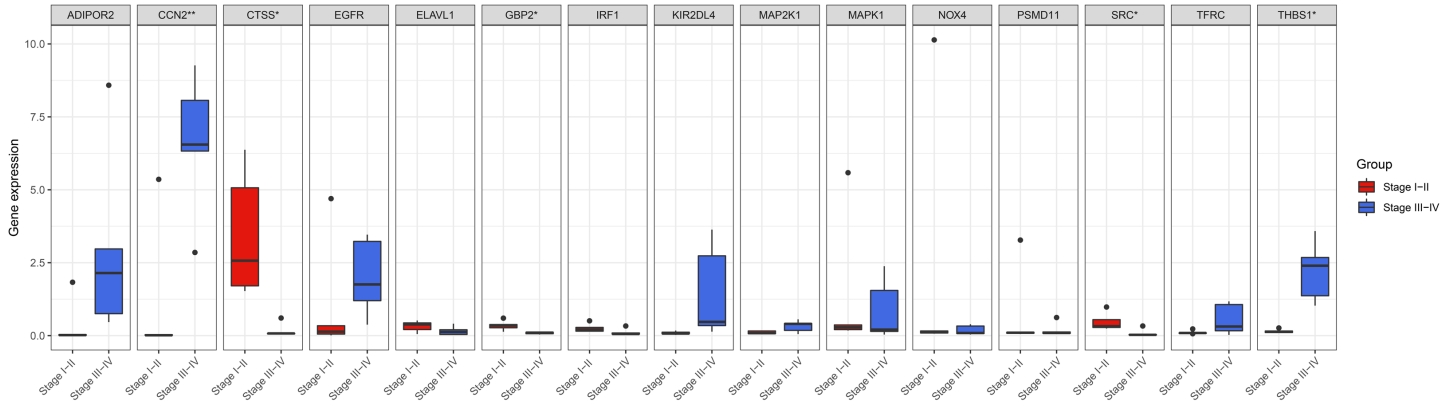
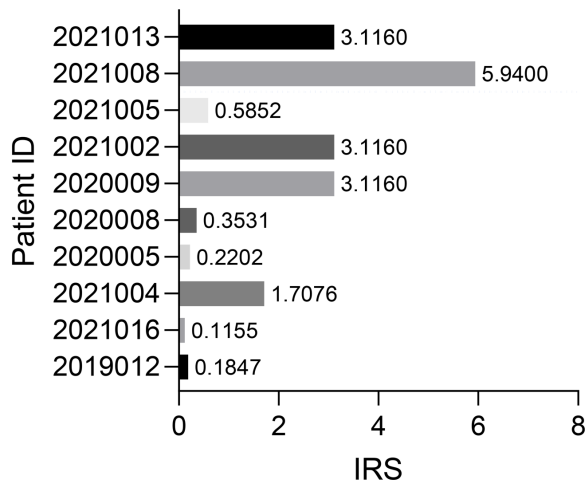
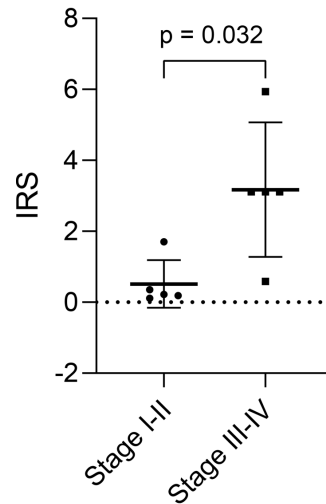


a**b****c**

Supplementary Figure 5 Validation of the prognostic value of IRS in the local cohort. (a) The expression level of the 15 genes in the risk signature in the subjects with BCa in stage I-II and the subjects with BCa in stage III-IV. (b) The calculated IRS of each patient. (c) The patients in stage III-IV exhibited higher IRS level compared with those in stage I-II.