4/1-701
PoptrARF16.5/1-536
PoptrARF10.2/1-713
PoptrARF17.2/1-592
PoptrARF2.6/1-724
PoptrARF2.5/1-614

FDEAPNRLT S FKNQF QDQGS ARHFS DP YYY VSP QP S L T VESS T QMHT DSKEL HF WNGQST VY GNSRDRP QNFRFE QN SSSW L NQSF ARPE QPRV I R
----- LSE I SOK I FQVTS NEAR I ATWP ARSAY QAEPT - - - - - SKL SSNTAACGV - - - - - SKL SSNTAACGV -
PQQQQQESHQQPQHQQMQQKHL SNFQT VPNAL SVFSQL SSTPQST PSTL QT VSPFSQQHNF PDTN I SCL SPSNVSSMHD L RSFPSEAASDL PG
----- L NVGFS T KNEP SAL SNKHF YWPMRETR - - - - - ANSYSAS I SKVPS - - - - - ANSYSAS I SKVPS -
----- QSSL MYPGLRRPDHVGPTS I PPR I STL TM - - - - - GSSPPARALS MGA - - - - - GSSPPARALS MGA -
SAVLPSP TSLPSHLREKFGSDPNANSSSF ITSSSSDNMLDSSFL QGS - - - - - SKAVDL SRFNQPV - - - - - SKAVDL SRFNQPV -
----- QPNL ML YAG I RLPPADRAAPAPRPRI I STLDT LG - - - - - SPKGPDIDAACSPS - - - - - SPKGPDIDAACSPS -
GEGNRRRST LVL PHVRSAPFHAVVP GAVGGQRRQ I VAAAVAE ADGNENGKNK HFTCHRENANYGLKSASENK I ELLRGAHSWADSN I EDVLASVNN - - - - - DV
----- P I GFSESL GFPEVL QGQEMS QV VPLFRGATFGARTQN - - - - - DRVVSANSVHRSA - - - - - DRVVSANSVHRSA -
QPTLTQT S QVQQA A EYQALL QSQQQQQQL QLQQL SQPEVQL QL L QK I QQQNML SQLNPQHQS QL I QQL SQKSQ E I LQQQ I LQHQFGGSDS I GQLK
QQQQQQSQCMQVPQHQQMQQKNMTNY QSVPNAL SPFSQL SSPSQSSPMTL QT VLPFSQP QSYPTD SMSSL SPSTNTST MHNAL RPFSS EAPSHLSM
----- QSSL SPHGLHQLDHGMQPRIAAGL IIG - - - - - HPAARDD I SC L L T - - - - - HPAARDD I SC L L T -
QQL SKMPAQL SSLANQQFQLTDQQL QLQL KQL QQQQSSQL SQPAVTLAQLPL I QEQQK L L LDMQQQL SNSOTL SQQQMMPQSTKVPSONT PLPL
PQQL HHQQQQT QQL QPVQVQVQSVQEHHQ I K I QPVHVSMDSAMNT QVADHQMKL QL LKAL QPQQPL I SEQQKML L DLQQV I NSQSA PQQCQVQTN - - - - - QA I SLHNS
AGHSNF VHTVNAQY QDQGS NHHMFPSSWSF MPNT RLGL NKQNYSM I QEAGVL SQRPGNT KFGNGVYAAL PGRGT EYQSGGW GHHMPPNSHMDDT Q
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GNRSNTWNT I NVHYPDQNAHNHMYPGTWL MPNTGFGVGNQNYL MTPDITL PQRSL NAKFGGNGAFTSLRAHGI DQRSSGWH G IEPSSH I DDS
SSKSPDGGKSVNSFPNQGCF SQ I DGLDMMTPQSY YQDSNV I QPAGVSEN I FSSSAD I PPSMI ADT METFQASCL SDCLPNS I QEF I SSPDL NSLT - - - - - F
----- CS GF GESHRLQKVL QGOEL FRPYRGT LVDASMG S - - - - - NGFHQQDSPRAPG - - - - - NGFHQQDSPRAPG -
----- CS GF GESQRFQEV LQGQEVFRPYRGGL SDAC I RG - - - - - SGFRQPDGNHAG - - - - - SGFRQPDGNHAG -
----- SVSFGES I GFPEVL QQGE I SQTVPFFQGM L P D ACSAK - - - - - SRYELKNYVCTPA - - - - - SRYELKNYVCTPA -
----- HLGL L HSGFSNRLDA I T PPSRI SKG FVVS - - - - - SAPAHON I SC L L S - - - - - SAPAHON I SC L L S -
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----- SLLV PPI L MHPQPOPPAD I QGARHNNG - - - - - HAYADI P SSS T P S - - - - - HAYADI P SSS T P S -
----- VLPR I PDL RVTNL QGGRW EFGHT WSDADT DRRSSSHTL AAGW - - - - - SAFVKAKRL CVGD - - - - - SAFVKAKRL CVGD -

