

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: RNA sequencing analysis of mRNA abundance in uninfected kidney cells (control), the total population (total) or sphere-selected population (spheres) of kidney cells after infection with lentivirus expressing shTsc2/shCdkn2a.

File Name: Supplementary Data 2

Description: Gene Set Enrichment Analysis: shTsc2/shCdkn2a total versus control, all genes >8 fold upregulated (154 identifiers converted to 126 Entrez genes).