

Provisional ID : NMRB01005527.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold5527,\_whole\_genome\_shotgun\_sequence\_47684

Score total : 34.8

Score for star read(s) : 3.9

Score for read counts : 29.2

Score for mfe : 2.3

Score for randfold : 0

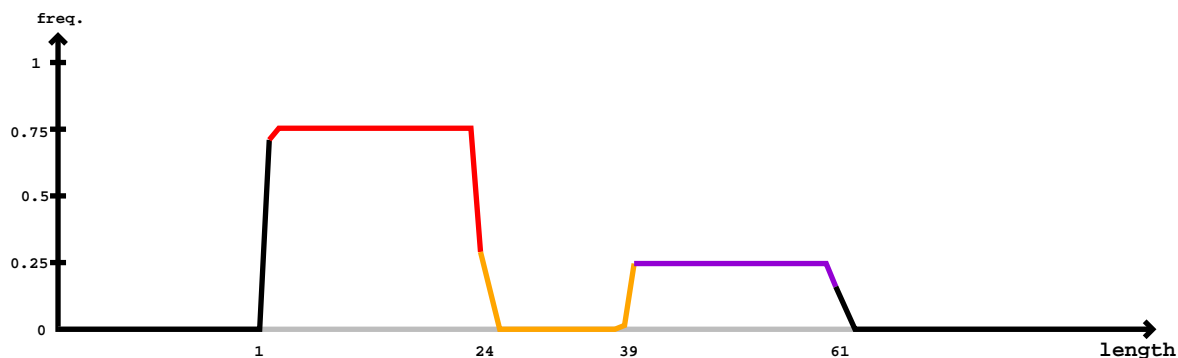
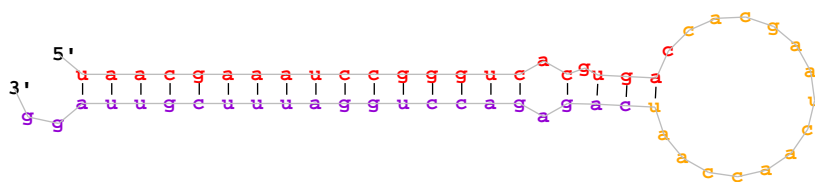
Score for cons. seed : -0.6

Total read count : 69

Mature read count : 52

Loop read count : 0

Star read count : 17



### Mature

### Star

5' -		-3'	obs	
			exp	
		reads	mm	sample
cgacguaccuguaccuggcc	uaacgaaaucgggucacgugac	cgagaccuggauuucguuag	1	Fm3
cgacguaccuguaccuggcc	uaacgaaaucgggucacgugac	cgagaccuggauuucguuag	12	Fm3
(((((((.....))))))))			1	Fm3
.....uaacgaaaucgggucacgugac			1	Fm3
.....uaacgaaaucgggucacgugac			1	Fm3
.....aacgaaaucgggucacgugac			1	Fm3
.....cagagaccuggauuucguuag			1	Fm3
.....cagagaccuggauuucguuag			1	Fm3
.....uaacgaaaucgggucacgugG			8	Mm2
.....uaacgaaaucgggucacguga			12	Mm2
.....uaacgaaaucgggucGcguga			1	Mm2
.....uaacgaaaucgggucacgugac			5	Mm2
.....aacgaaaucgggucacgugG			1	Mm2
.....aacgaaaucgggucacgugac			1	Mm2
.....ucagagaccuggauuucguuag			1	Mm2
.....cagagaccuggauuucguuU			2	Mm2
.....cagagaccuggauuucguuag			2	Mm2
.....cagagaccuggauuucguuagU			1	Mm2
.....cagagaccuggauuucguuagA			2	Mm2
.....cagagaccuggauuucguuagg			5	Mm2
.....cagagaccuggauuucguuaggU			1	Mm2
.....cagagaccuggauuucguuagga			1	Mm2
.....uaacgaaaucgggucacguga			7	Mm1
.....uaacgaaaucgggucacgugG			1	Mm1