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Clinical implications of TMB in GC

Dong-yan Zhao, Xi-zhen Sun, Shu-Kun Yao

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TMB has been described as a powerful predictor of **tumor** behavior and response to immunotherapy. Methods: A total of 443 bladder **cancer** samples obtained from **The Cancer Genome Atlas** (TCGA) were analyzed for **mutation** types, TMB values, and prognostic value of TMB. Differentially expressed genes (DEGs) were identified from the TMB groupings.

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