Figure S8. TCF-4 occupies the *PDK* and *MCT1* promoters. ChIP-seq data compiled from the UCSC Genome Browser (assembly Feb. 2009, hg19; [http://genome.ucsc.edu](http://genome.ucsc.edu)\(^{76}\)), showing TCF-4 occupancy in four different cell lines over the promoter regions of *PDK1*, *PDK2*, *PDK3*, *PDK4*, and *MCT1*. These data were generated and analyzed by the labs of Michael Snyder at Stanford University; Mark Gerstein and Sherman Weissman at Yale University; Peggy Farnham at University of Southern California; and Kevin Struhl at Harvard. RT-qPCR analysis of MCT-1 and GLUT1 was performed on RNA prepared from freshly-extracted xenograft tumors. RT-qPCR data shown are representative of two replicate tumor sets. Error bars represent the standard error between three internal replicates.