----- HGGNKSFGVSI GSAFWPTNADSAAESFA - - - - - SAFNNESTEKKQT - - - - - SAFNNESTEKKQT -
----- GTEFGDSYGFHKVL QGOETVPAYS I T DHRQQ - - - - - HGLSQRN I WCGFP - - - - - HGLSQRN I WCGFP -
----- CSGVMDLDRFP RVL QQGE I CSLKSFQFAGSPAAAPNPFAYQANK - - - - - SSYPPLALHGIRS - - - - - SSYPPLALHGIRS -
----- SQVTTSTVCNEEKVNQL KQPGASSPVQADQCLDI THQ I YQPSDPI NGSFLE TDELT SQVSSSQSLAGSYKQPF I LSSQDSSAVVLPD
SQPNTSPL QSMTSLCHQQFSVSDTNGGNNP I SPLHTLLNSFSQDESSQLHL TRTNSAMTSSGWPSKRPADVSSFHQSAGAGNNNT QSVLQL QGSHT
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----- NEDAKSDMLNNSYSVSNVAKDSTLN - - - - - DQMSVPEQK KPE - - - - - DQMSVPEQK KPE -
----- LNLSSYTGNK L HSPAMFLSSFNPRHHYQAR - - - - - DSENSNN I SC L L T - - - - - DSENSNN I SC L L T -
----- SQSHESNP SVKLLF QDPATERNSN - - - - - KS VFS SGLQCK I T - - - - - KS VFS SGLQCK I T -
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----- NA I EDSKFLSGLLLNHSLLA I P NENYNS - - - - - DQMI QPRKEDI TT - - - - - DQMI QPRKEDI TT -
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----- ETTEDAMNPSRLMSYPVQPM PKRNYN - - - - - NQMVQ I EEN I TT - - - - - NQMVQ I EEN I TT -
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----- STFPAQMGGARQYDFGSEFN - - - - - TGF I GGNPPQLFT - - - - - TGF I GGNPPQLFT -
----- SGGYSSNNSKFKETPPP - - - - -
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----- GFNESYRFHKVL QQGE I PPRSPYRR I PNANEAREN - - - - - CGLGLSDGQRSS - - - - - CGLGLSDGQRSS -
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----- NASMLNYS TFPAGMQGARQDPSTFSL - - - - - SNL I SENASQVFG - - - - - SNL I SENASQVFG -
----- SAPMMYTGHFPMNLASSMEVQVAKDKDTQQHGSRF - - - - - LFPFYADSSPHPS - - - - - LFPFYADSSPHPS -
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----- LQSGFLSSLQRFNPRTKNSENYPTG - - - - - HPDSNKNLSC L L T - - - - - HPDSNKNLSC L L T -

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----- QTGLLRTGF P L L DHTASLTKASNVT I P - - - - - KPSMCE DVSCELT - - - - - KPSMCE DVSCELT -
----- QQSGMFLSSLQRFNPHSRNSETYLTG - - - - - HTNSNEN I SC L L T - - - - - HTNSNEN I SC L L T -
----- NASMLNYS TFPAGMQGARQDPSTFGL - - - - - SNF I SENAPQVFS - - - - - SNF I SENAPQVFS -
NIARLG I P NSRHSTFSYGVHDNAVGSRLSVPNVSHNSGSQKWG - - - - - SELKHANEVPLAA - - - - - SELKHANEVPLAA -
----- SSGVHDNAVGSRLSVPFVSHNSGSQKW - - - - - RGELKHANEVPL - - - - - RGELKHANEVPL -

Consensus



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-----EKKQEPSSAGCRL FGI E I SSAVEAT SPLAAVSGVGQDQPAASV
-----KKPDDAKPPGLML FGQR I L TERQMSL SGTTSPAATGNSSL NWN
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-----MVRTQQL FPRDLQ I LVPHTDI VTPQNGSPDPNPVNTPLASADGMKT
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-----NGNV CRL FGFEL VENVN VDECF SAASVS GAVAVDQPV
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Consensus



