

## Figure 4-figure supplement 4.

(A) qRT-qPCR was performed from LS180 cells transfected for 48 hr with control siRNAs (siCTL) or BRI3BP siRNAs Smartpool (siBRI3BP). *GAPDH* was to normalize the data and UBC was used as negative control. (B) PI staining and FACS analysis was performed from LS180 cells transfected for 48 hr with control siRNAs (siCTL) or BRI3BP siRNAs and then treated with DTT for 2 hr. Error bars represent SD from 3 experiments. \*\*p<0.005.