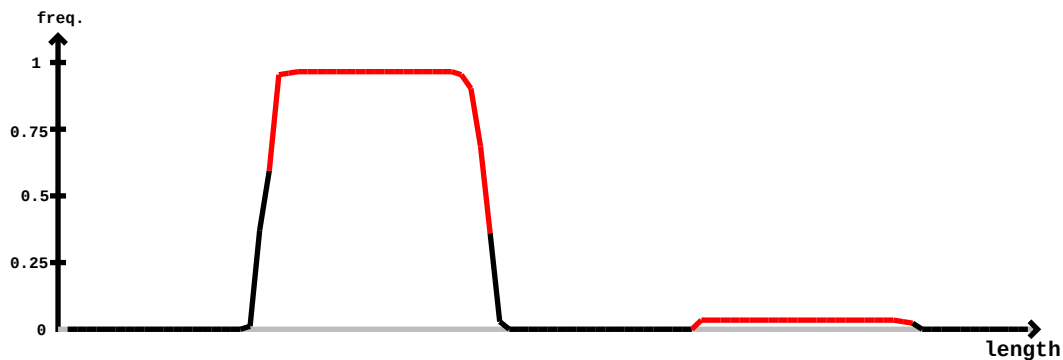
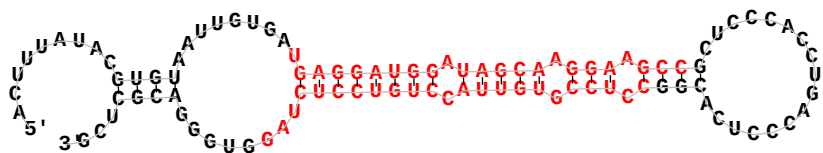


miRBase precursor : hsa-mir-3605
 Total read count : 175
 hsa-miR-3605-5p read count 167
 hsa-miR-3605-3p read count 6
 remaining reads : 8



hsa-miR-3605-3p

hsa-miR-3605-5p		hsa-miR-3605-3p			
5'	acuuuauacguguaauugugaugaggauaggauagcaaggaagccgcuccaccugaccucacggccuccguguuaccuguccucuaugguggagcgcucg	-3'	exp		
	reads	mm	sample		
.....(((.....(((.....(((.....(((.....)))))).....)))).....)))).....	1	0	hsa		
.....ugaggauaggauagcaagga.....	1	0	hsa		
.....ugaggauaggauagcaaggaU.....	1	1	hsa		
.....gaugaggauaggauagcaagga.....	1	0	hsa		
.....gaugaggauaggauagcaagga.....	5	0	hsa		
.....gaugaggauaggauagcaagga.....	1	1	hsa		
.....gaugaggauaggauagcaaggaA.....	1	1	hsa		
.....gaugaggauaggauagcaaggaag.....	20	0	hsa		
.....Uaugaggauaggauagcaaggaag.....	1	1	hsa		
.....gauAaggauaggauagcaaggaag.....	1	1	hsa		
.....Caugaggauaggauagcaaggaagc.....	1	1	hsa		
.....gaugaggauaggauagcaaggaagc.....	17	0	hsa		
.....gaugaggauaggauagcaaggaagcc.....	10	0	hsa		
.....gaugaggauaggauagcaaggaagcU.....	1	1	hsa		
.....gaugaggauaggauagcaaggaagcA.....	2	1	hsa		
.....gaugaggauaggauagcaaggaagccU.....	1	1	hsa		
.....gaugaggauaggauagcaaggaagccA.....	1	1	hsa		
.....augaggauaggauagcaaggaag.....	9	0	hsa		
.....augaggauaggauagcaaggaagc.....	17	0	hsa		
.....augaggauaggauagcaaggaagcc.....	12	0	hsa		
.....augaggauaggauagcaaggaagcU.....	1	1	hsa		
.....ugaggauaggauagcaaggaag.....	1	0	hsa		
.....ugaggauaggauagcaaggaagA.....	1	1	hsa		
.....ugaggauaggauagcaaggaag.....	5	0	hsa		
.....ugaggauaggauagcaaggaagc.....	21	0	hsa		
.....ugaAgauggaugauagcaaggaagc.....	1	1	hsa		
.....ugaggauaggauagcaaggaagcU.....	1	1	hsa		
.....ugaggauaggauagcaaggaagcc.....	29	0	hsa		
.....ugaggauaggauagcaaggaagcA.....	2	1	hsa		
.....ugaggauaggauagcaaggaagccU.....	1	1	hsa		
.....ugaggauaggauagcaaggaagcUg.....	1	1	hsa		
.....gaUgauggaugauagcaaggaag.....	1	1	hsa		
.....aggauaggauagcaaggaagccg.....	1	0	hsa		
.....ccuccguguuaccuguccucu.....	1	0	hsa		
.....ccuccguguuaccuguccucua.....	1	0	hsa		

hsa-miR-3605-3p

hsa-miR-3605-5p

acuuuauacguguaauugugaugaggauggauagcaaggaagccgcuccaccugaccucacggccuccguguuaccugucccuaggugggacgcucg

.....ccuccguguuaccugucccuag..... 4

0

hsa