

hsa-miR-3177-3p

hsa-miR-3177-5p

ccacgugccauguguacacacgugccaggcgcugucuuugagacauucgcgagugcacggcacuggggacacgugggcacugg

.....ugcacggcacuggggacacgug.....	18	0	GDM
.....ugcacggcacuggggacacgGg.....	2	1	GDM
.....ugcacggcacuggggacacgug.....	18	0	GDM
.....ugcaUggcacuggggacacgug.....	1	1	GDM
.....ugAacggcacuggggacacgug.....	1	1	GDM
.....ugcacggcacugggAgacacgug.....	3	1	GDM
.....ugcacggcacuggggacacgug.....	18	0	GDM