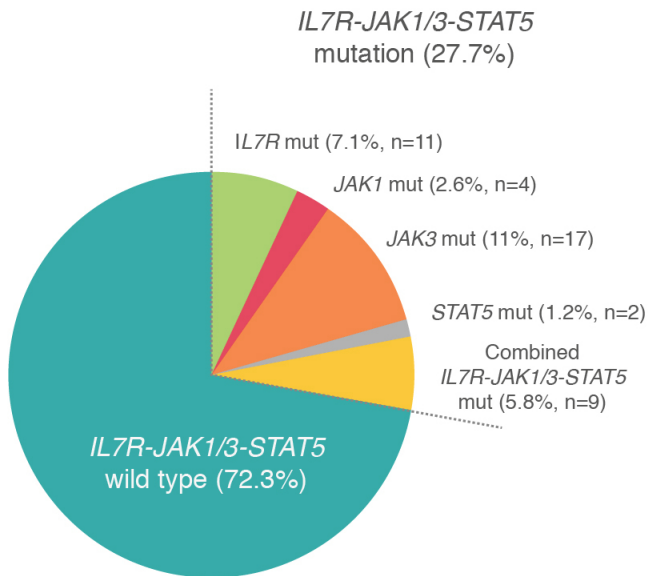


The *IL7R-JAK* pathway is mutated in 27.7% of T-ALL



Mutations in *IL7R-JAK* genes and epigenetic modulators are significantly associated

