

Identification of key ferroptosis-related targets in colorectal cancer: A multiomics study via machine learning and AUcell analysis of single-cell RNA-sequencing

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Fig. S1. Flow chart of this study

Fig. S2. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the AQP8 protein.

Fig. S3. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the NOX4 protein.

Fig. S4. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the NR5A2 protein.

Fig. S5. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the SCD protein.

Fig. S6. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the TIMP1 protein.

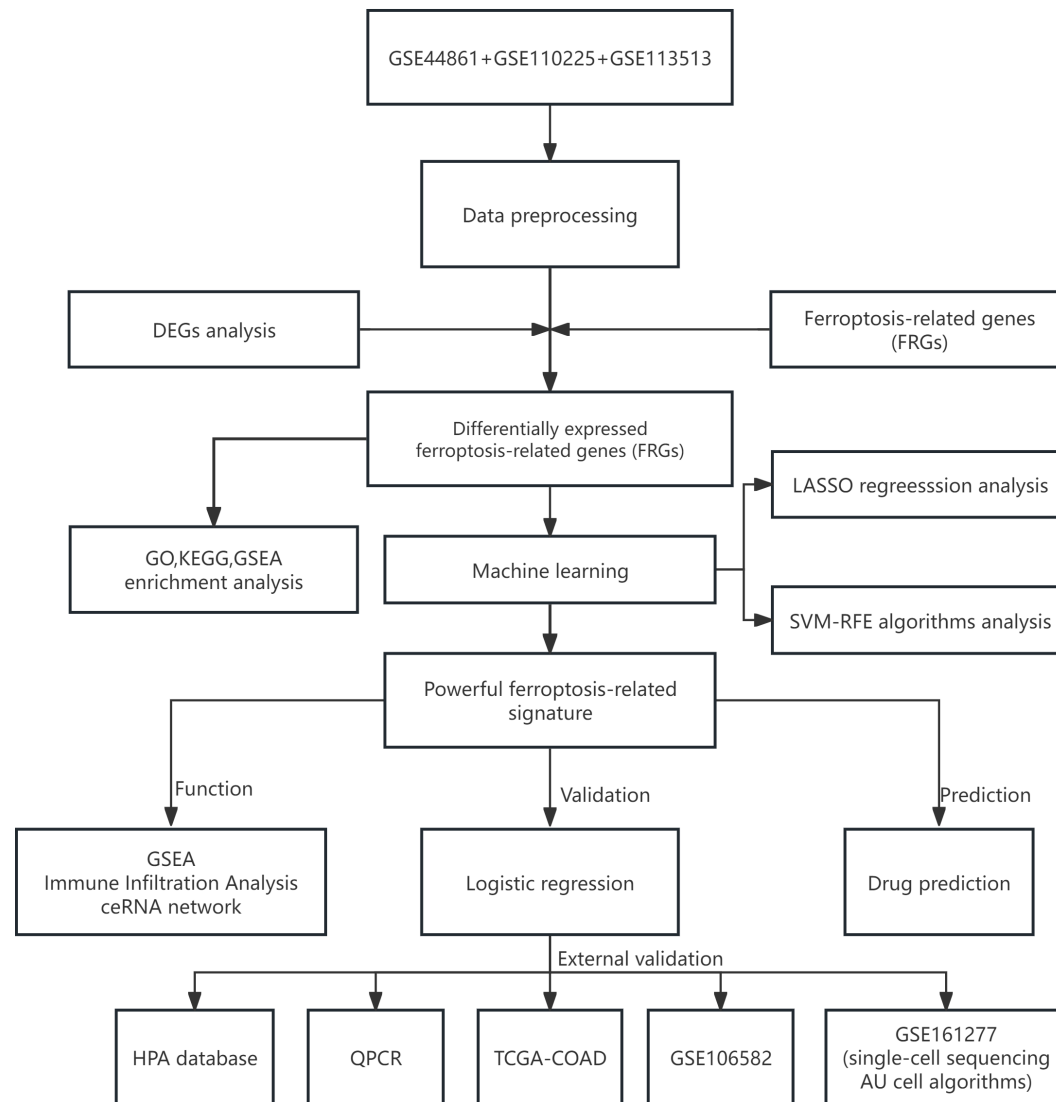


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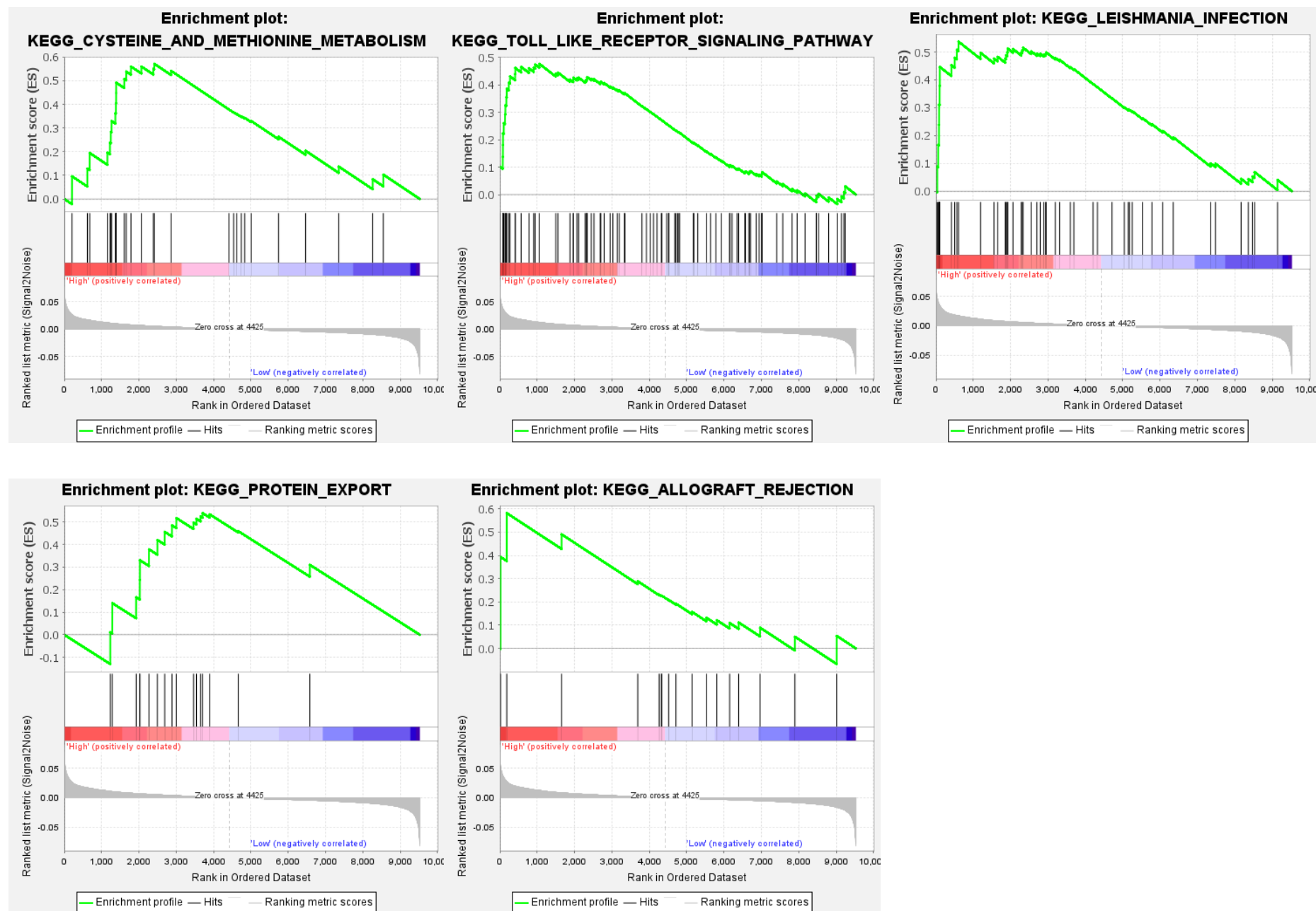


Fig. S2. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the AQP8 protein (An examination of the top five routes using).

