Mutational Landscape of T-ALL Samples 10 -0 5 10 15 20 CDKN2A 50% NOTCH1 PHF6 CDKN2B 38% MED12 29% WT1 18% PTEN 9% DNM2 21% EZH2 18% JAK1 18% 18% CHD4 **Alterations** 18% JAK3 Missense Nonsense CTCF 15% Frameshift Splice_Site NRAS 15% Other Amplification
Hemizygous_Deletion
Homozygous_Deletion KAT6A 15% 9% FLT3 Structural_Variant AHNAK2 12% TYRO3 12% CENPB 12% 12% TLX3 CNOT1 9% ZNF99 9% CCND3 9% **ACAN** 9% FBXW7 9% TAL1 9% 6% LMO2