



Figure S14: Significance test of genome-wide somatic amplification and LOH for urothelial cancer samples.

This figure illustrates statistical significance of somatic LOH (a) and amplification (b) provided by GIANT. The germline aberrations are excluded (such as LOH in chromosome 9p12-11 and 9q12 in sample UC288, as shown in Figure 7), and only somatic aberrations are reserved for further analysis. Statistical significance was evaluated by permutation test with pre-determined threshold (dashed line) for q -value.