

Discovery and annotation of novel microRNAs in the porcine genome by using a semi-supervised transductive learning approach

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Fig. S1: (A) Receiver Operating Characteristics (ROC) and (B) Precision-Recall (PR) curves computed for each pre-defined imbalance scenario using Ensembl annotation for positive (miRNAs) and negative (other hairpin-like non-coding RNAs) datasets.

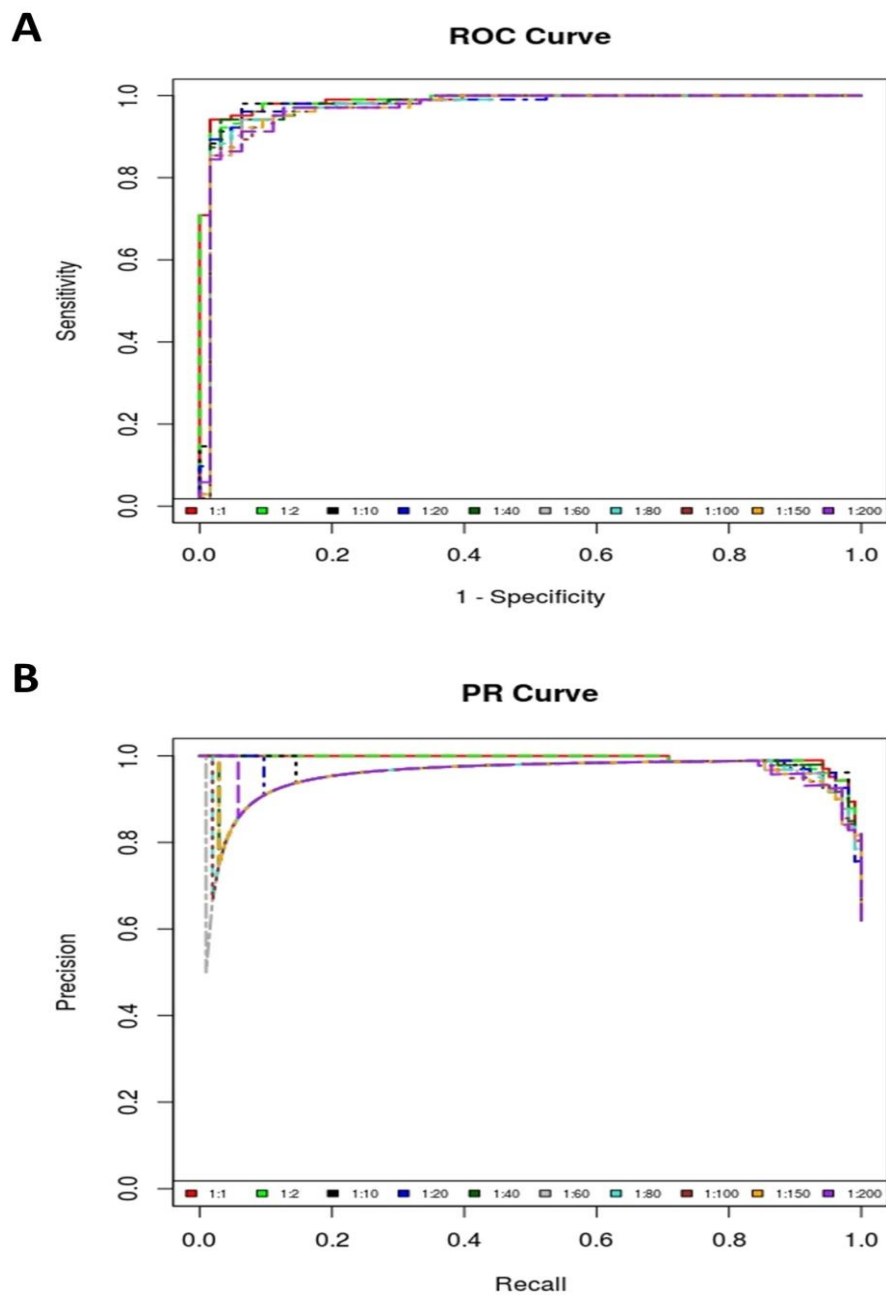


Fig. S2: RT-qPCR of selected novel miRNAs successfully profiled in **(A)** the *longissimus dorsi* skeletal muscle and **(B)** liver tissues from 7 Göttingen minipigs.

