

		10	20	30	40	50
<i>AT1G01060.3_9/1-48</i>	TA	LQFENFHVWGD	QRWFLFYTLLVREDL	KQR	-----IAAPVLLWIL	IHLEKNY
<i>ALy_fgenesh2_kg.1_131_AT1G01060.3_9/1-</i>	TA	LQFGNFHVWGD	RRWFLYYILLVREDL	KQR	-----IAATVLLWIL	IYLEKNY
<i>BSt_Bostr.5325s0026.1_12/1-48</i>	TA	LQFGNFDVWGD	QRLFLYYTLLIREDL	KQR	-----IAAKILLWII	IHLEKNY
<i>CGr_Cagra.1968s0051.2_8/1-49</i>	TAL	LQFGNFHVLGD	QRWFLYYTLLVREDL	KQR	-----IAAKILLWIL	IHLEKNY
<i>CRu_Carubv10008551m_12/1-49</i>	TAL	LQFGNFHVLGD	QRWFLYYTLLVREDL	KQR	-----IAAKILLWIL	IHLEKNY
<i>RCa_30076.m004464_12/1-30</i>	TA	-LFVVSDFTRD	ERRFLFYPLFIREDL	KQH	-----	
<i>MES_Manes.01G239000.2_15/1-30</i>	TA	-LFVAFDIARN	ERRFLFYPLLREDL	KQR	-----	
<i>GRa_Gorai.008G132000.2_34/1-31</i>	TA	LLFVVLNVQRN	ERRFVFYPLLREDL	KQR	-----	
<i>GMa_Glyma.03G261800.4_10/1-31</i>	TA	LLFGVFHVTKN	ERRYSFYPLLREDL	KQR	-----	
<i>PVu_PhvuL.009G259600.2_11/1-47</i>	TA	LPFAVFHVTKN	ERRYSFYPLLREDL	KQR	-HTSSNGRLLW-	-RRSGY
<i>Ptr_Potri.014G106800.10_19/1-36</i>	TA	-LFVVL SVARN	EGRFLFYPLLREDL	KQR	-----	-W- -LPKVH
<i>SPu_SapurV1A.0162s0390.6_11/1-36</i>	TA	-LFVVL SVARN	EGRFPFYPLLREDL	KQR	-----	-W- -LPKVR
<i>TCa_EOX95555_33/1-50</i>	TA	LLFGVLDVQRN	ERRFVFYPLLGEDL	KQHYRCPGPVNGHILLW-	-RRTRY	
<i>CSa_Cucsa.349800.2_15/1-31</i>	TA	SPFR IYI PNN	ERRSFYPLLREDL	KQR	-----	
<i>CCI_Ciclev10018963m_12/1-29</i>	TA	LSFDT - - VRRN	EGRVLFYPLLREDL	KQR	-----	
<i>CSI_orange1.1g004257m_12/1-29</i>	TA	LSFDT - - ARRN	EGRVLFYPLLREDL	KQR	-----	
<i>PPe_EMJ20128_33/1-49</i>	TA	LLVIKLAAFIN	ERRSLFYPLLREDL	KQR	-KLRRSNGHTV IW-	-RRFGH
<i>LJa_Lj3g3v2518940.3_8/1-31</i>	SA	QFQPCFASQKN	ERRYSLYTLFVREDL	KQR	-----	
<i>SLo_Solyc10g005080.2.1_13/1-33</i>	TDR	LYLIVT DVVGR I	GRRFVLYPLFVPEDL	KQR	-----	