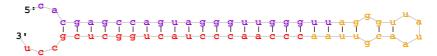
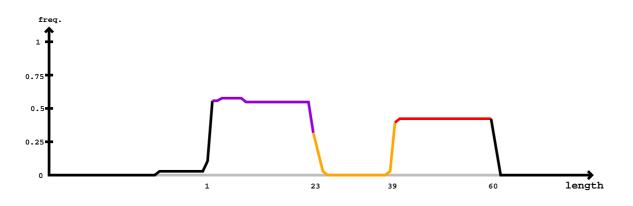
$: NMRB01007015.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold 7015, \_whole\_genome\_shotgun\_sequence\_48788$ Provisional ID

51.7 Score total

Score for star read(s) 3.9 45.5 Score for read counts 2.9 Score for mfe Score for randfold 0 -0.6



Score for cons. seed Total read count 101 44 Mature read count Loop read count 0 Star read count 57



	Star	Mature
5'- gggcgacgcagaaugacgccacagggaccuga	acacgagccaguaggguugg	gguuaggguuauaacguuaacccaacccuacuggcucgccucagguuccuguggauga

gggcgacgcagaaugacgccacagggaccugacacgagccaguaggguuggguuaggguuauaacguuaacccaacccuacuggcucgccucagguuccuguggaugacgc	ex	p	
	reads	mm	sample
cccaacccuacuggcuGgccu	1	1	Fm2
cccaacccuacuggcucgccu	5	0	Fm2
cagggaccugacacgagA	3	1	Fm3
acGcgagccaguaggguugggu	3	1	Fm3
cacgagccaguGggguuggguu	1	1	Fm3
cacgagccaguaggguuggguu	5	0	Fm3
cacgagccaguaggguuggguA	2	1	Fm3
.acccaacccuacuggcucgccu	1	0	Fm3
cccaacccuacuggcucgccu	14	0	Fm3
ccaGcccuacuggcucgccu	1	1	Fm3
acGcgagccaguaggguugggu	2	1	Mm2
acacgagccaguGggguugggu	1	1	Mm2
acacgagccaguaggguugggG.	1	1	Mm2
acacgagccaguaggguugggu	1	0	Mm2
cacgGgccaguaggguugggu	2	1	Mm2
cGcgagccaguaggguugggu	1	1	Mm2
cGcgagccaguaggguuggguu	5	1	Mm2
cacgagccaguaggguuggguu	6	0	Mm2
cacgagccaguaggguuU	1	1	Mm2
CacgagccaguaggguuaU	2	1	Mm2
CacgagccaguaggguuaA	1	1	Mm2
cgagccaguaggguua	2	0	Mm2
.acccaacccuacuggcucgccu.	1	0	Mm2
.acccaacccuGcuggcucgccu.	1	1	Mm2
cccaacccuacuggcucgccu	9	0	Mm2
cccaGcccuacuggcucgccu	1	1	Mm2
cccaacccuacuggcucgccG	1	1	Mm2
cccaacccuacuggcucgccuc	1	0	Mm2
ccaGcccuacuggcucgccu	2	1	Mm2

.....cacgagccagAaggguugggu.....

Star Mature

gggcgacgcagaaugacgccacagggaccugacacgagccaguaggguuggguuaggguuauaacguuaacccaacccuacugggcucgccucagguuccuguggaugacgc			
cGcgagccaguaggguugggu.	1	1	Mm1
cacgagccaguaggguugggu	11	0	Mm1
cacgagccaguaggguu	4	0	Mm1
cacgagccaguaggguuggguuU	1	1	Mm1
cacgGgccaguaggguu	1	1	Mm3
cacgagccaguGggguu	1	1	Mm3
cacgagccaguaggguuggguuŪ	1	1	Mm3
cccaacccuacuggcucgccu	5	0	Mm3
consGeneral condensation of the constant of th	1	1	Mm 3