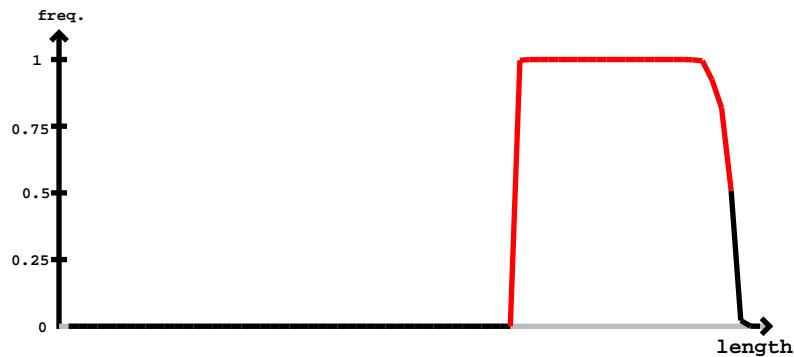
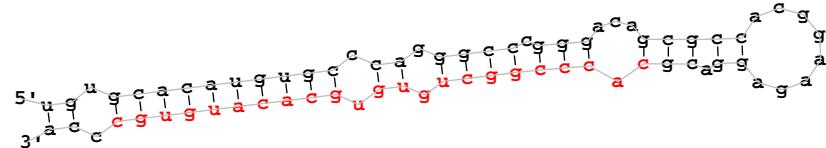


miRBase precursor	:	hsa-mir-941-3
Total read count	:	14188
hsa-miR-941 read count	:	14188
remaining reads	:	0



5' -	hsa-miR-941	-3'	exp	
		reads	mm	sample
.....	cacccggcugugugcacaugugccca	1	1	Z2_-
.....	cacccggcCgugugugcacaug.....	1	1	Z2_-
.....	cacccggcugugugcacaug.....	40	0	Z2_-
.....	Gaccggcugugugcacaugu.....	2	1	Z2_-
.....	cacccggcugugAgcacaugu.....	1	1	Z2_-
.....	cacccggcugugugugcacaugu.....	2	1	Z2_-
.....	cacccggcugugugugcacaugG.....	1	1	Z2_-
.....	cacccggcugugugugcacGugu.....	1	1	Z2_-
.....	cacccggcugugugugcacaugu.....	80	0	Z2_-
.....	cacccggcugugugugcacaugC.....	7	1	Z2_-
.....	cacccgAgcugugugcacaugu.....	1	1	Z2_-
.....	cacccggcugugugGgcacaugug.....	1	1	Z2_-
.....	cacccggcugugugAcacaugug.....	1	1	Z2_-
.....	cacccggcugCgugcacaugug.....	2	1	Z2_-
.....	cUccggcugugugugcacaugug.....	1	1	Z2_-
.....	caccUggcugugugugcacaugug.....	1	1	Z2_-
.....	cacccAgcugugugugcacaugug.....	1	1	Z2_-
.....	cacccggcugugugugAcacaugug.....	1	1	Z2_-
.....	cacccggcugugugugugcacaugug.....	1	1	Z2_-
.....	cacccggcAuggugugcacaugug.....	1	1	Z2_-
.....	cacccggcCgugugugcacaugug.....	1	1	Z2_-
.....	cacccggcugugugugugcacaugug.....	1	1	Z2_-
.....	cacccggcugugugugcacCugug.....	1	1	Z2_-
.....	cacccggcugugugugcacaugugCg.....	1	1	Z2_-
.....	cGcccgguugugugugcacaugug.....	1	1	Z2_-
.....	cacccggcugugCgcacaugug.....	1	1	Z2_-
.....	Uacccggcugugugugcacaugug.....	1	1	Z2_-
.....	cacccUgcugugugugcacaugug.....	1	1	Z2_-
.....	cacccggcugugugGcacaugug.....	11	1	Z2_-
.....	cacccggcAuuggugugcacaugug.....	1	1	Z2_-
.....	cacccggUuggugugugcacaugug.....	1	1	Z2_-
.....	cacccggcugugugugcacaugu.....	3	1	Z2_-
.....	cacccggcugugugugcacaugAg.....	2	1	Z2_-
.....	cacccggcugugugugcacauguA.....	14	1	Z2_-
.....	cacccggcugugugugcacaugug.....	277	0	Z2_-
.....	cacccggcugugugugCcacaugug.....	1	1	Z2_-
.....	cacccggcugugugugcacaugugG.....	1	1	Z2_-

ugugcacauugugcccccaggggccccgggacagcgccacggaaaggaggacgcacccggcugugugcacauugugccca
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.....	cacccggcugugugcacauA.....	24	1	GDM
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hsa-miR-941

