

hme-miR-2788

	Sequence	abundance	
GTCGCTGTATTGCTGGGGTTTCCTAGCGGCATGTGCCTTCTTCTATGCAATGCCCTTGGAAATCCCAAACGTGCTGGCGAC			
.....TGGGGTTTCCTAGCGGCAT.....		16	
.....TGGGGTTTCCTAGCGGCATG.....		102	
.....TGGGGTTTCCTAGCGGCATGT.....		234	
.....TGGGGTTTCCTAGCGGCATGTG.....		306	
.....TGGGGTTTCCTAGCGGCATGTGC.....		33	
.....TGGGGTTTCCTAGCGGCATGTGCCT.....		20	
.....CAATGCCCTTGGAAATCCCAA.....		2	
.....CAATGCCCTTGGAAATCCCAAA.....		27	hme-miR-2788*
.....CAATGCCCTTGGAAATCCCAAAC.....		2	
.....AATGCCCTTGGAAATCCCAAA.....		2	

hme-miR-193

CCAGCCTTGGTGAGGGTCTTGGCGGTCTAGTGGGTGTGCTCAGTTCTTACTGGCCTGCTAAGTCCCAAGCTATGGTGG			
.....AGGGTCTTGGCGGTCTAGT.....		1	
.....AGGGTCTTGGCGGTCTAGTG.....		20	
.....AGGGTCTTGGCGGTCTAGTGG.....		30	hme-miR-193*
.....AGGGTCTTGGCGGTCTAGTGGG.....		26	
.....TACTGGCCTGCTAAGTCCCA.....		2	
.....TACTGGCCTGCTAAGTCCCAA.....		39	
.....TACTGGCCTGCTAAGTCCCAAG.....		45	hme-miR-193