1	Association of polymorphisms at the microRNA binding site of the caprine KITLO
2	3'-UTR with litter size
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4	Xiaopeng An ^{1a} , Yuxuan Song ^{1a} , Shuhai Bu ^{1b} , Haidong Ma ^a , Kexin Gao ^a , Jinxing Hou ^a
5	Shan Wang ^a , Zhang Lei ^a , Binyun Cao ^a *
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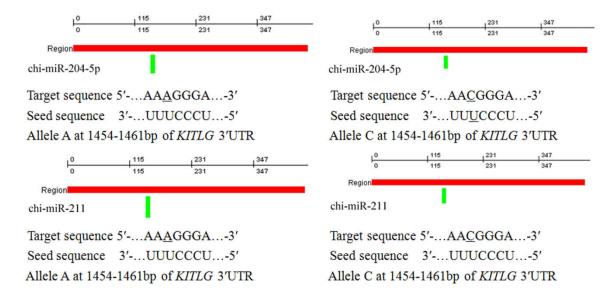


Figure S1 Effects of SNP c.1457A>C on miRNA binding sites in the caprine *KITLG* 3'UTR. The green boxes represent potential miRNA target sites. The underline indicates the position of base mutation

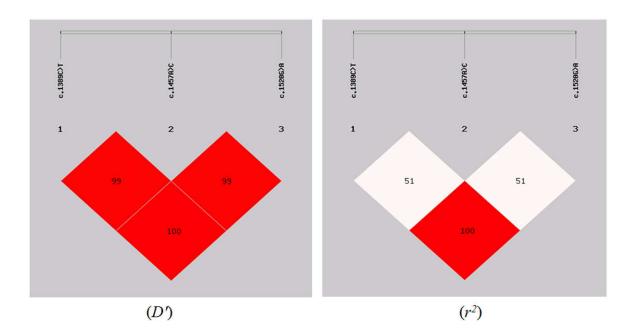


Figure S2 Linkage disequilibrium (LD) analysis of three SNP loci in the GD goats. Numbers in each cell represent pairwise D' and r^2 values (%).

Note: When $r^2 > 0.33$ and D' > 0.5, the linkage disequilibrium was considered strong 22,23 .

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