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Correction: Construction and validation of a prognostic model of angiogenesis-related genes in multiple myeloma

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Following publication of the original article [1], it was reported that Fig. 6 is a duplicate of Fig. 5. This is a result of a typographical error. The correct Fig. 6 is given below. Further to this, in the original article [1], only one corresponding author was erroneously listed. Xuezhong Gu and Zhixiang Lu are both corresponding authors. The original article [1] has been updated.

 ${}^{\dagger}\text{Rui}$ Hu and Fengyu Chen contributed equally to this work.

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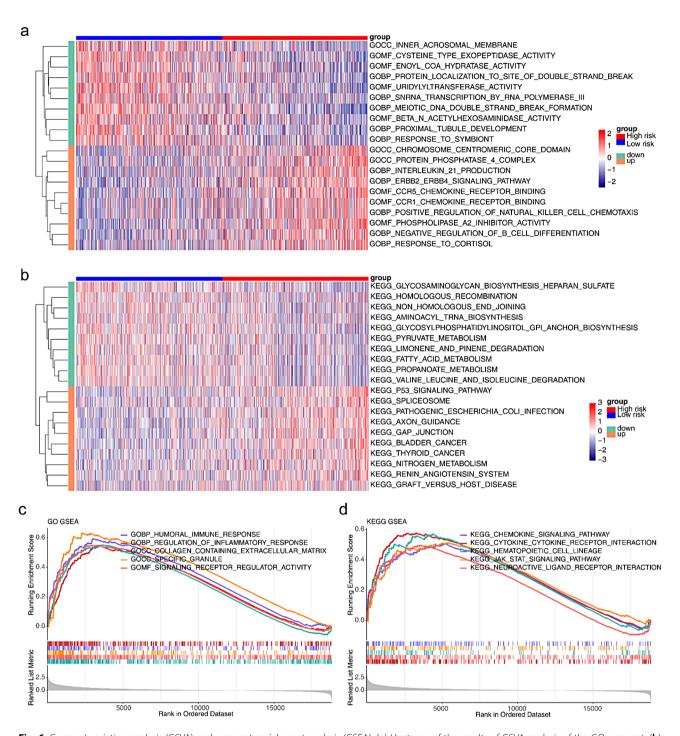


Fig. 6 Gene set variation analysis (GSVA) and gene set enrichment analysis (GSEA). (a) Heatmap of the results of GSVA analysis of the GO gene set. (b) Heatmap of the results of GSVA analysis of the KEGG gene set. (c) GSEA results for the GO gene set. (d) GSEA results for the KEGG gene set. GO, Gene Ontology; KEGG, Kyoto Encyclopedia of Genes and Genomes

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