Consensus

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Consensus



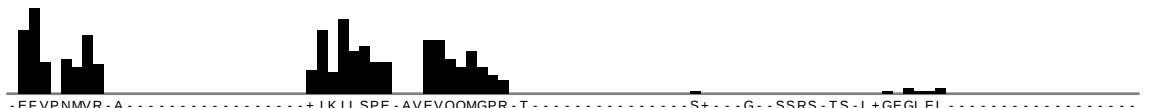
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LOC_Os02g41800.1|11668.m03990/1-784
LOC_Os04g57610.1|11670.m05695/1-189
LOC_Os04g43910.1|11670.m04249/1-696
LOC_Os04g49230.1|11670.m04840/1-294
LOC_Os04g36060.1|11670.m03490/1-1674
LOC_Os05g43920.1|11682.m04181/1-700
LOC_Os06g09660.1|11680.m00929/1-1056
LOC_Os06g46410.1|11680.m04579/1-918
LOC_Os06g47150.1|11680.m04656/1-731
LOC_Os06g48950.1|11680.m04850/1-1116
LOC_Os08g40900.1|11674.m04106_altsplice/1-1056
LOC_Os11g32110.1|11687.m02952/1-812
LOC_Os12g41950.1|11686.m04180/1-900
LOC_Os12g29520.1|11686.m02861_altsplice/1-842
LOC_Os04g56850.1|11670.m05610_altsplice/1-956
LOC_Os05g48870.1|11682.m04683_altsplice/1-696
LOC_Os01g48060.1|11667.m04657_altsplice/1-723
LOC_Os01g54990.1|11667.m05421/1-656
LOC_Os10g33940.1|11676.m02996/1-699
LOC_Os02g04810.1|11668.m00419/1-1094
LOC_Os04g59430.1|11670.m05892/1-530
LOC_Os07g08520.1|11673.m00789/1-729
LOC_Os07g08530.1|11673.m00790/1-408
LOC_Os07g08600.1|11673.m00797/1-525
TFVHKQGVAVGRS
IKVQMHGNAVGRA
VKVYKSG-TVGRLL
TKVIMQGMVAVGRA
CKVFIESDTVGRN
VKVYKSG-SVGRS
CKVFVESETVGRS
AESYLTGSAEADTKPREVSVAUVSSIMDIPSAWDACAVDSVDPDNFNPFLT
SEITSNQFALARI
TKVQKRG-SVGRS
VKVYKSG-TYGRS
CKVFMQSEDVGR
TKVYKRG-AVGRS
TKVHKRG-AVGRS
KKVHKQGIALGRS
VKVYKSG-SLGRS
KKVHKQGIALGRS
TKVQKQG-SVGRS
GEVIDGQVTVEKS
TRNEKLHSPVSGA
SETSDWQFPIKGA
CKVFMESEDVGR
TKVYKRG-AVGRS
GDTNGAF
VRLLMNPDDARRR
IDDTASTSVS
TKVHKQGSVAVGRA
DHHGSGSRC
TKVHKQGSQVGRA
TKVQKTG-SVGRS
VKVYKSG-SFGRS
TKVQKRG-SVGRS
VKVYKSG-SVGRS
TKVQMQGVPVGRA
CKVFMESEDVGR
IKVQMQGTAVGRA
TKVQMQGVTIGRA
IKVHKQGVAVGRA
TKVQMQGVTIGRA
TKVQMQGVTIGRA
TKVQMQGVTIGRA
CKVFMESEDVGR
NSIQLFGKI
TKVQMQGTAVGRA
TKVQKRG-SVGRS
TKVQMQGVTIGRA
TKVQMQGVTIGRA
TKVQMQGVTIGRA
TKVQMQGVTIGRA
TKVHKQGIALGRS
VKVYKSG-SVGRS
TKVHKQGVAVGRA
TKVYKRG-AVGRS
VKVYKQGTAVGRS
TKVQKTG-SVGRS
TKVYKRG-AVGRS
CKVFMESEDVGR
TKVHKQGVAVGRA
TKVQKQIAGVRA
TKVHKQGIALGRS
CKVFLESEDVGR
SHEHLQHGISQPV
VKVYKSG-SFGRS
VKVHKSG-SYGRS
QSSAHSFGK
VKVHKQGTAVGRS
CKVFLESEDVGR
TKVHKQGIALGRS
KAFLESEDLGRT
TKVQMQGTAVGRA
VKVYKSG-SFGRS
TKVHKQGSVAVGRA
VKVYKSG-SFGRS
SHEHLQHGVSQPI
AMVQMKGIAVGRA
VKVHKSG-SYGRS
TKVQMQGVAVGRA
TKVQKRG-SVGRS
TKVQKRG-SVGRS
VKVYKSG-SVGRS
CKVFMESEDVGR
CKVFMDSEDVGR
CKVFLESEDVGR
QSSAYSFG
TKVLKYGTVPGRS
TKVLKYGTVLGRS

