

LncRNAs variability in skeletal muscle of Polish Landrace boars



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MATERIAL & METHODS

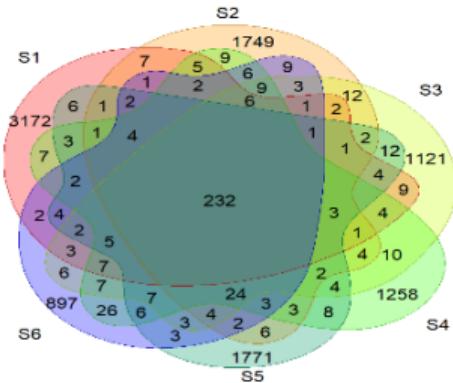
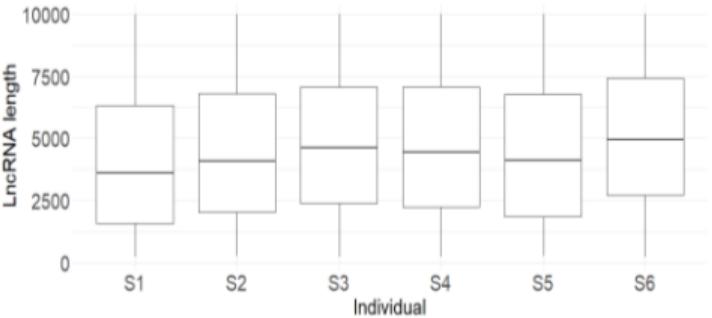
- RNA-seq data
- 6 Polish Landrace boars

Bioinformatics pipeline

- Alignment to *Sscrofa11.1* reference genome
- Quantification of gene expression
- LncRNA filtration

Statistical analysis

- Descriptive statistics
- Statistical tests (χ^2 , Dunn, Kolmogorov-Smirnov, Quade, Wilcoxon, Kruskal-Wallis)



- Individuals significantly varied in LncRNA length, numbers and expression
- No consistent pattern between pairs of half-brothers