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.....	cacccggcugugugGcaugug.....	17	1	GDM
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.....	cacccggcugugugGacaugug....	45	1	GDM
.....	Aacccggcugugugcacauug....	2	1	GDM
.....	cacccggcugugugcacauu....	7	1	GDM
.....	caAcggcugugugcacauug....	1	1	GDM
.....	cacccggcugugugcacauAg....	2	1	GDM
.....	cacccggcuAuggcacauug....	2	1	GDM
.....	cacccggcuguguAcacauug....	3	1	GDM
.....	cacccggcugugugcacauu....	113	1	GDM
.....	cacccggcuguguUcacauug....	1	1	GDM
.....	cacccggcugugugcacauug....	71	1	GDM
.....	cacccggcuguguAcacauug....	5	1	GDM
.....	cacccggcugugugcacacGug....	9	1	GDM
.....	cacccggcugugugcacauugC....	4	1	GDM
.....	cacccggcuguUugcacauug....	1	1	GDM
.....	caccAgcugugugcacauug....	4	1	GDM
.....	cacccggcugugugcacauu....	1	1	GDM
.....	cacccggcugugugUcaugug....	7	1	GDM
.....	cacccggcugugugCcaugug....	1	1	GDM
.....	cacccggcugugugcacauug....	6	1	GDM
.....	cacccggcuguguAcacauug....	5	1	GDM
.....	cacccggcugugAgcacauug....	3	1	GDM
.....	cacccggcugugugcacauu....	113	1	GDM
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.....	cacccgUcugugugcacauug....	1	1	GDM
.....	cacccggUugugugcacauug....	5	1	GDM
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.....	cacUcgcugugugcacauug....	3	1	GDM
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.....	cacccggcugugugcacacGug...	5	1	GDM
.....	cacccggcugugugGcaugugc...	5	1	GDM
.....	cacccggcugugugGcacauugc...	28	1	GDM
.....	cGcccgcgugugugcacauugc...	9	1	GDM
.....	cacccggcugugugGacaugugc...	28	1	GDM
.....	cacUcgcugugugcacauugc...	1	1	GDM
.....	caGcgcugugugcacauugc...	3	1	GDM
.....	cacccggcugugugcacauugc...	1538	0	GDM
.....	cacccggcCgugugcacauugc...	11	1	GDM
.....	caccoCgcugugugcacauugc...	1	1	GDM
.....	cacccggcugugugGcaugugc...	5	1	GDM
.....	caUcgcugugugcacauugc...	3	1	GDM
.....	cacccggcugugugcacauugU...	842	1	GDM
.....	cacccggcugugugcacauu....	407	1	GDM
.....	cacccggcugCgugcacauugc...	4	1	GDM
.....	caccGgcugugugcacauugc...	2	1	GDM
.....	cacccggcugugugcacauugG...	11	1	GDM
.....	cacccggcugugugcacauu....	3	1	GDM

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.....	caccGGGcugugugcacauugCg..	6	1	GDM
.....	caccGGGcugugugcacauugG..	11	1	GDM
.....	cacAAGggcugugugcacauugc..	3	1	GDM
.....	Uacccggcugugugcacauugc..	6	1	GDM
.....	cacccggcugugugUacauugc..	4	1	GDM
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.....	cacccggcugugugGacauugc..	28	1	GDM
.....	cacccggcugugugcaUauguc..	4	1	GDM
.....	cacccggcugugugcacauguA..	3	1	GDM
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.....	cacccggcugugugcacGuguc..	11	1	GDM
.....	cacccggcuaAugugcacauugc..	2	1	GDM
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.....	Uacccggcugugugcacauugc..	6	1	GDM
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.....	cacccggcugugugcacGuguc..	11	1	GDM
.....	cacccggcugugugcacauugA..	407	1	GDM
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.....	acccggcugugugcacauugc..	9	0	GDM
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.....	acccggcugugugcacauugU..	3	1	GDM

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ugugcacauugugccaggggccccgggacagcgccacggaaaggaggacg	cacccggcugugugcacauugugcca			
.....accggcugugugcacauugc...	9	0	GDM
.....accggcugugugcacauugU...	3	1	GDM
.....accggcugugugcacauugc...	9	0	GDM
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.....cacccggcugugugcacau...	3	1	Z3_
.....cacccggcugugCgcacau...	8	1	Z3_

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.....	Gacccggcugugugcacauuguc...	1	1	Z3_-
.....	cUcccggcugugugcacauuguc...	3	1	Z3_-
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.....	cacccggCugugugcacauuguc...	1	1	Z3_-
.....	cacccggcugugugGcacauuguc...	7	1	Z3_-
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.....	cGcccggcugugugcacauuguc...	1	1	Z3_-
.....	cacccggcugugugcacauugcA...	2	1	Z3_-
.....	cacccggcugugugcacauugcU...	28	1	Z3_-
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.....	acccggcCugugugcacauuguc...	1	1	Z3_-