Consensus



	1810	1820	1830	1840	1850	1860	1870	1880	1890	1900	
LOC_Os01g70270.1 11667.m07038_altsplice/1-809	-EFC-SIVR		-K I Y I Y T K E	-E V Q K M N S K							
LOC_Os01g13520.1 11667.m01333/1-700	-EFC-QMVR		-K I V L Y P I E	-D E K K I E P H							
LOC_Os02g06910.1 11668.m00642/1-909	-EFV-NSVS		-C I K I L S P Q	-E V Q Q M G K P							
LOC_Os02g35140.1 11668.m03331/1-757	W L T T - T L L Q		-C L I L L S R V	-L V N A A L L I L Y I Q R A V L F N G L C F Q I H W I N A G G F G G W K E G Y S T K R A S G V C R E P							
LOC_Os02g41800.1 11668.m03990/1-784	R D F N - K K A Q S A Q V S Y C S P V S F F I I I N N D Q L R V L V D S		-R V L R C K N P V T T H Q R L E C R G P L A Q G T S A C S M Q T Q T Q R C A I S D L S G G L T S R								
LOC_Os04g57610.1 11670.m05695/1-819	-S F V - N S V W		-Y I K I L S P E	-D V H K M G K Q							
LOC_Os04g43910.1 11670.m04249/1-696	-A F V - K S A R		-K L R I L T D A	-G S D N L G D X							
LOC_Os04g49230.1 11670.m04840/1-294	-S T L		-R C T T R A P X								
LOC_Os04g36060.1 11670.m03490/1-1674	G E F V F D Q S W A E A Y Y S Y G L E Y Y P		-K L Q S C V P F T P V T G Q R I L V R D T P Y R D Q V F E A L V K E L K S L T T K L K V S S L H I T F P S E G E F S K L K D S E F L Q R I G M Q Y H W R N R								
LOC_Os05g43920.1 11682.m04181/1-700	-L F G H S H S T		-P G A L H A L C	-A A A P L G M X							
LOC_Os06g09660.1 11680.m00929/1-1056	-E F V - G C V K		-S I K I L S A A	-E V Q Q M S L D							
LOC_Os06g46410.1 11680.m04579/1-918	-E F V - N S V S		-C I K I L S P Q	-E V Q Q M G K P				-F E			
LOC_Os06g47150.1 11680.m04656/1-731	-E F T - K T A R		-R L N I L T D T	-S G D N L A S W							
LOC_Os06g48950.1 11680.m04850/1-1116	-E F V - G C V K		-C I R I L S P Q	-E V Q Q M S L E							
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056	-D F V - K C V R		-C I R I L S P Q	-E E M Q M R L V							
LOC_Os11g32110.1 11687.m02952/1-812	-F A - P V Q P		-D G T G Y L S C	-Y L A T X							
LOC_Os12g41950.1 11686.m04180/1-900	-E F A - N S V W		-C I K I L S P Q	-E V Q Q L V R G					G D G L		
LOC_Os12g29520.1 11686.m02861_altsplice/1-842	-E F C - N M V H		-K I F I Y T R E	-E V Q K M N P G					-A L		
LOC_Os04g56850.1 11670.m05610_altsplice/1-956	-E F I - N C V R		-C I R I L S P S	-E V Q Q M S E N							
LOC_Os05g48870.1 11682.m04683_altsplice/1-696	-K S L G N S C A		-T V S V L X								
LOC_Os01g48060.1 11667.m04657_altsplice/1-723	-Y H S		-R L Q S L K P Q	-M P K S L G S S							
LOC_Os01g54990.1 11667.m05421/1-656	-L F G Y N C S T		-P G A L H A L C	-A A P L G I X							
LOC_Os10g33940.1 11676.m02996/1-699	-D F M - K A A R		-R L T I I A G D	-R E R I E R P							
LOC_Os02g04810.1 11668.m00419/1-1094											
LOC_Os04g59430.1 11670.m05892/1-530	-D -		-S L T N G H N Q	-D G A R L X							
LOC_Os07g08520.1 11673.m00789/1-729	-E A A A A T T S T A H A G		-D A T T S A P S L A L Q L Q T M A S S								
LOC_Os07g08530.1 11673.m00790/1-408			-R L E A L I P D	-N I X							
LOC_Os07g08600.1 11673.m00797/1-525	-E V I		-S I E N Y A T S	-I G A X							
ARF1/1-665	-E F C - G M V R		-K I F I Y T P E	-E V K K L S P K							
ARF3/1-608			-C V E G N S M K G A S A V Q S N H H H								
ARF4/1-788	-D F C - N V V W		-K I H L Y T K E	-E V E N A N D D							
ARF5/1-902	-E F V - G C V R		-C I R I L S P T	-E V Q Q M S E E							
ARF6/1-933	-E F V - S S V W		-C I K I L S P Q	-E V Q Q M G K R				-G L E L L N S A P			
ARF7/1-1165	-E F V - N C V Q		-S I K I L S S A	-E V Q Q M S L D							
ARF8/1-811	-S F V - N N V W		-Y I K I L S P E	-D V H Q M G D H							
ARF9/1-638	-E F C - N M V K		-R I F I W S K E	-E V K K M T P G							
ARF10/1-693	-D F M - K A T K		-R L T I K M D I	-G G D N V R K T							
ARF11/1-601	-E F C - K M A K		-K L F I Y P S D	-E V K K M R S K							
ARF12/1-593	-E F C - N M V K		-K I F I Q K R R								
