

Supplementary Table 3 Percentages of methylated CpG in total CpGs identified using bisulfite-sequencing method (in *RANKL* gene, stable transfectant cell line).

Group names	Left to right: -700 to TSS site (<i>RANKL</i> promoter)
blank	10%, 20%, 40%, 10%, 30%, 50%, 70%, 50%, 40%
-sgRNA	0%, 30%, 30%, 20%, 20%, 60%, 90%, 80%, 80%
R3: dCas9-CD+MS2-CD	0%, 30%, 10%, 10%, 10%, 20%, 60%, 40%, 50%
R8: dCas9-CD+MS2-CD	0%, 0%, 0%, 10%, 10%, 40%, 90%, 70%, 50%