

Supplementary Figure 3. Validation of INPP5F methylation in DNMT3A KO cells. Methylation percentages for cg10053779, located within the INPP5F gene body, are shown for non-targeting control (NT) and DNMT3A knockout HEC1B cells. Each point represents an independent replicate, and horizontal bars indicate group means. Differential methylation was assessed using limma with Benjamini–Hochberg FDR correction.