

Using International Information In National Single Step Genomic BLUP In Swiss Dairy Cattle

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Introduction

More stuff here

In Swiss dairy cattle breeding, genomic breeding values are estimated using marker-effect models (MEM) and a Bayesian Regression approach (Bayes $A - C$ and C_π). Estimated genomic breeding values are combined with traditional BLUP-based breeding values via an index procedure. MEM do not provide breeding value estimates for animals that are not genotyped.