

## Supplementary Information

### **iTRAQ-based proteomic analysis reveals key proteins affecting muscle growth and lipid deposition in pig**

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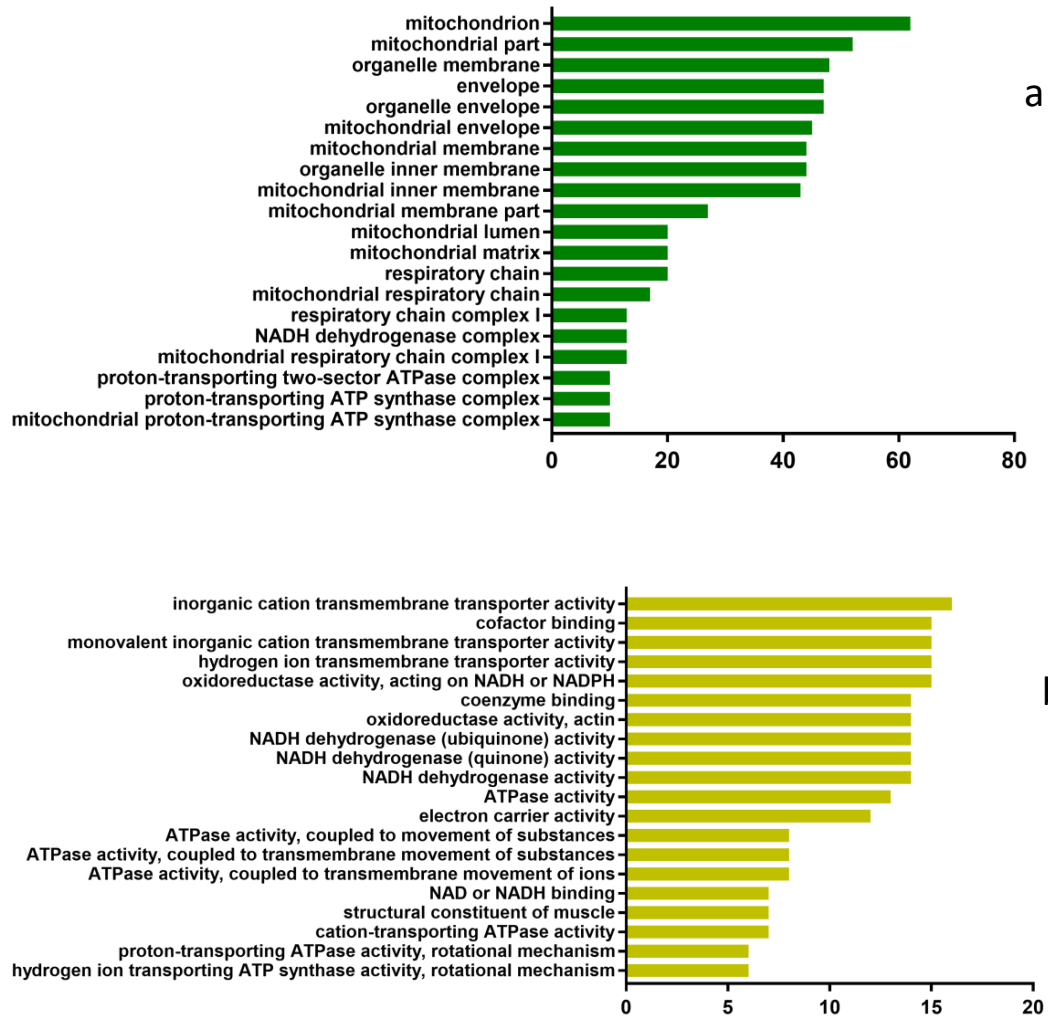
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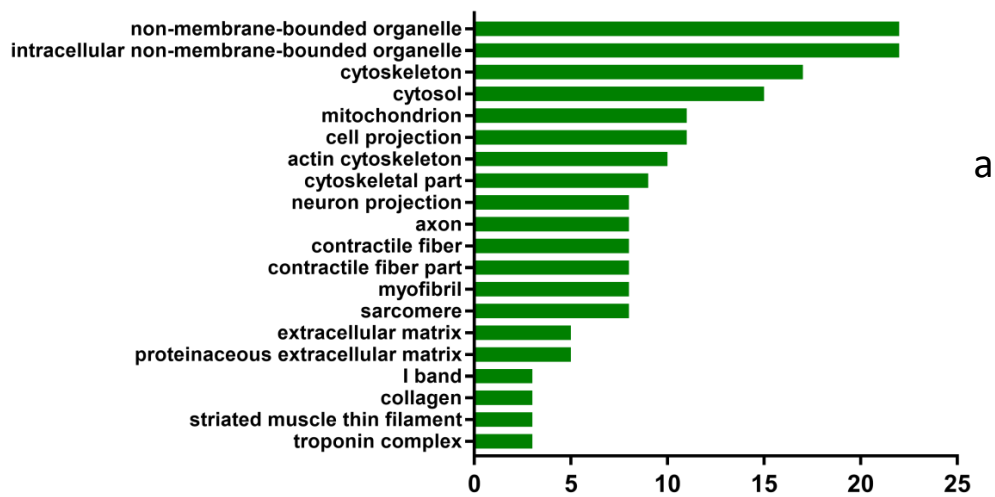
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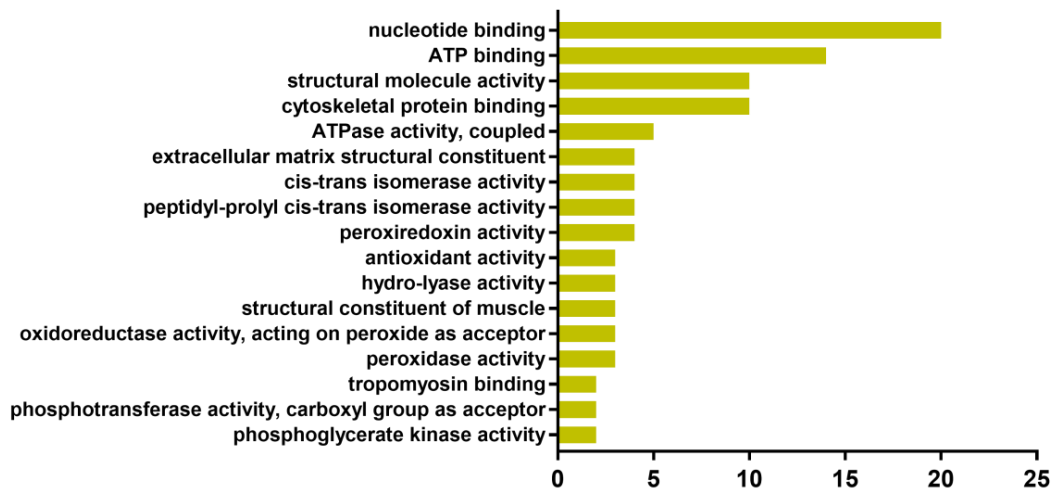
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**Supplementary figure S1. Top terms of cellular component and molecular function of up-regulated DEPs. Panel (a) shows the top terms of cellular component. Panel (b) shows the top terms of molecular function.**