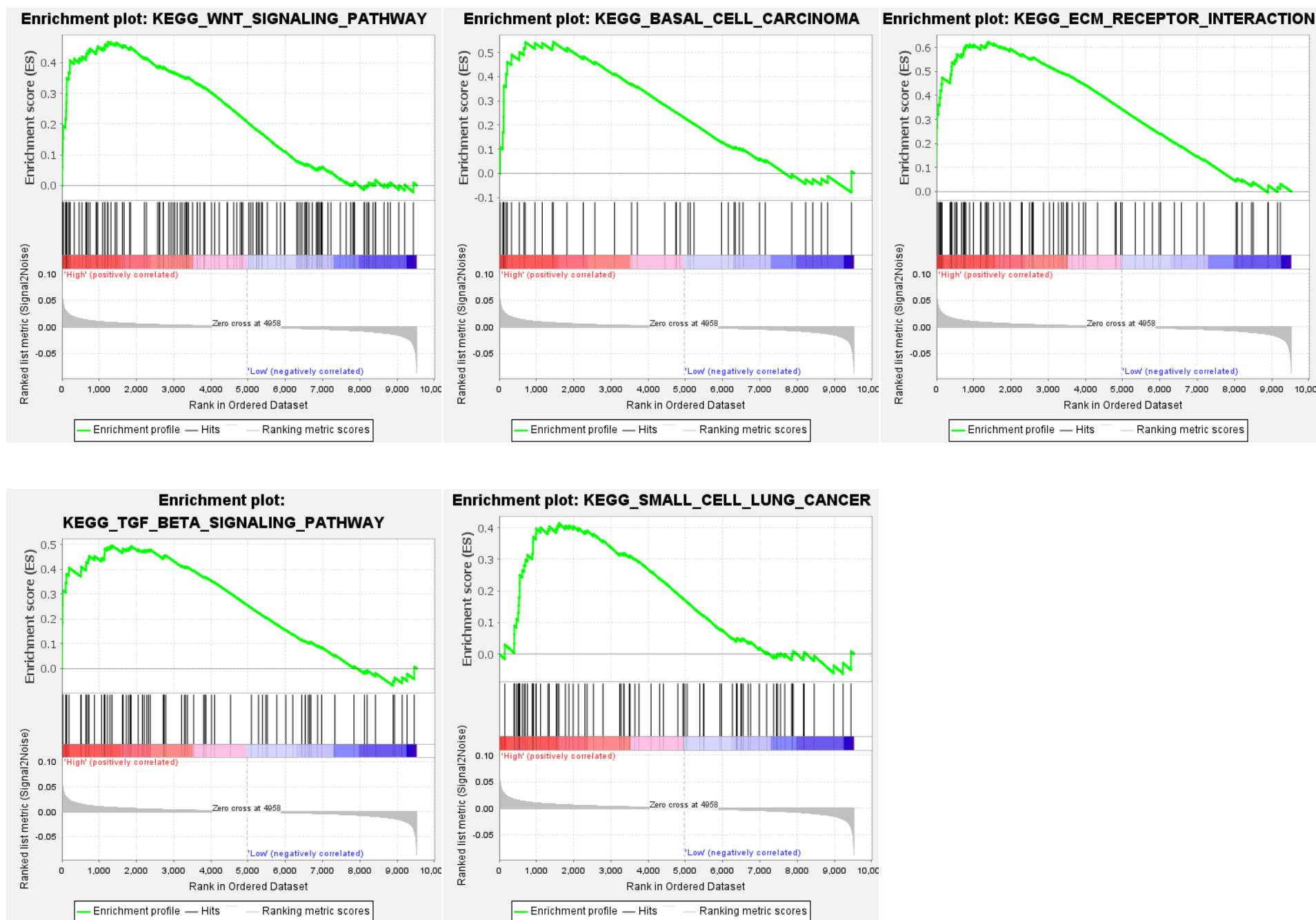


Fig. S3. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the NOX4 protein (An examination of the top five routes using).

