



907 bulk large B cell lymphoma (LBCL) transcriptomes



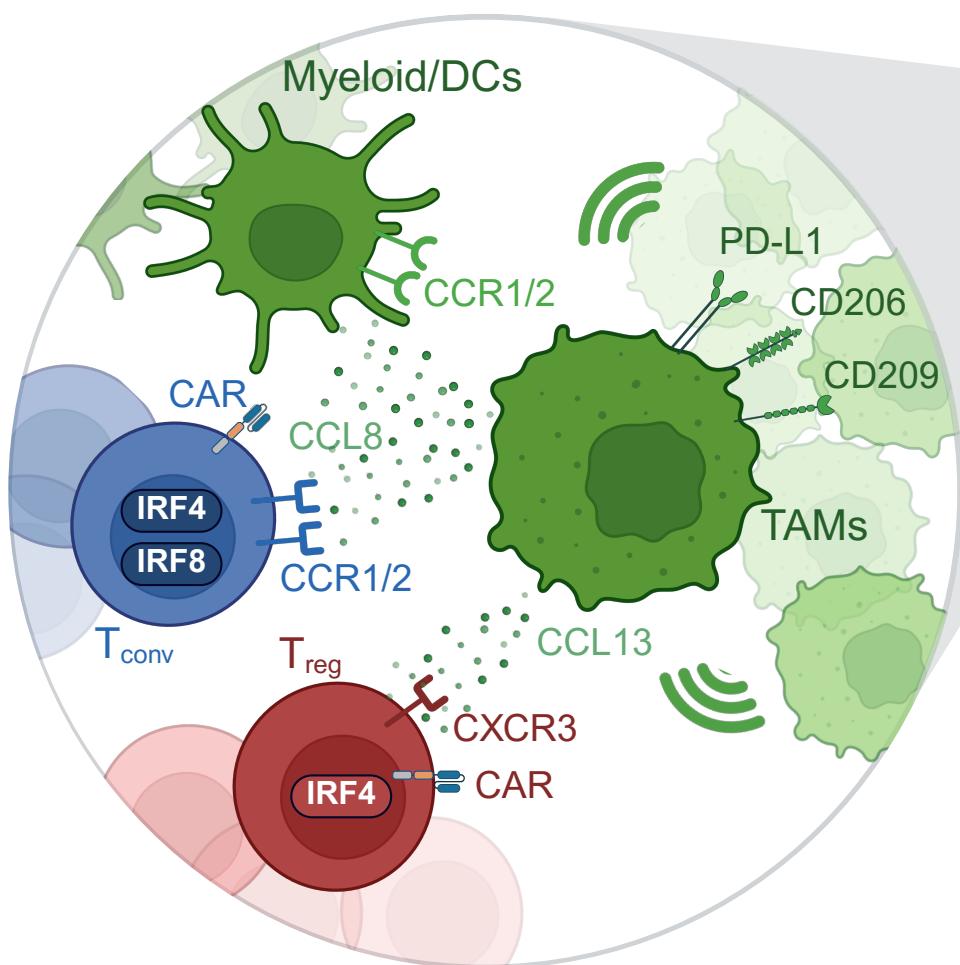
23,670 single cells from 18 on-treatment CAR T TME samples

15,935 single cells from 4 baseline LBCL samples

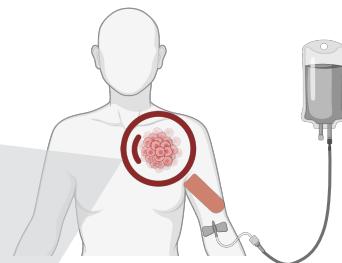


Discovery and validation of the CAR T tumor microenvironment

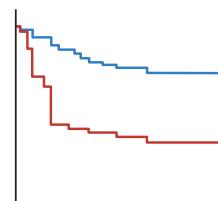
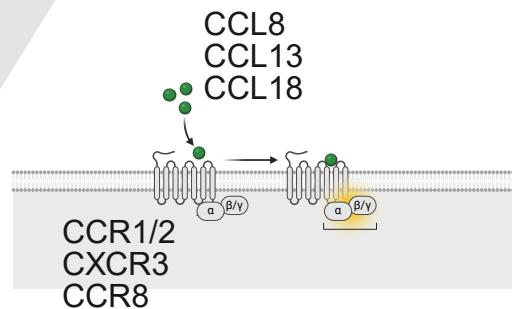
The CAR T TME of LBCL



18 LBCL patients receiving CD19-CAR or bispecific CD19/CD22-CAR therapy



Cell-cell interactomes identify ligand-receptor pairs in the post-CAR TME



Overexpression leads to poor outcomes after (R)-CHOP and CAR T therapies