



Figure S5. PCR amplification efficiency at the untreated and UNG-treated telomeric and *36B4* loci. qPCR amplification efficiency in the untreated/treated samples was normalized to the average expression level for each target genomic region. Since no significant differences between BRCA groups were found, all FBOC cases were grouped together (n=328). Bars show the mean \pm standard error of the mean (SEM). Unpaired *t*-tests were performed for statistical significance. a. u.: arbitrary units.