

cmr 7 - At5g35360 locus

	exon 2 \ / exon 3		
At2g31700	GGU <u>AUCGGUUCGGGUUCGGGUCGAU</u> : : ccaUAGCCAAGCCCAAGCCCAU <u>aau</u>	(186 210)	exon 2, exon 3
		-42.9	
	\ / exon 3		
	CGAU <u>UCGGUUCGGGUUCGGUUCGGG</u> : : : o ccaUAGCCAAGCCCAAGCCCAU <u>aau</u>	(207 231)	exon 3
		-30.9	
At1g75980	gggAUCGGUUCGGGUUCGGGU <u>Uugg</u> : : o ccaUAGCCAAGCCCAAGCCCAU <u>aau</u>	-38.3	3'UTR
At1g28540	GGAGAU <u>CGGAUCGGGUUCGGGUAAAA</u> : cc-aUAGCCAAGCCCAAGCCCAU <u>aau</u>	-34.9	
At3g16760	AGCUUCGGGAUCGGGUUCGGGU <u>AUUG</u> : o ccaUAG-CCAAGCCCAAGCCCAU <u>aau</u>	-34.4	
At5g05250	AGGAUCCGGGUCGGGUUUGGGU <u>AUGC</u> : : o : ccaUA-GCCAAGCCCAAGCCCAU <u>aau</u>	-31.2	
At1g13100	UGUGUCGGAU <u>CGGAUUCGGGUUAAA</u> o : ccaUAGCCAAGCCCAAGCCCAU <u>aau</u>	-29.6	

cmr 214 (MIR393) - At2g39890 locus (miR393a)

At3g62980	CAAUGCGAUCCCUUGGAUGUC GUUACGCUAGGGAAACCU <u>Agga</u>	-33.8
At3g26810	CAAUGCGAUCCCUUGGAUGUC GUUACGCUAGGGAAACCU <u>Agga</u>	-33.8
At1g12820	CAAUGCGAUCCCUUGGAUGUC GUUACGCUAGGGAAACCU <u>Agga</u>	-33.8
At4g03190	CCAUGCGAUCCCUUGGAUGUC GUUACGCUAGGGAAACCU <u>Agga</u>	-30.7
At4g19780	GUCUUCGAUCCCUUGGAUCCA : o GUUACGCUAGGGAAACCU <u>Agga</u>	-28.8
At4g17450	GUCUUCGAUCCCUUGGAUCCA : o GUUACGCUAGGGAAACCU <u>Agga</u>	-28.8

cand 1 - At5g58470 locus

At1g30040	GGGUCG CUGUCCCUCUGAUCA o: o uaCCGCGAUAGGGAGGACUCga	-34.4	
At5g53470	UGGGUGCUAUGCCUCCUGAGGA : o uaCCGCGAUAGGGAGGACUCga	-35.0	
At4g13430	AUGGUGUCAUCCCUCUGAUGC o o uaCCGCGAUAGGGAGGACUCga	-33.2	
At5g14210	UUGGCUCUGUCCCUCUGAUAU : o : uaCCGCGAUAGGGAGGACUCga	-33.1	
At4g08850	CUGGUGAAAUCCCUCUGAGAU o : : uaCCGCGAUAGGGAGGACUCga	-32.6	(964-985)
	CUGGAGCCAUCCACCUGAGAU : : uaCCGCGAUAGGGAGGACUCga	-26.9	(1612-1633)
At4g03205	AUGGUGUUAUGCCUCCUGAAGC o o uaCCGCGAUAGGGAGGACUCga	-31.9	
At1g03475	AUGGUGUUAUGCCUCCUGAAGC o o uaCCGCGAUAGGGAGGACUCga	-31.9	
At5g07280	CCGGCGAAAUCCCUCUGAAAU : : uaCCGCGAUAGGGAGGACUCga	-31.2	
At1g60720	UCGGCGCCUCCCUCUCAGAA : : : uaCCGCGAUAGGGAGGACUCga	-30.2	
At1g56130	UAGGACCUAUCCCUCAGAGCU : : uaCCGCGAUAGGGAGGACUCga	-29.9	
At1g61520	CUGGAGCUAUUGCUCUGAGAU : o : uaCCGCGAUAGGGAGGACUCga	-29.6	

cand 2 (MIR394) - At1g20380 locus (miR394a)

At1g27340	AGGAGGUUGACAGAAUGCCAAA uCCUCCACCUGUCUUACGGUUU	-35.2	
At1g10200	AGGAGGAGGAGAGAAUGCCGCC o uCCUCCACCUGUCUUACGGUUU	-31.3	
At4g24850/60	ACGAGGUGGACAGCAUGCUGG o oo uCCUCCACCUGUCUUACGGUUU	-30.9	overlapping transcripts
At5g20580	UGGAGGUGGAGAGAAUAUCAA : o uCCUCCACCUGUCUUACGGUUU	-29.2	

anti-cand 3 (MIR398) - At2g03450 locus (miR398a)

At3G15640	AAGGUGUGACCUGAGAAUCAC o : UUCCCACUGGACUCUUGUGU	-28.1	
At3G06370	UAGGGGUGACUUGAGUACACC : o : UUCCCACUGGACUCUUGUGU	-32.1	
At1G08830	AAGGGGUUCCUGAGAUACA : : UUCCCACUGGACUCUUGUGU	-28.8	