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Corrigendum: Immunogenomic landscape in breast cancer reveals immunotherapeutically relevant gene signatures

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A Corrigendum on

Immunogenomic landscape in breast cancer reveals immunotherapeutically relevant gene signatures

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In the published article, there was an error in the IHC images of Figure 5B as published. Four repeat IHC images were displayed cursorily. The corrected Figure 5B and its caption “Construction of the immune biological signature. (A) DEGs were used to construct the immune biological signature. (B) Representative immunohistochemical images of infiltrated immune cells in Bengbu cohort between high- and low- ITBscoore groups. (C) Kaplan-Meier curves for patients in the BRCA cohort divided into high and low ITBscoore subgroups. (D) Alluvial diagram indicating immune subtypes in groups with different BRCA subtypes (basal,

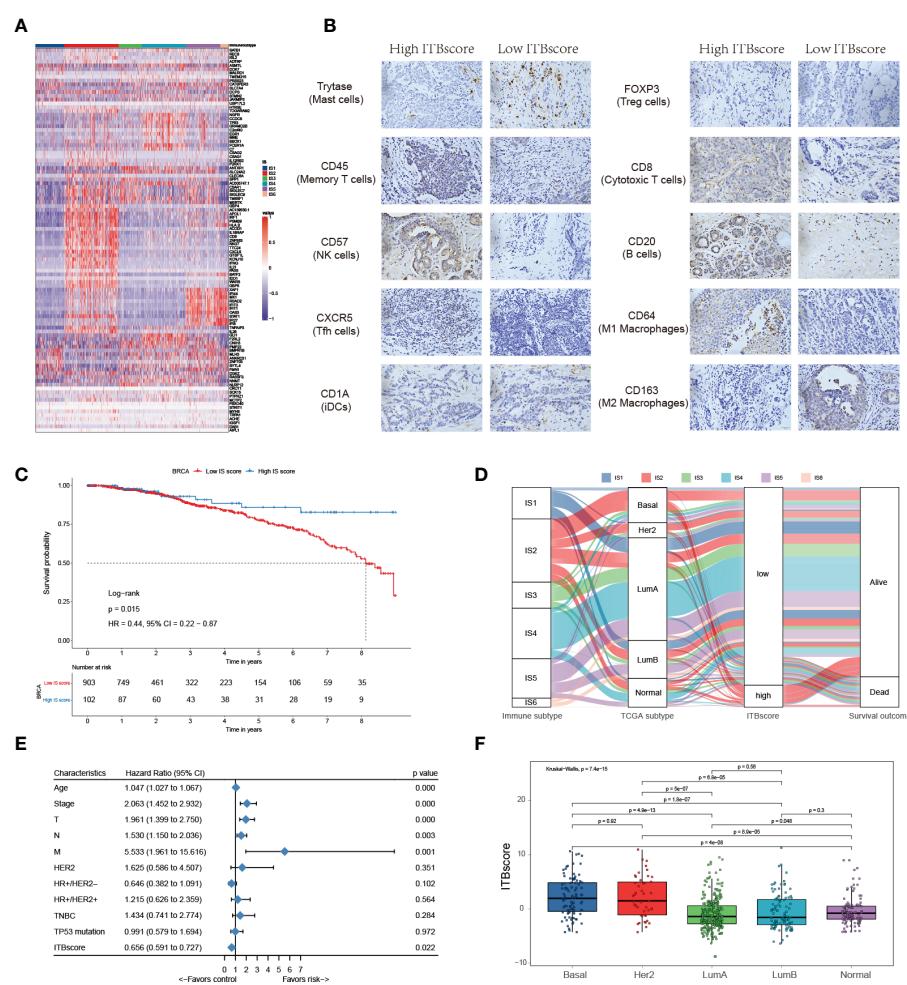


FIGURE 1

Construction of the immune biological signature. **(A)** DEGs were used to construct the immune biological signature. **(B)** Representative immunohistochemical images of infiltrated immune cells in Bengbu cohort between high- and low- ITBscore groups. **(D)** Alluvial diagram indicating immune subtypes in groups with different BRCA subtypes (basal, Her2, LumA, LumB, and normal), ITBscores, and survival outcomes. **(E)** Prognostic value of the ITBscore and classic clinicopathological covariates in the high/low ITBscore subgroups. **(F)** Distribution of the ITBscore among TCGA-BRCA molecular subtypes. “ appear below.

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The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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