



Supplementary Fig. S3. mNC-derived cell lines share transcriptional profiles with tumor tissues that are characterized by MYC pathway upregulation. Heatmaps plotting the adjusted p-value from GSEA (Hallmark or Genome Ontology (GO) gene sets) of ranked log fold-change values that are corrected for standard-error (Wald statistic). up, differentially upregulated genes; down, differentially downregulated genes.