

Identifying prognostic biomarkers based on aberrant DNA methylation in kidney renal clear cell carcinoma

SUPPLEMENTARY TABLES

Supplementary Table S1: Subpathways enriched by differentially methylated genes

See Supplementary File 1

Supplementary Table S2: Subpathways enriched by differentially expressed genes

See Supplementary File 2

Supplementary Table S3: subpathways under the regulation of both DMGs and DEGs

See Supplementary File 3