ARF13/1-623	-E F C - N M A K		-R I F I C S K E	-E I K K M K L K							
ARF14/1-605	-E F C - N M V K		-K I F I Y S K E	-E V K N L K S R							
ARF15/1-598	-E F C - N M V K		-R I Y I Q K R R								
ARF16/1-670	-E F L - K T A R		-R L T I L T E Q	-G S E S V V V							
ARF17/1-585			-S V P K H S N S	-N A G S S S Q G							
ARF18/1-602	-E F C - K M A K		-K I F I Y S S D	-E V K K M T T K							
ARF19/1-1086	-E F V - N C V Q		-N I K I L S S V	-E V Q Q M S L D							
ARF20/1-615	-E F C - K M V K		-K I L I Y S K E	-E V K N L K S S							
ARF21/1-606	-E F C - K M V K		-K I L I Y S K E	-E V K N L K S S							
ARF22/1-600	-E F C - N M V K		-K I L I F K R G	-G Q K L E V Q							
ARF23/1-222											
ARF2/1-859	-E F C - C M V R		-K I F I Y T K E	-E V R K M N P G							
PoptrARF8.1/1-827	-L F V - N N V W		-Y I K I L S P E	-D V L K L G E Q							
PoptrARF1.1/1-660	-E F C - S M V K		-K I F I Y A S E	-E V K R L S P K							
PoptrARF7.3/1-1113	-E F V - N C V R		-C I K I L S P Q	-E V Q Q M S L D							
PoptrARF2.3/1-792	-E F C - S M V R		-R I F V F T R E	-E I N R M E P R				-S L N			
PoptrARF5.1/1-933	-E F V - G C V R		-C I R I L S P S	-E V Q Q M S E E							
PoptrARF5.2/1-944	-E F V - G C V R		-C I R I L S P S	-E V Q Q M S E E							
PoptrARF7.4/1-1137	-E F V - N C V R		-C I K I L S P Q	-E V Q Q M S L D							
PoptrARF16.1/1-669	-E F L - K T A R		-R L T I L S D A	-S S D N V G R							
PoptrARF1.2/1-662	-E F C - G M V K		-K I F I Y T S E	-E V K R L S P K							
PoptrARF9.1/1-666	-E F C - N M V R		-R I F I C S S Q	-D V K K M S P G							
PoptrARF2.1/1-854	-E F V - G M V R		-K I V I Y T R E	-E V Q R I K P G							
PoptrARF16.3/1-700	-N F S - K T A R		-R L T I V T R F	-R Q							
PoptrARF3.1/1-709	-S F L P R V G E		-Q L H P K P P A	-I N N A V G S S							
PoptrARF6.2/1-914	-E F V - N S V W		-C I K I L S P Q	-E V Q Q M G K R					G L E L		
PoptrARF6.4/1-953	-E F V - N N V W		-Y I K I L S P L	-E V Q Q M G K E				G L T S A A S V P S Q K L S N S T S D			
PoptrARF17.1/1-594	-L E L		-S L T S S Y T E	-L L N R I D V Q							
PoptrARF2.4/1-879	-E F C - S M V R		-R I F I Y T K E	-E I N R M E P R				-S L N L E A E G N S R S T D Q M V D L E N R I S			
PoptrARF16.2/1-708	-E F L - K T A R		-R L T I L S Y A	-S R D N F G R							
PoptrARF2.2/1-852	-E F V - G M V R		-K I V I Y T K E	-E A Q K I K P G							
PoptrARF10.1/1-708	-V F M - K T A K		-R L T I L M N R	-A S G N S V G R R							
PoptrARF3.3/1-109											
PoptrARF9.2/1-670	-E F C - N M V R		-R I Y I C S S Q	-D V K R M G P G							
PoptrARF6.1/1-884	-E F V - N S V W		-Y I K I L S P Q	-E V Q Q M G K R					G L E L		
PoptrARF4/1-713	-E F C - N V A T		-K I H I Y T Q E	-E V E K M T L F							
PoptrARF6.3/1-163	-E F V - N S V W		-C I K I L S P Q	-E V Q Q M G K R					G L E L		
PoptrARF3.2/1-714	-S F L P R A G E		-H F H Q S L Q Q								
PoptrARF9.3/1-579	-E F C - K M V K		-K I F I Y S S E	-E V K K T G T R							
PoptrARF6.5/1-907	-E F V - N N V W		-Y I K I L S P L	-E V Q Q M G K E					G L S P		
PoptrARF9.4/1-632	-E F C - K M V K		-K I F I Y S S E	-E V K K M S T R							
PoptrARF7.1/1-1047	-E F M - S C V Q		-S I K I L S S A	-E V Q Q M S L D							
PoptrARF7.2/1-1093	-E F V - S C V Q		-S I K I L S S A	-E V Q Q M S L D							
PoptrARF8.2/1-816	-L F V - N N V W		-Y I K I L S P E	-D V L K M G E Q							
PoptrARF16.6/1-91											
PoptrARF16.4/1-701	-D F F - K T A R		-R L T I V T D S	-S S G N V G I							
PoptrARF16.5/1-536											
PoptrARF10.2/1-713	-V F M - K T A K		-R L T I L M N Q P S T E I D D L S C R								
PoptrARF17.2/1-592	-L S L T S S Y T		-E L L N R I D A	-Q C Q R A S P V							
PoptrARF2.6/1-724	-E F Q - L T V R		-R I F I S P K E	-D I G K L N P L							
PoptrARF2.5/1-614	-E F L - S A V H		-R I F I C P K E	-E T G K L N P							

