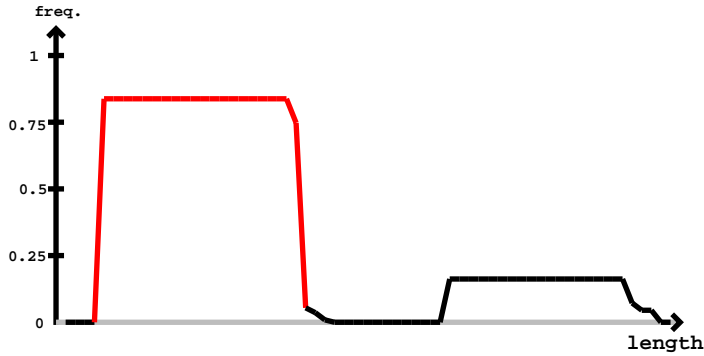
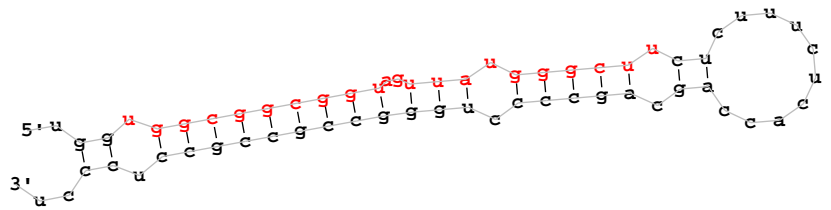


miRBase precursor : hsa-mir-4467
 Total read count : 111
 hsa-miR-4467 read count : 93
 remaining reads : 18



hsa-miR-4467

5'-	3'	exp	reads	mm	sample
UGGUGGCGGCGGUAGUUAUGGGCUUCUCUUUCUCACCAGCAGCCCCUGGGCGCGCCUCCU					
..(((((((((((..((((((((((.....))))))))))))))))))..			2	1	Z3_
..UGGCGGCGGUAGUUAUGGGCA.....			3	0	Z3_
..UGGCGGCGGUAGUUAUGGGCU.....			2	0	Z1_
..UGGCGGCGGUAGUUAUGGGCC.....			1	1	Z1_
..UGGCGGCGGUAGUUAUGGGC.....			1	1	GDM
..UGGCGGCGGUAGUUAUGGGCU.....			23	0	GDM
..UGGCGGCGGAGUUAUGGGCU.....			1	1	GDM
..UGGCGGCGGUAGUUAUGGGCC.....			4	1	GDM
..UGGCGGCGGUAGUUAUGGGCU.....			23	0	GDM
.....CAGCCCCUGGGCGCGCGCU.....			2	1	GDM
.....CAGCCCCUGGGCGCGCGCC.....			7	0	GDM
.....CAGCCCCUGGGCGCGCGCC.....			7	0	GDM
.....CAGCCCCUGGGCGCGCGCC.....			7	0	GDM
.....CAGCCCCUGGGCGCGCGCU.....			2	1	GDM
.....CAGCCCCUGGGCGCGCGCCU.....			2	0	GDM
.....CAGCCCCUGGGCGCGCGCCUCC.....			3	0	GDM
.....CAGCCCCUGGGCGCGCGCCUCU.....			1	1	GDM
..UGGCGGCGGUAGUUAUGGGU.....			2	1	Z2_
..UGGCGGCGGUAGUUAUGGGC.....			6	0	Z2_
..UGGCGGCGGUAGUUAUGGGAA.....			1	1	Z2_
..UGGCGGCGGUAGUUAUGGGA.....			1	1	Z2_
..UGGCGGCGGUAGUUAUGGGCA.....			3	1	Z2_
..UGGCGGCGGUAGCUAUGGGCU.....			1	1	Z2_
..UGGCGGCGGUAGUUAUGGGCU.....			27	0	Z2_
..UGGCGGCAGUAGUUAUGGGCU.....			2	1	Z2_
..UGGCGGCGGUAGUUAUGGGCC.....			7	1	Z2_
..UGGCGGCGGUAGUUAUGGGCUU.....			2	0	Z2_
..UGGCGGCGGUAGUUAUGGGCUUA.....			3	1	Z2_
..UGGCGGCGGUAGUUAUGGGCUUCU.....			1	0	Z2_
.....CAGCCCCUGGGCGCGCGCC.....			1	0	Z2_
.....CAGCCCCUGGGCGCGCGCCU.....			1	0	Z2_
.....CAGCCCCUGGGCGCGCGCCUCU.....			1	1	Z2_

hsa-miR-4467

ugguggcggcgguagguuagggcuucucuuucucaccagcagccccugggcgcgcucccu