The application of mixed linear models (LMM) for the estimation of functional effects on bovine stature based on SNP summary statistics from a whole-genome association study

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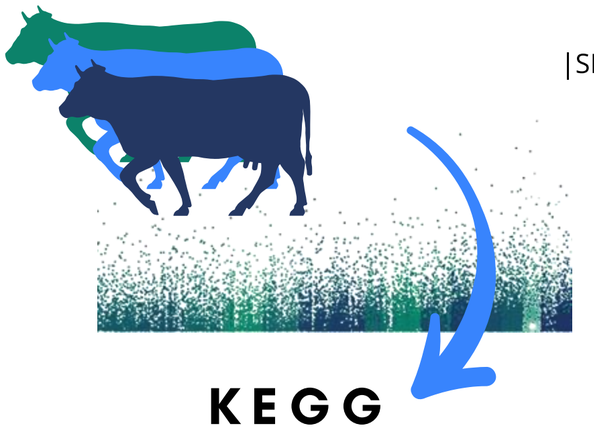
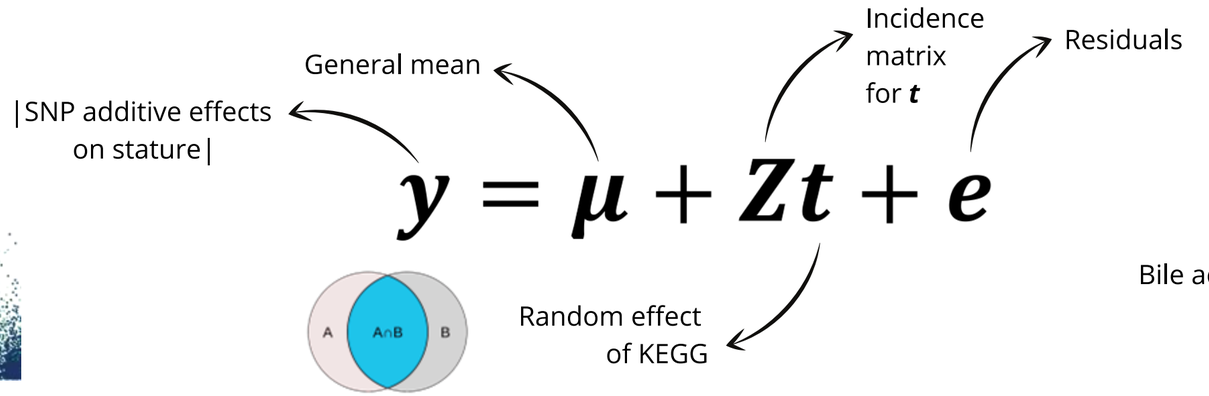
1 GWAS VS. UNITS



Three different bulls' breeds:

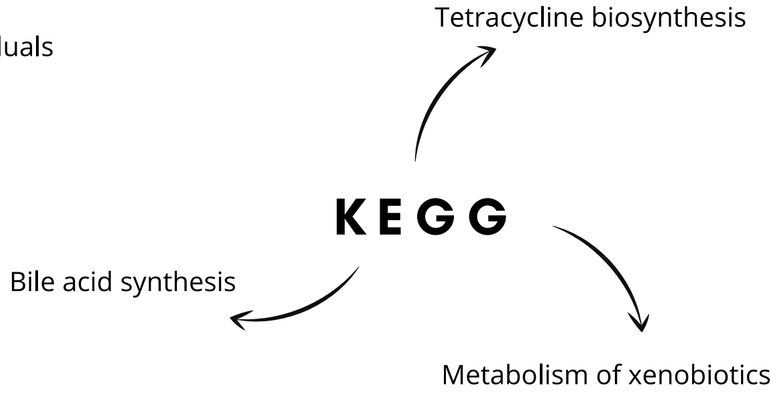
- Danish Holstein (366 877 SNPs)
- Danish Red Dairy (299 723 SNPs)
- Finnish Red Dairy (396 224 SNPs)

2 LMM



- The similarity between KEGGs → Jaccard Coefficients
- Henderson mixed model equations → Random effects
- Calculations → NumPy

3 RESULTS



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