Genomic selection for growth, reproductive and conformation traits in Zebu Brahman in Colon





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Aim

 Estimate the increase of breeding value accuracies in traditional breeding scheme compared to genomic selection

Background

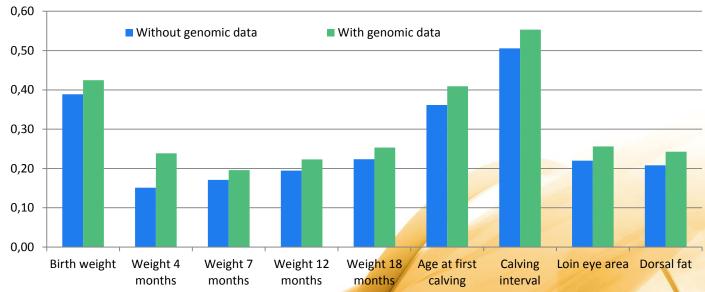
- Colombia a country with beef production focus
- Main beef breed Zebu Brahaman cattle breeds and their crosses
- Current selection pedigree based only
- Improvement under way Openness of farmers and breeding organizations towards genomic technologies

Conclusions

- Substantial increase in accuracies when genomic data was used to estimate breeding values
- Increase higher in growth traits
- Increased accuracies expected to increase genetic gain



Results



- Results based on data from 120,000 animals from 35 farms in Colombia
- Genotypes available from 4,250 animals
- Single step genomic BLUP to estimate breeding values, using the via the BlupF90 program (Misztal and Aguilar 2010)
- Focus on increasing the accuracies of breeding value estimates for weight, conformation and reproduction traits
- Considerable responses in all studied traits
- Weight related traits +9 to +55% increase
- Reproduction related traits +8 to +16% increase
- Conformation related traits +16% increase

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