



Figure S5. Differentially Expressed Genes between HIR and QT subgroups of HCC in three cohorts

(A) Venn Diagram of genes whose expression levels are significantly different between the HIR and QT subgroups of non-tumor surrounding tissues from patients with HCC (cohorts 1, 2, and 3). A P value of < 0.001 was required for a gene to be retained.

(B) Expression patterns of the 177 up- or down-regulated in all three cohorts. Colored bars at the top of the heat map represent samples as indicated.