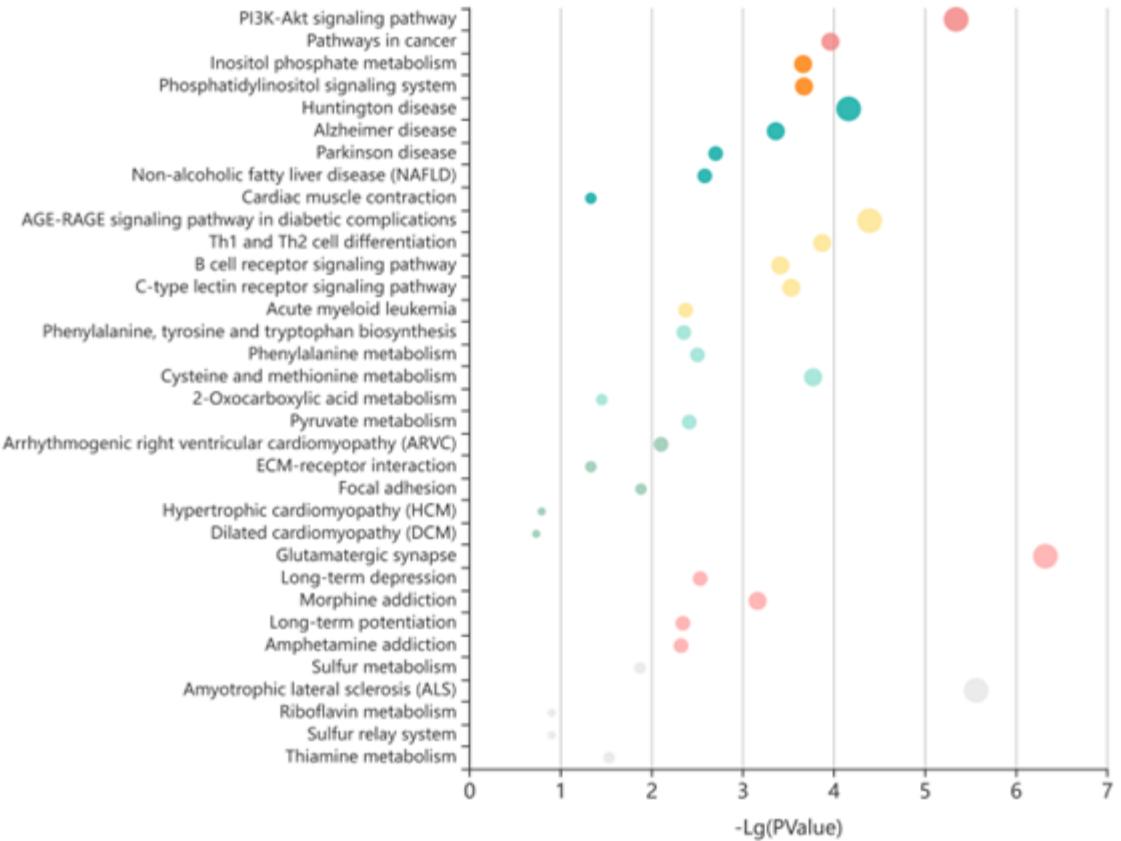
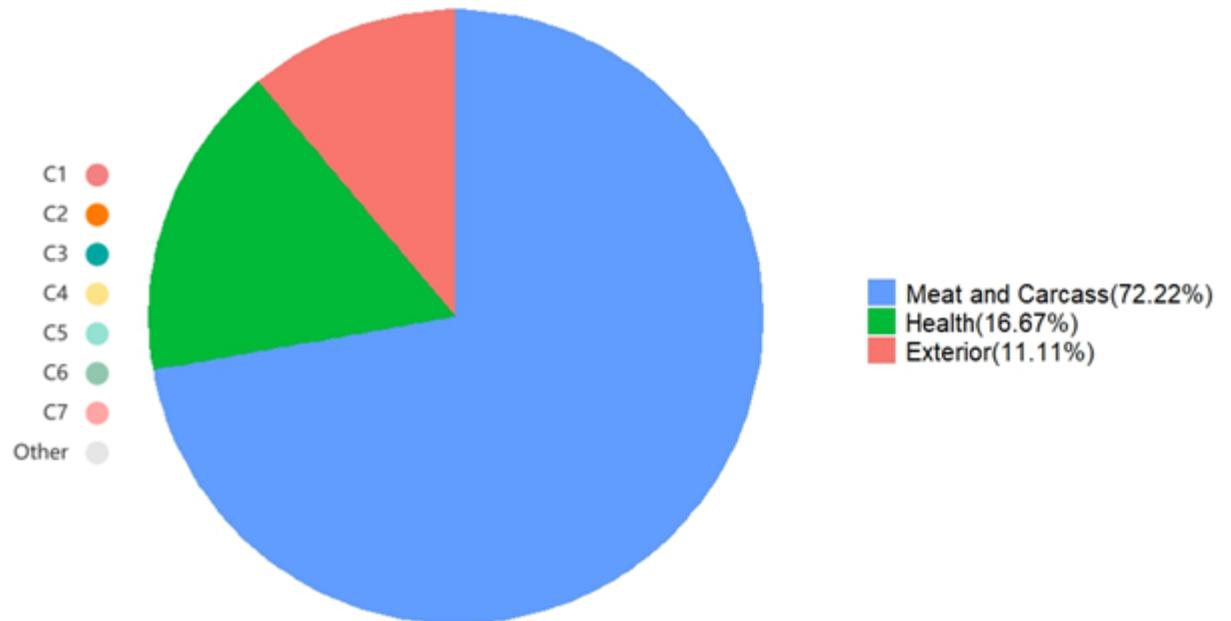


**a****b**

565

566 **Supplementary Material**

567 **Supplementary Figure S1.** Kyoto encyclopedia of genes and genomes (KEGG) and  
568 gene ontology (GO) analysis. **a.** Significant GO terms and KEGG pathways for genes  
569 **b.** Quantitative trait loci enriched results.

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571 **Supplementary Table S1.** p-values for fixed effects

572

Fixed effects	PC1 <sup>b</sup>	PC2 <sup>b</sup>	PC3 <sup>b</sup>	PC4 <sup>b</sup>	PC5 <sup>b</sup>	Slaughter_batch	Live_weight
p_value <sup>1)</sup>	0	7.51175e <sup>-287</sup>	1.34436e <sup>-175</sup>	3.66312e <sup>-176</sup>	5.52682e <sup>-174</sup>	<2e <sup>-16</sup>	<2e <sup>-16</sup>

573 <sup>1)</sup> p-value is calculated by EIGENSTRAT software and the R language b. PC1-PC5 are the five principal components are  
574 calculated using PLINK.

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