Consensus



	1910	1920	1930	1940	1950	1960	1970	1980	1990	2000	2010
LOC_Os01g70270.1 11667.m07038_altsplice/1-809											
LOC_Os01g13520.1 11667.m01333/1-700											
LOC_Os02g06910.1 11668.m00642/1-909											
LOC_Os02g35140.1 11668.m03331/1-757											
LOC_Os02g41800.1 11668.m03990/1-784											
LOC_Os04g57610.1 11670.m05695/1-819											
LOC_Os04g43910.1 11670.m04249/1-696											
LOC_Os04g49230.1 11670.m04840/1-294											
LOC_Os04g36060.1 11670.m03490/1-1674											
LOC_Os05g43920.1 11682.m04181/1-700											
LOC_Os06g09660.1 11680.m00929/1-1056											
LOC_Os06g46410.1 11680.m04579/1-918											
LOC_Os06g47150.1 11680.m04656/1-731											
LOC_Os06g48950.1 11680.m04850/1-1116											
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056											
LOC_Os11g32110.1 11687.m02952/1-812											
LOC_Os12g41950.1 11686.m04180/1-900											
LOC_Os12g29520.1 11686.m02861_altsplice/1-842											
LOC_Os04g56850.1 11670.m05610_altsplice/1-956											
LOC_Os05g48870.1 11682.m04683_altsplice/1-696											
LOC_Os01g48060.1 11667.m04657_altsplice/1-723											
LOC_Os01g54990.1 11667.m05421/1-656											
LOC_Os10g33940.1 11676.m02996/1-699											
LOC_Os02g04810.1 11668.m00419/1-1094											
LOC_Os04g59430.1 11670.m05892/1-530											
LOC_Os07g08520.1 11673.m00789/1-729											
LOC_Os07g08530.1 11673.m00790/1-408											
LOC_Os07g08600.1 11673.m00797/1-525											
ARF1/1-665											
ARF3/1-608											
ARF4/1-788											
ARF5/1-902											
ARF6/1-933											
ARF7/1-1165											
ARF8/1-811											
ARF9/1-638											
ARF10/1-693											
ARF11/1-601											
ARF12/1-593											
ARF13/1-623											
ARF14/1-605											
ARF15/1-598											
ARF16/1-670											
ARF17/1-585											
ARF18/1-602											
ARF19/1-1086											
ARF20/1-615											
ARF21/1-606											
ARF22/1-600											
ARF23/1-222											
ARF2/1-859											
PoptrARF8.1/1-827											
PoptrARF1.1/1-660											
PoptrARF7.3/1-1113											
PoptrARF2.3/1-792											
PoptrARF5.1/1-933											
PoptrARF5.2/1-944											
PoptrARF7.4/1-1137											
PoptrARF16.1/1-669											
PoptrARF1.2/1-662											
PoptrARF9.1/1-666											
PoptrARF2.1/1-854											
PoptrARF16.3/1-700											
PoptrARF3.1/1-709											
PoptrARF6.2/1-914											
PoptrARF6.4/1-953											
PoptrARF17.1/1-594											
PoptrARF2.4/1-879											
PoptrARF16.2/1-708											
PoptrARF2.2/1-852											
PoptrARF10.1/1-708											
PoptrARF3.3/1-109											
PoptrARF9.2/1-670											
PoptrARF6.1/1-884											
PoptrARF4/1-713											
PoptrARF6.3/1-163											
PoptrARF3.2/1-714											
PoptrARF9.3/1-579											
PoptrARF6.5/1-907											
PoptrARF9.4/1-632											
PoptrARF7.1/1-1047											
PoptrARF7.2/1-1093											
PoptrARF8.2/1-816											
PoptrARF16.6/1-91											
PoptrARF16.4/1-701											
PoptrARF16.5/1-536											
PoptrARF10.2/1-713											
PoptrARF17.2/1-592											
PoptrARF2.6/1-724											
PoptrARF2.5/1-614											

