

Supplementary Figure 1: Unsupervised clustering analysis using an irinotecan metabolism pathway gene expression signature of 5 colorectal cancer tumorgrafts treated with irinotecan. The 3 susceptible tumor clustered together (CCR-004, CCR-003H, POS 0805) and were characterized by lower expression of gene involved in the intracellular catabolism of irinotecan and SN38 such as UGT1A1 and CYP3A4 and 5.

