Supplementary Materials: Establishment and Validation of an Individualized Cell Cycle Process-Related Gene Signature to Predict Cancer-Specific Survival in Patients with Bladder Cancer

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Figure S1. A detailed screening diagram to establish a robust gene signature for CSS.



Sample Cluster Dendrogram

Figure S2. Sample clustering showed one outlier was detected and excluded in WGCNA.



Figure S3. A power of β = 9 was chosen as the optimal soft threshold to ensure a scale-free coexpression network.



Figure S4. Scatter diagram showed a highly positive correlation between GS and MM in the brown module (r = 0.87, $p < 1 \times 10^{-200}$).



Figure S5. Significant difference of overall survival in TCGA MIBC patients (*P*=0.0002) was observed among different risk subgroups defined by the decision tree.



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