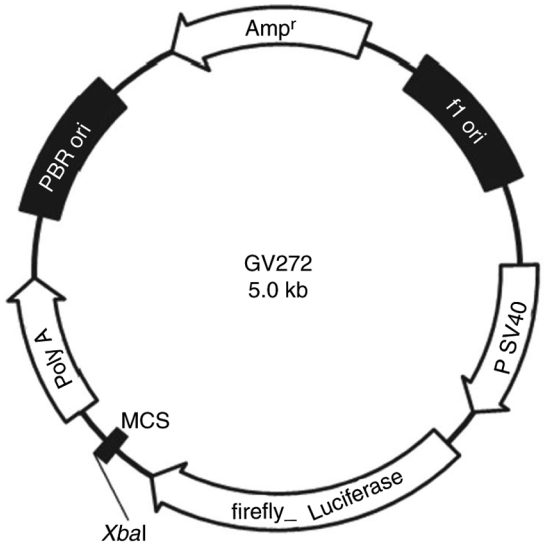
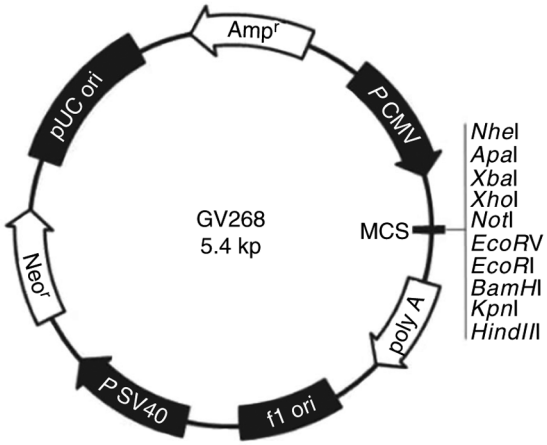
Figure S1. GV272 plasmid information. MCS, multiple cloning site.



General: 5010 bp
SV40_promoter: 48–250
firefly_Luciferase: 280–1932
SV40 late poly A signal: 1964–2185
pBR322_origin: 3117–2498
Ampicillin : 4132–3272
f1 ori : 4719–4264
Primer locations and sequences:
Luc-C-F (1820–1839) : GAGGAGTTGTGTTTGTGGAC
RVprimer4 (2272–2253) : GACGATAGTCATGCCCCGCG

Figure S2. GV268 plasmid information. MCS, multiple cloning site.



General: 5427 bp
CMV promoter: 232–819
BGH poly A: 1027–1251
f1 ori: 1297–1725
SV40 promoter: 1730–2073
Neomycin: 2135–2929
pUC ori: 4286–3616
Ampicillin: 5291–4431
Primer locations and sequences:
CMV-F (769–789): CGCAAATGGGCGGTAGGCGTG
pcDNA-SEQR (1121–1101): TTATTAGGAAAGGACAGTGGG

Figure S3. Reverse transcription-quantitative PCR analysis of 293T cells transfected with GV268/miR-146a and negative control. Data are presented as the mean ± standard error of the mean (n=3). **P<0.01.