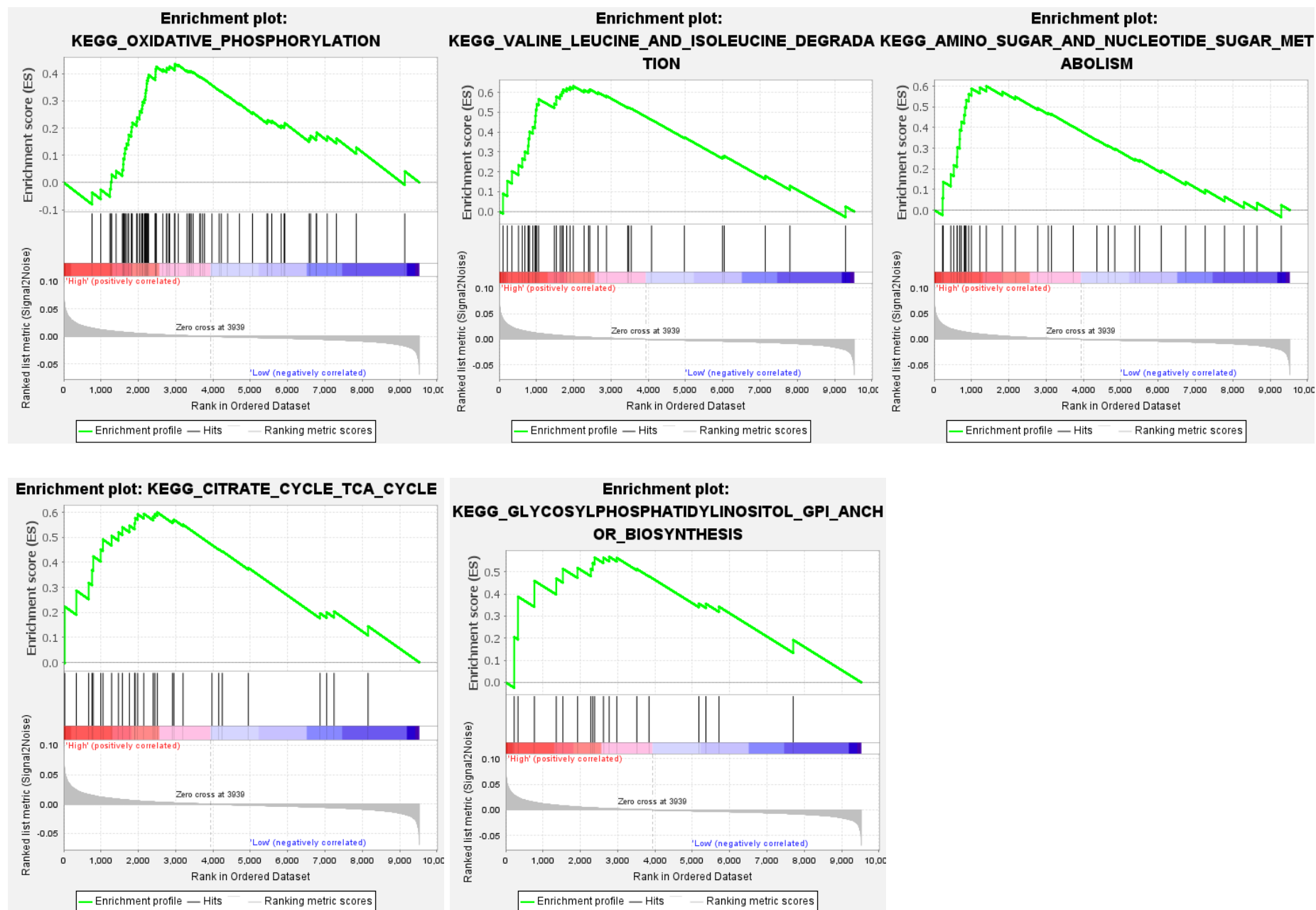


Fig. S4. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the NR5A2 protein (An examination of the top five routes using).