Consensus

	2020	2030	2040	2050	2060	2070	2080	2090	2100	2110	
LOC_Os01g70270.1 11667.m07038_altsplice/1-809											
LOC_Os01g13520.1 11667.m01333/1-700											
LOC_Os02g06910.1 11668.m00642/1-909											
LOC_Os02g35140.1 11668.m03331/1-757											
LOC_Os02g41800.1 11668.m03990/1-784											
LOC_Os04g57610.1 11670.m05695/1-819											
LOC_Os04g43910.1 11670.m04249/1-696											
LOC_Os04g49230.1 11670.m04840/1-294											
LOC_Os04g36060.1 11670.m03490/1-1674											
LOC_Os05g43920.1 11682.m04181/1-700											
LOC_Os06g09660.1 11680.m00929/1-1056											
LOC_Os06g46410.1 11680.m04579/1-918											
LOC_Os06g47150.1 11680.m04656/1-731											
LOC_Os06g48950.1 11680.m04850/1-1116											
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056											
LOC_Os11g32110.1 11687.m02952/1-812											
LOC_Os12g41950.1 11686.m04180/1-900											
LOC_Os12g29520.1 11686.m02861_altsplice/1-842											
LOC_Os04g56850.1 11670.m05610_altsplice/1-956											
LOC_Os05g48870.1 11682.m04683_altsplice/1-696											
LOC_Os01g48060.1 11667.m04657_altsplice/1-723											
LOC_Os01g54990.1 11667.m05421/1-656											
LOC_Os10g33940.1 11676.m02996/1-699											
LOC_Os02g04810.1 11668.m00419/1-1094											
LOC_Os04g59430.1 11670.m05892/1-530											
LOC_Os07g08520.1 11673.m00789/1-729											
LOC_Os07g08530.1 11673.m00790/1-408											
LOC_Os07g08600.1 11673.m00797/1-525											
ARF1/1-665											
ARF3/1-608											
ARF4/1-788											
ARF5/1-902											
ARF6/1-933											
ARF7/1-1165											
ARF8/1-811											
ARF9/1-638											
ARF10/1-693											
ARF11/1-601											
ARF12/1-593											
ARF13/1-623											
ARF14/1-605											
ARF15/1-598											
ARF16/1-670											
ARF17/1-585											
ARF18/1-602											
ARF19/1-1086											
ARF20/1-615											
ARF21/1-606											
ARF22/1-600											
ARF23/1-222											
ARF2/1-859											
PoptrARF8.1/1-827											
PoptrARF1.1/1-660											
PoptrARF7.3/1-1113											
PoptrARF2.3/1-792											
PoptrARF5.1/1-933											
PoptrARF5.2/1-944											
PoptrARF7.4/1-1137											
PoptrARF16.1/1-669											
PoptrARF1.2/1-662											
PoptrARF9.1/1-666											
PoptrARF2.1/1-854											
PoptrARF16.3/1-700											
PoptrARF3.1/1-709											
PoptrARF6.2/1-914											
PoptrARF6.4/1-953											
PoptrARF17.1/1-594											
PoptrARF2.4/1-879											
PoptrARF16.2/1-708											
PoptrARF2.2/1-852											
PoptrARF10.1/1-708											
PoptrARF3.3/1-109											
PoptrARF9.2/1-670											
PoptrARF6.1/1-884											
PoptrARF4/1-713											
PoptrARF6.3/1-163											
PoptrARF3.2/1-714											
PoptrARF9.3/1-579											
PoptrARF6.5/1-907											
PoptrARF9.4/1-632											
PoptrARF7.1/1-1047											
PoptrARF7.2/1-1093											
PoptrARF8.2/1-816											
PoptrARF16.6/1-91											
PoptrARF16.4/1-701											
PoptrARF16.5/1-536											
PoptrARF10.2/1-713											
PoptrARF17.2/1-592											
PoptrARF2.6/1-724											
PoptrARF2.5/1-614											

