

Provisional ID : NMRB01000592.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold592,\_whole\_genome\_shotgun\_sequence\_16585

Score total : 225.9

Score for star read(s) : 3.9

Score for read counts : 216.8

Score for mfe : 2.2

Score for randfold : 0

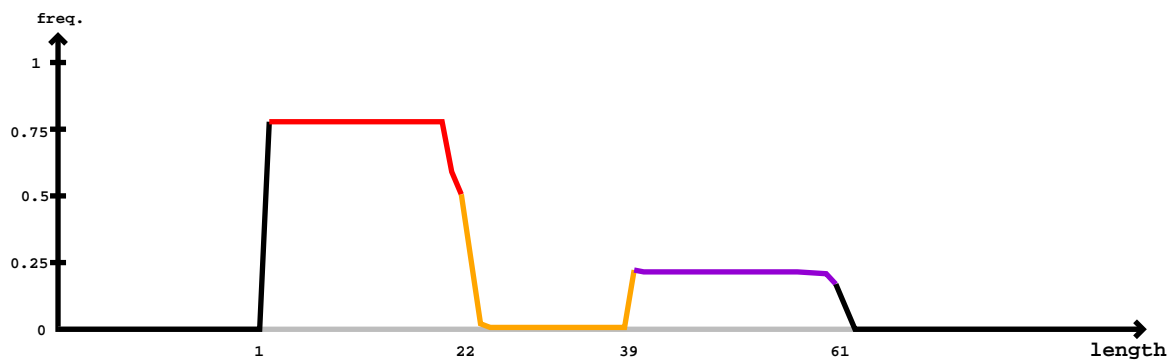
Score for cons. seed : 3

Total read count : 437

Mature read count : 340

Loop read count : 3

Star read count : 94

[illegible]

Star

Sequence	Count	Frequency	Category
ugaauuaccucucccuguggugcauugguaguugcauugcaucugaucuuuaagaaugcaauguauucugcagugcaaaacaagagggaggguccgucugccugaagaauucg	1	0	Fm1
.....caauguauucugcagugcaaaac.....	9	0	Fm1
.....caauguauucugcagugAaaac.....	1	1	Fm1
.....caauguauucugcaUugcaaaac.....	1	1	Fm1
.....caauguauucugcagugCCaaaca.....	1	1	Fm1
.....caauguauucugcagugcaaaaca.....	33	0	Fm1
.....caauguauucugcagugcUaaca.....	1	1	Fm1
.....caauguauucUAcagugcaaaaca.....	1	1	Fm1
.....caauguauucugcagugcaaaacC.....	2	1	Fm1
.....caauguauucugcagugcaaaacaU.....	1	1	Fm1
.....gugcauugguaguugcauug.....	7	0	Fm2
.....gugcauugguaguugcauugca.....	3	0	Fm2
.....gugcauugguaguugcauugcau.....	3	0	Fm2
.....ucugaucuuuaagaaugU.....	2	1	Fm2
.....gugcauugguaguugcauug.....	1	0	Mm2
.....gugcauugguaguugcauugca.....	7	0	Mm2
.....gugcauugguaguugcauugcau.....	1	0	Mm2
.....caauguauucugcagugcaaaaca.....	4	0	Mm2
.....caauguauucugcagugcUaaca.....	1	1	Mm2
.....gugcauugguaguugcauugca.....	20	0	Mm1