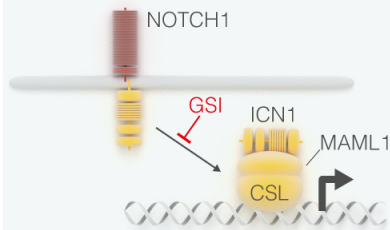
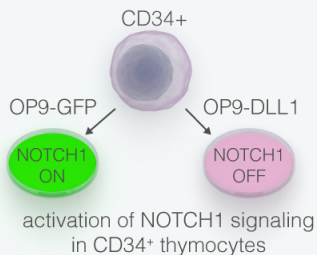


The Notch driven long non-coding RNA repertoire in T-cell acute lymphoblastic leukemia

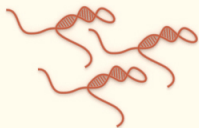
RNA-sequencing



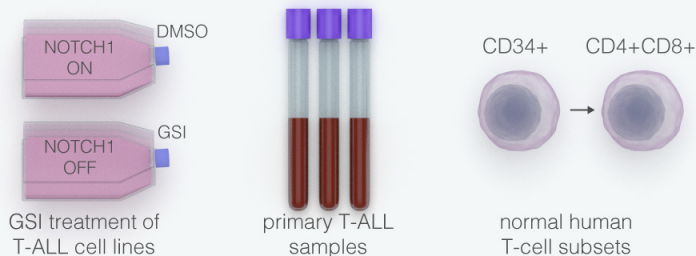
GSI treatment of T-ALL cell line CUTLL1



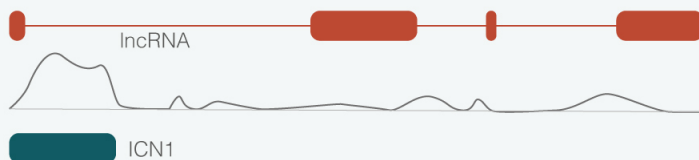
selection of NOTCH regulated lncRNAs



micro-array expression data



ICN1 ChIP-Seq data



functional annotation through guilt-by-association