Consensus

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	2130	2140	2150	2160	2170	2180	2190	2200	2210
LOC_Os01g70270.1 11667.m07038_altsplice/1-809	-----	-----	-----	-----	-----	SNAPRKDDSSENEKGHL PMPNKSDNX	-----	-----	-----
LOC_Os01g13520.1 11667.m01333/1-700	-----	-----	-----	-----	-----	PKLLSSANPEQDQKTGF X	-----	-----	-----
LOC_Os02g06910.1 11668.m00642/1-909	-----	-----	-----	-----	-----	G I E L F S T S A R R L G N S C D N Y M S R Q E S R S L S T G I A S V G S V E F X	-----	-----	-----
LOC_Os02g35140.1 11668.m03331/1-757	-----	-----	-----	-----	-----	V V D A I D V E G P Q G I D M L A I A L R K H S L N V A Q G S N L E A L P F I T E L I I E I L V V Q T Q R K T Y I E V L S X	-----	-----	-----
LOC_Os02g41800.1 11668.m03990/1-784	-----	-----	-----	-----	-----	L Q P P G E A I P L G S L A L A A S S L T H D D E I C D L H R A A Y P W L V E S X	-----	-----	-----
LOC_Os04g57610.1 11670.m05695/1-819	-----	-----	-----	-----	-----	-----	G N D P R Y L S X	-----	-----
LOC_Os04g43910.1 11670.m04249/1-696	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os04g49230.1 11670.m04840/1-294	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os04g36060.1 11670.m03490/1-1674	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os05g43920.1 11682.m04181/1-700	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os06g09660.1 11680.m00929/1-1056	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os06g46410.1 11680.m04579/1-918	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os06g47150.1 11680.m04656/1-731	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os06g48950.1 11680.m04850/1-1116	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os08g40900.1 11674.m04106_altsplice/1-1056	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os11g32110.1 11687.m02952/1-812	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os12g41950.1 11686.m04180/1-900	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os12g29520.1 11686.m02861_altsplice/1-842	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os04g56850.1 11670.m05610_altsplice/1-956	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os05g48870.1 11682.m04683_altsplice/1-696	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os01g48060.1 11667.m04657_altsplice/1-723	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os01g54990.1 11667.m05421/1-656	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os10g33940.1 11676.m02996/1-699	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os02g04810.1 11668.m00419/1-1094	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os04g59430.1 11670.m05892/1-530	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os07g08520.1 11673.m00789/1-729	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os07g08530.1 11673.m00790/1-408	-----	-----	-----	-----	-----	-----	-----	-----	-----
LOC_Os07g08600.1 11673.m00797/1-525	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF1/1-665	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF3/1-608	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF4/1-788	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF5/1-902	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF6/1-933	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF7/1-1165	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF8/1-811	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF9/1-638	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF10/1-693	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF11/1-601	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF12/1-593	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF13/1-623	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF14/1-605	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF15/1-598	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF16/1-670	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF17/1-585	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF18/1-602	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF19/1-1086	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF20/1-615	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF21/1-606	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF22/1-600	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF23/1-222	-----	-----	-----	-----	-----	-----	-----	-----	-----
ARF2/1-859	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF8.1/1-827	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF1.1/1-660	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF7.3/1-1113	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF2.3/1-792	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF5.1/1-933	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF5.2/1-944	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF7.4/1-1137	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF16.1/1-669	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF1.2/1-662	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF9.1/1-666	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF2.1/1-854	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF16.3/1-700	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF3.1/1-709	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF6.2/1-914	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF6.4/1-953	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF17.1/1-594	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF2.4/1-879	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF16.2/1-708	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF2.2/1-852	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF10.1/1-708	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF3.3/1-109	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF9.2/1-670	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF6.1/1-884	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF4/1-713	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF6.3/1-163	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF3.2/1-714	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF9.3/1-579	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF6.5/1-907	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF9.4/1-632	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF7.1/1-1047	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF7.2/1-1093	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF8.2/1-816	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF16.6/1-91	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF16.4/1-701	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF16.5/1-536	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF10.2/1-713	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF17.2/1-592	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF2.6/1-724	-----	-----	-----	-----	-----	-----	-----	-----	-----
PoptrARF2.5/1-614	-----	-----	-----	-----	-----	-----	-----	-----	-----

Consensus

-----LGD+GNSPL PNQACSLSDAGEA+ RNSSDS I SSVGSL DYXEI SS- E-----T-R-----X