

E-cadherin promoter region

-250 GGC GGGGCTG GGATT CGA ACC CAGT GGAAT CAGA ACC GTGCAGG **TCCCAT**
-200 **AACC** CACCT **AGACCCTAGCA** ACTCCAGGCTAGAGGGTCACCGCGTCTATG
-150 CGAGGCCGGGTGGGCGGGCCGTCAGCTCCGCCCTGGGGAGGGGTCCGCGC
-100 TGCTGATTGGCTGTGGCCGGCAGGTGAACCCTCAGCCAATCAGCGGTACG
-50 GGGGGCGGTGC**CTCCGGGGCT** CACCT **GGCTGCAGCCACGCACCCCTCTC**

: ZEB1 binding site; Red: EMSA probe

Fig. S1

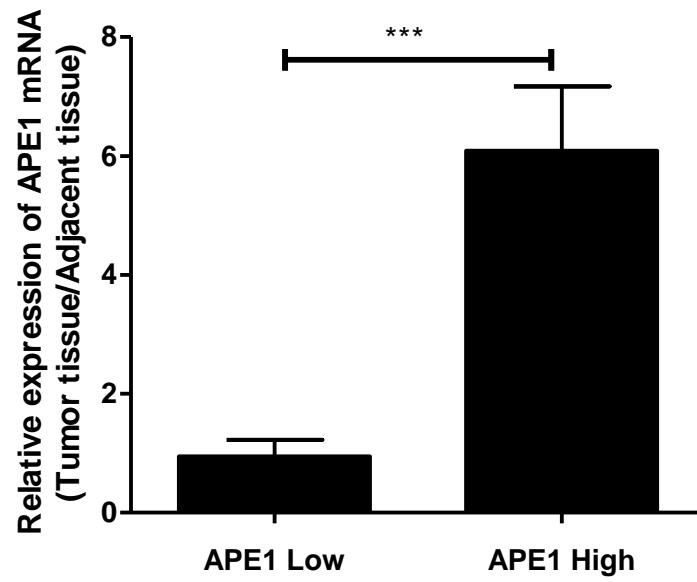


Fig. S2

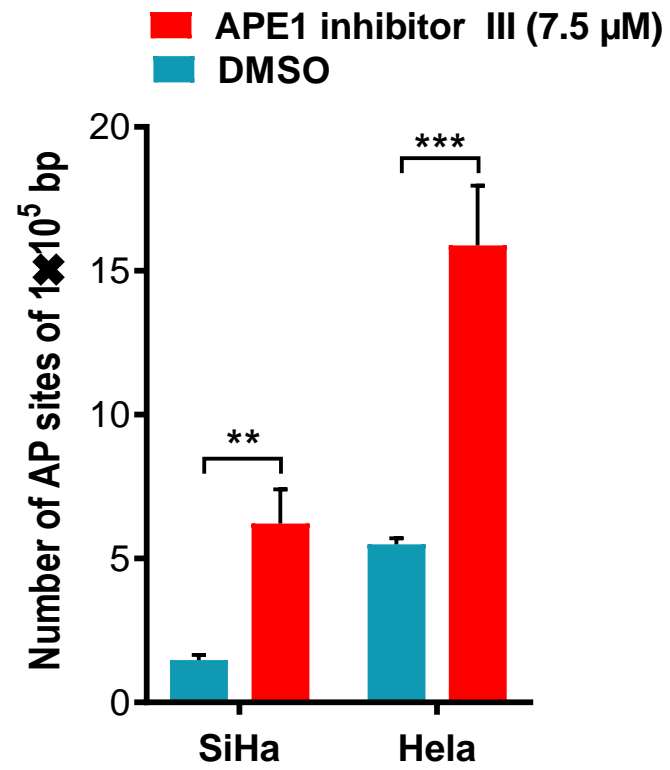


Fig. S3

561 TGDGNLSPSQ PPLKNLLSLL KAYYALNAQP **Wild type**
561 TGDGNLSPSQ PPLKNLL**PIV** KAYYALNAQP **Mutant**

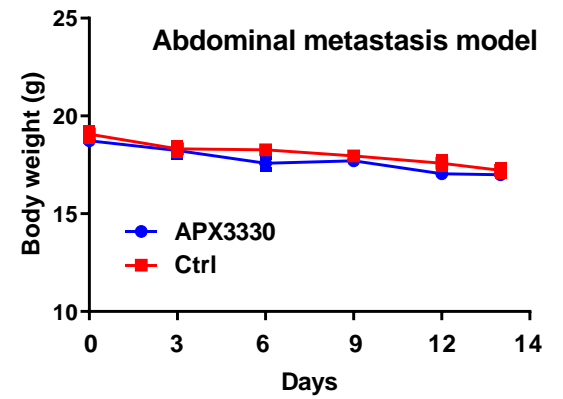
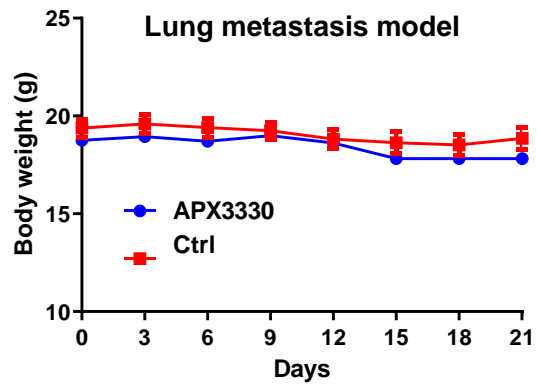
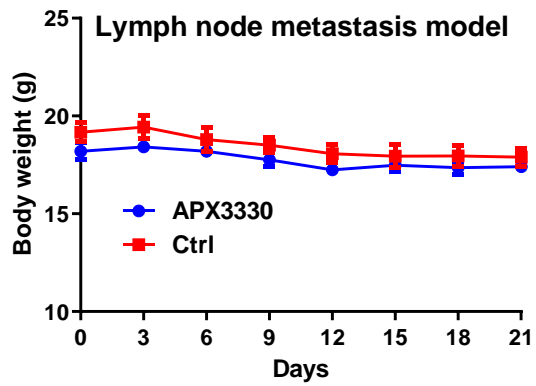


Fig. S6