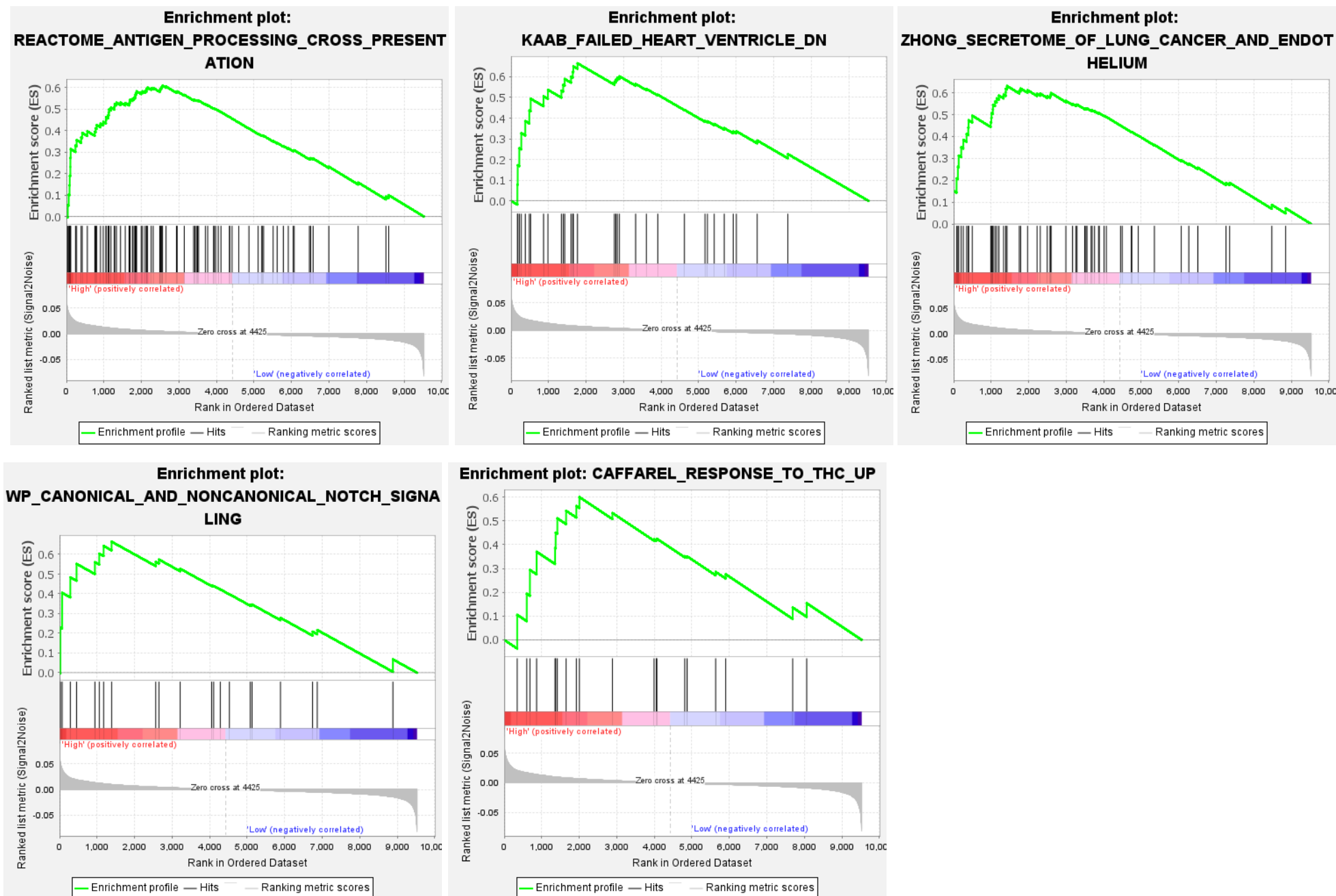


Fig. S5. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the SCD protein (An examination of the top five routes using).

