

Employing an orthotopic model to study the role of epithelial-mesenchymal transition in bladder cancer metastasis

Supplementary Materials

Supplementary Table S1: Differentially expressed genes in CTCs versus the other tumor sites with $FDR < 0.05$, $p < 0.001$, and fold-change ≥ 1.5 using the class comparison tool within BRB array tools. See Supplementary_Table_S1