Cite this as: Dongyun XIN, Yangyang BAI, Yi BI, Libang HE, Yuxin KANG, Chuanying PAN, Haijing ZHU, Hong CHEN, Lei QU, Xianyong LAN. Insertion/deletion variants within the *IGF2BP2* gene identified in reported genome-wide selective sweep analysis reveal a correlation with goat litter size[J]. Journal of Zhejiang University Science B, 2021, 22(9): 757-766. http://doi.org/10.1631/jzus.B2100079

Insertion/deletion variants within the *IGF2BP2* gene identified in reported genome-wide selective sweep analysis reveal a correlation with goat litter size

Keywords: goat, IGF2BP2, litter size, indel, MAS

Research Summary

◆ The primary aims of the study were to verify the relationship between indels in *IGF2BP2* and Shaanbei white cashmere (SBWC) goat litter sizes and to select the favorable alleles. We also wanted to lay a foundation for the application of MAS in goat breeding, to improve the economic efficiency of the goat livestock industry.



- → SNP (Single nucleotide polymorphism)
- ◆ InDel (Insertion/Deletion)
 - ◆ CNV(Copy number variation)

Marker-assisted selection (MAS), which is characterized as an alternative selection of DNA mutations, is not affected by the microenvironment. In addition, compared with traditional breeding techniques, MAS is a more accurate, efficient and convenient genetic selection method

Innovation points

The InDel locus of *IGF2BP2* gene in goats was screened and verified, and the correlation and function of *IGF2BP2* gene with lambing traits of goats were determined, so as to provide a theoretical basis for improving economic traits and breeding new varieties of goats.

- ◆ In this study, the mRNA expression level of *IGF2BP2* was found to be significantly higher in a single-lamb group than in a multilamb group. Insertions/deletions (indels) within the goat *IGF2BP2* gene, including P4-Ins-13bp and P5-Del-12bp, were verified in 918 Shaanbei white cashmere (SBWC) female goats.
- ♦ Analysis using the Chi-square ($χ^2$) test showed that the genotype ($χ^2$ = 14.479, P = 0.006) and allelotype ($χ^2$ = 5.228, P = 0.022) distribution of P4-Ins-13bp was significantly different between the single-lamb and multilamb groups.

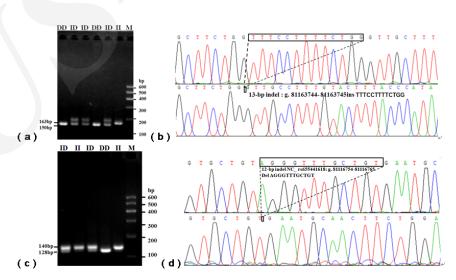


Fig 3

Innovation points

- ♦ Correlation analysis demonstrated that P4-Ins-13bp was significantly associated with goat litter size (P = 0.022), and individual goats with the deletion/deletion (DD) genotype produced more litters than other goats
- **◆**The JASPAR online database predicted that the binding of TFs GATA-binding factor 2 (GATA2) [MA0036.3] and Forkhead box L1 (FOXL1) [MA0033.2] might bind to the P4-Ins-13bp and P5-Del-12bp insertion sequences in the *IGF2BP2*, respectively



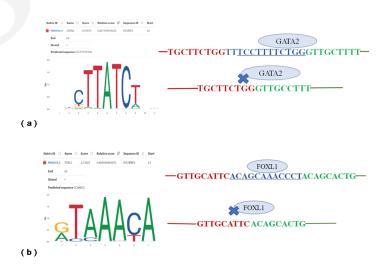


Fig 7