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.....ugaagcgccugugcucugccU...	4	1	GDM
.....ugaagcgGcugugcucugccg...	18	1	GDM
.....Agaagcgccugugcucugccg...	2	1	GDM
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.....ugaagcgccugugcucCgcccga...	30	1	GDM
.....ugaagcgccugugcucugccgU...	161	1	GDM
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.....ugaagcgccugugcCucgcccga...	26	1	GDM

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.....ugaagcgccugugcucugccgC..	16	1	GDM
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.....ugaagcgccugugcucugccgC..	16	1	GDM
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.....ugaagcgccAugcucugccga..	5	1	GDM
.....ugGagcgccugugcucugccga..	37	1	GDM
.....ugaaCcgccugugcucugccga..	3	1	GDM
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.....ugaagcgccugugcucAccga..	12	1	GDM
.....ugaagAgccugugcucugccga..	3	1	GDM
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.....ugaagcgccugugcucCgccga..	30	1	GDM
.....ugaGggcgccugugcucugccga..	24	1	GDM
.....ugaagcgccugugUucugccga..	5	1	GDM
.....ugUagcgccugugcucugccga..	11	1	GDM
.....ugaagcgccCgugcucugccga..	21	1	GDM
.....ugaagcgccugugcucugccAa..	10	1	GDM

Sequence	Count	Score	Method
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.....ugaagcgccugugcucugccga..	4840	0	GDM
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.....Agaagcgccugugcucugccga..	15	1	GDM
.....ugaagcgccugugcGcugccga..	3	1	GDM
.....ugaaCcgccugugcucugccga..	3	1	GDM
.....ugaagcgccuguaAcucugccga..	7	1	GDM
.....ugaagcgUcugugcucugccga..	6	1	GDM
.....ugaagcgccugugcucugcUga..	10	1	GDM
.....ugaagcgccAugucucugccga..	5	1	GDM
.....ugaagcgccugugcucugcUga..	10	1	GDM
.....ugaagcgccugugcGcugccga..	3	1	GDM
.....ugaagcgccugugcucugccgC..	16	1	GDM
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.....ugaagcgccugugcucugccAag..	3	1	GDM
.....ugaagcgccugugcucugccgaC..	73	1	GDM
.....ugaagcgccugugcucugccgaA..	281	1	GDM
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.....ugaagcgUugugcucugccgag..	2	1	GDM
.....ugaagcgccugugcucuaAccgag..	8	1	GDM
.....ugaagcgccugugcucugUcgag..	1	1	GDM
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.....ugaagcgccugugcucugccgaU..	763	1	GDM
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.....ugaagcgccugugcuUugccgag..	9	1	GDM
.....ugaagcgUcugugcucugccgag..	2	1	GDM

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.....ugaagcgccugugcucugccgaU.	763	1	GDM
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.....ugaagcgccugugcucugccgagU	403	1	GDM
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.....ugaagcgccugugcucugccgagG	4	1	GDM
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