

NGS-based analysis of Copy Number Variations in various cattle breeds

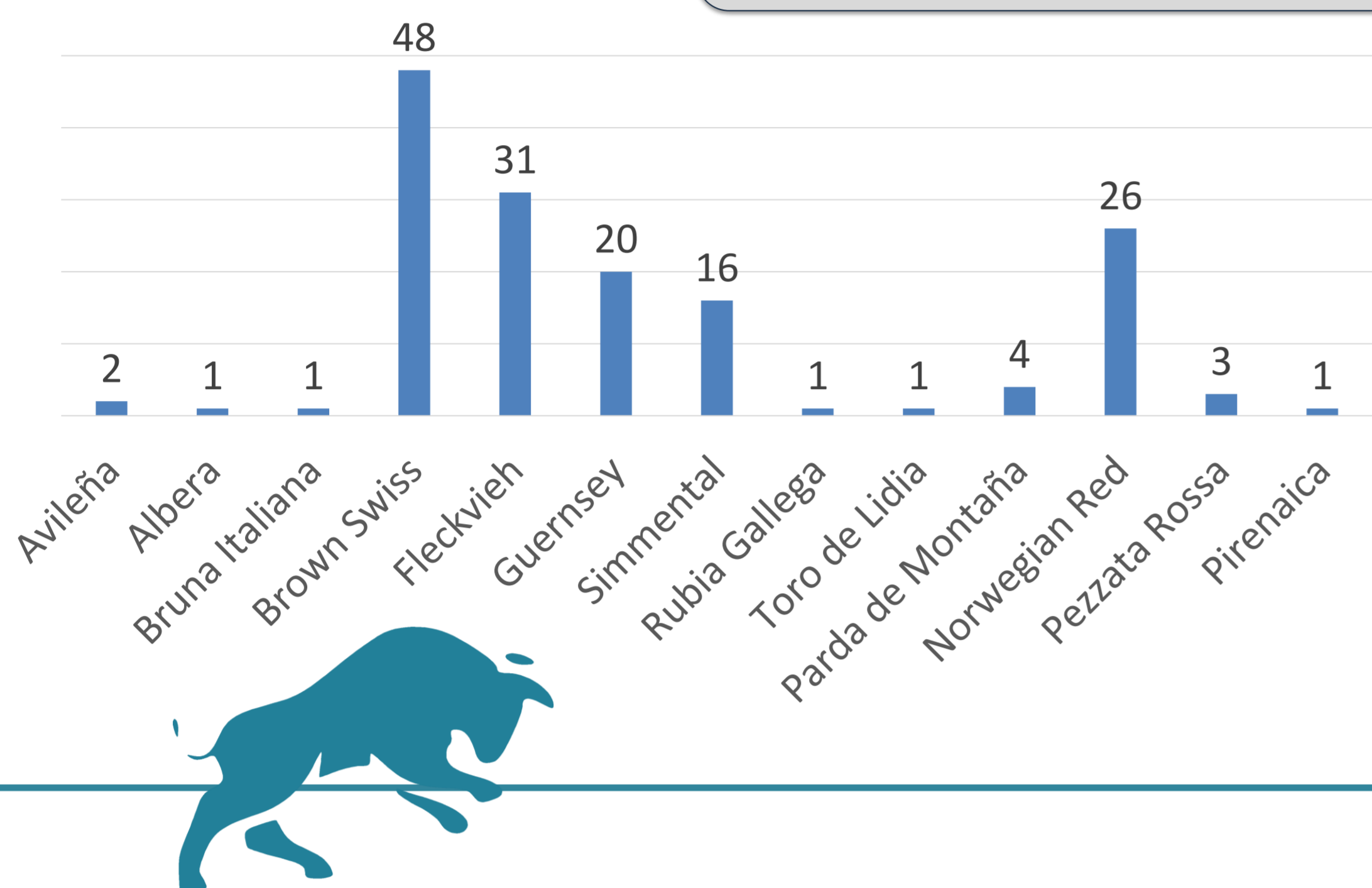


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Data set

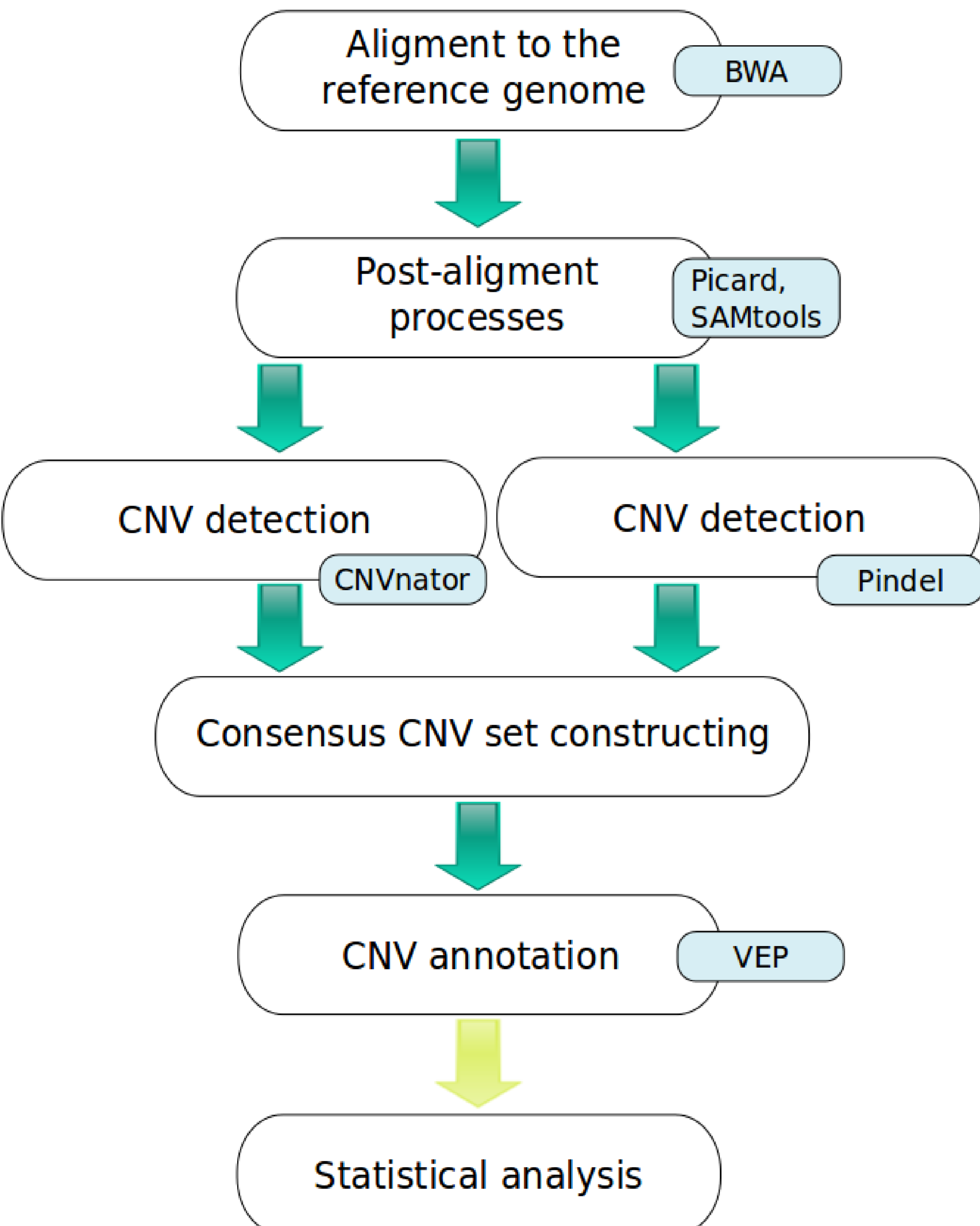


Conclusions

A highly significant variation between all animals was observed both in the number and in the length of CNVs.

The inter-breed variation was also significant and the breed-specific phenomenon was especially strongly emphasized for Fleckvieh breed.

Methods



Results

