keywords: 3977 Expectation: 3.0 UPE: 25.0 Search Sort miRNA Acc. by: Expectation(E)

e.g. AT1G27360, miR156, transcription factor ... Range: 0.0 - 3.0 Range: 0.0 - 25.0

List of Predicted miRNA/Target Pairs [#Session ID: 1332363803601223]

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miRNA Acc.	Target Acc.	Expectation (E)	Target Accessibility (UPE)	Alignment	Target Description	Inhibition	Multiplicity
C17893406.Lu0910.pre-miRNA:241.miRNA:1349	Contig3977	3.0	9.165	miRNA 21 GCAGCGGUAGGAGGUAAAGAA 1 .::: :: ::.:::.::::: Target 544 UGUCCCCCUCUUCCGUUUCUU 564		Cleavage	1
C17893406.Lu0910.pre-miRNA:241.miRNA:1353	Contig3977	3.0	9.089	miRNA 21 CAGCGGUAGGAGGUAAAGAAG 1 ::: :: ::.:::.::::::: Target 545 GUCCCCCUCUUCCGUUUCUUC 565		Cleavage	1
scaffold521.Lu0910.pre-miRNA:2295.miRNA:13924	Contig3977	3.0	12.025	miRNA 20 UUGUUGGAAAUUUCGAUUUU 1 ::::::::::::.::. :::: Target 161 AACAACCUUUGGAGUGAAAA 180		Cleavage	1
scaffold521.Lu0910.pre-miRNA:2296.miRNA:13930	Contig3977	3.0	12.025	miRNA 20 UUGUUGGAAAUUUCGAUUUU 1 ::::::::::..::. :::: Target 161 AACAACCUUUGGAGUGAAAA 180		Cleavage	1
scaffold521.Lu0910.pre-miRNA:2297.miRNA:13936	Contig3977	3.0	12.025	miRNA 20 UUGUUGGAAAUUUCGAUUUU 1 ::::::::::::.::. :::: Target 161 AACAACCUUUGGAGUGAAAA 180		Cleavage	1