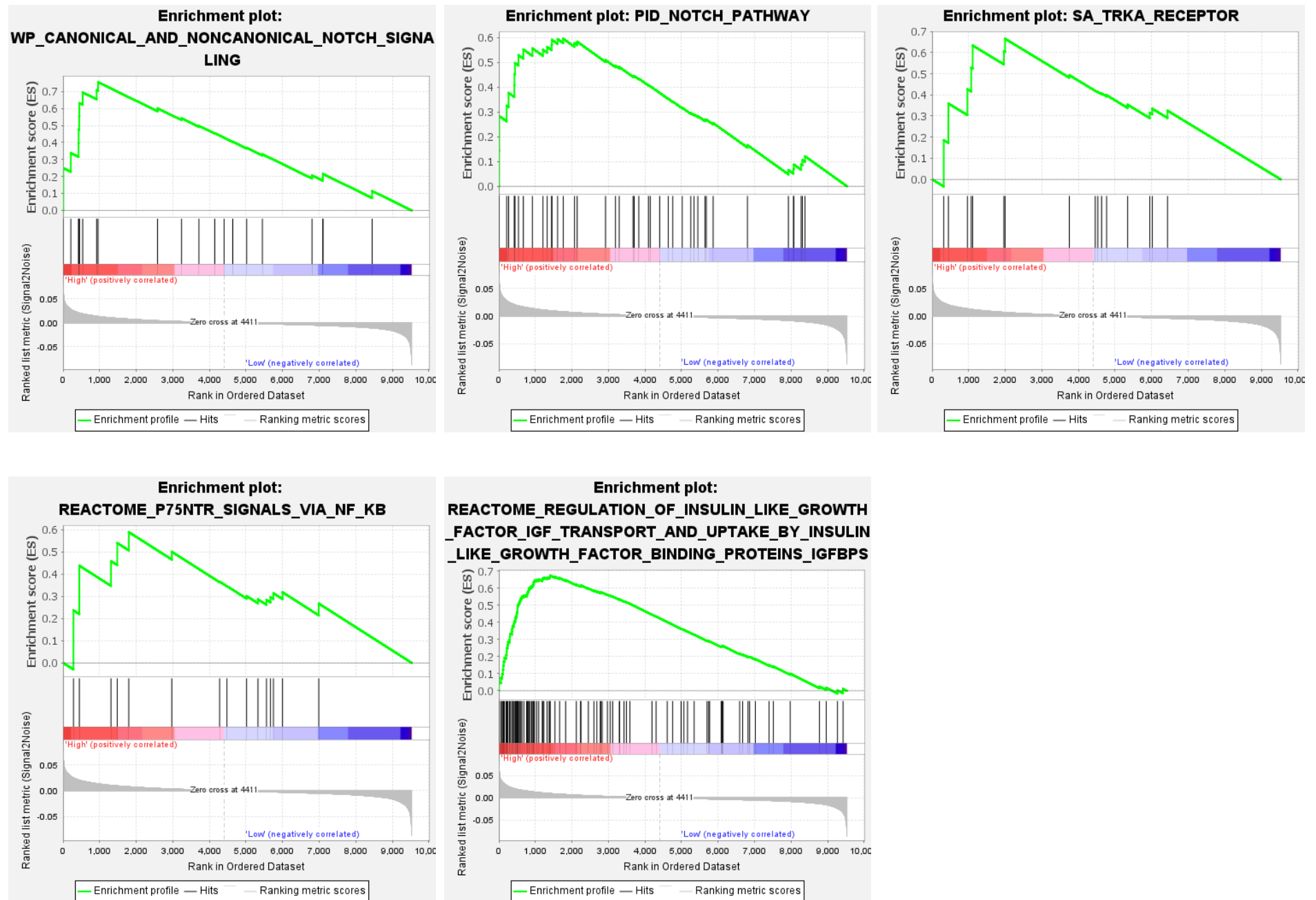


Fig. S6. The results of Gene Set Enrichment Analysis - Kyoto Encyclopedia of Genes and Genomes (GSEA-KEGG) pathway analyses for the TIMP1 protein (An examination of the top five routes using).