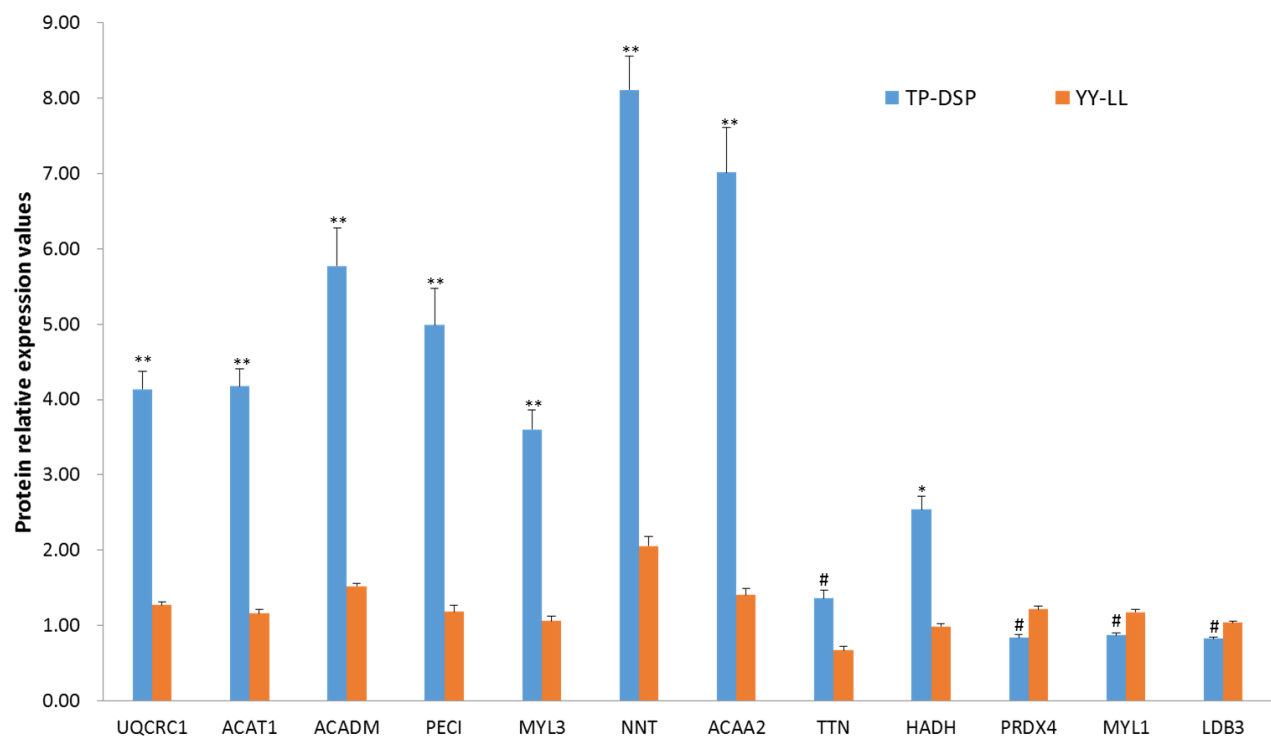


a



b

**Figure S2. Top terms of cellular component and molecular function of down-regulated DEPs. Panel (a) shows the top terms of cellular component. Panel (b) shows the top terms of molecular function.**



**Figure S3. Protein relative expression values in the PRM experiment. \*\* represents  $P < 0.01$ , \* represents  $P < 0.05$ , # represents  $P < 0.1$  (n=4 animals of each breed).**

**Supplementary Table S1. Identified peptides in iTRAQ (xml)**

**Supplementary Table S2. Identified proteins in iTRAQ (xml)**

**Supplementary Table S3. Differentially expressed proteins between TP-DSP and YY-LL groups (xml)**

**Supplementary Table S4. Functional annotation of up-regulated DEPs in the TP-DSP group (xml)**

**Supplementary Table S5. Functional annotation of down-regulated DEPs in the TP-DSP group (xml)**