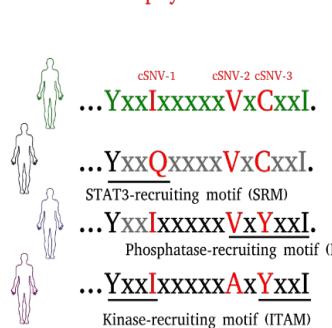
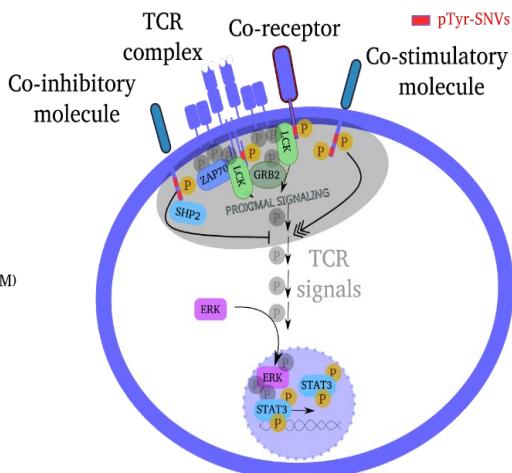


A

pTyr-motif altering SNPs
(pTyr-SNVs)

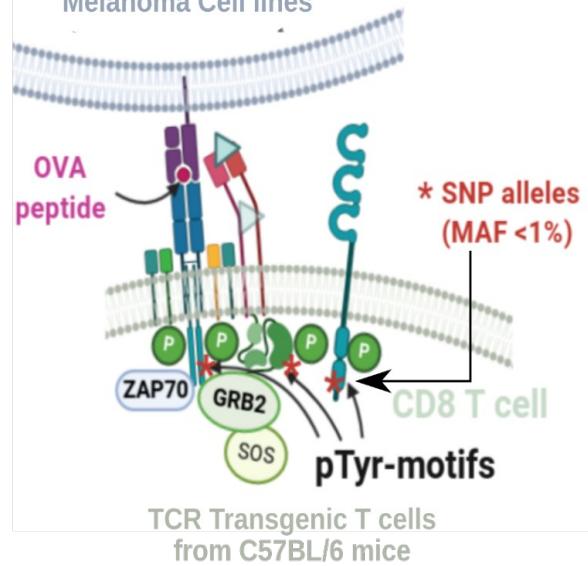


B

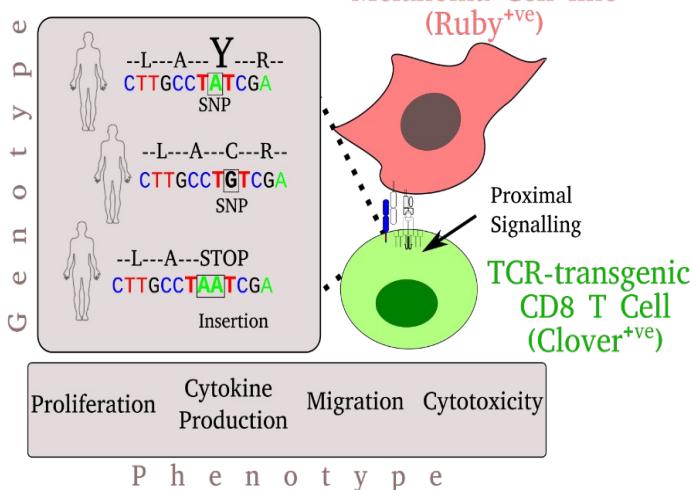


C

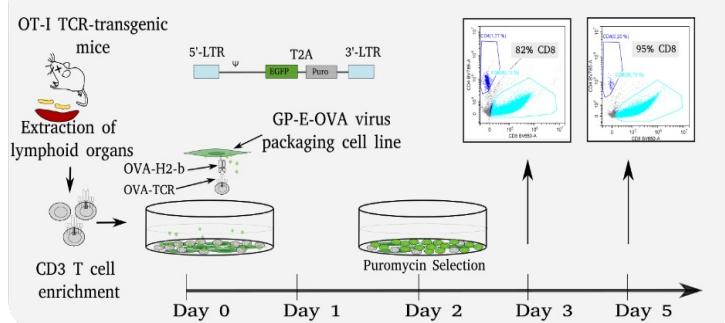
C57BL/6
Melanoma Cell lines



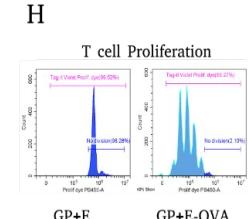
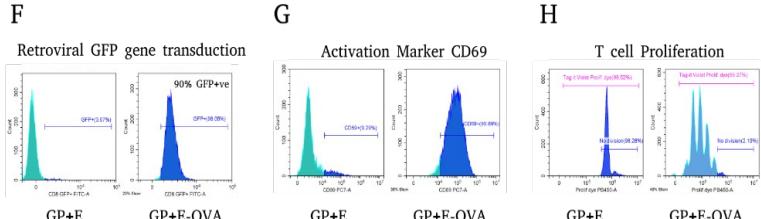
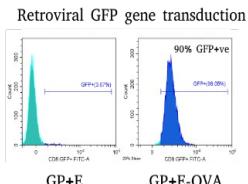
D



E



F



Online Figure-7. A new methodology to elucidate molecular functions of human pTyr-SNVs. (A) Rare germline pTyr-SNVs enriched in cancer cohorts are identified, (B) Example of rare coding region SNPs (cSNPs) and allelic variants which create proximal signalling motifs unique to an individual. Such cSNPs are referred to as pTyr-SNVs (C) TCR-induced proximal signalling pathways and pTyr-SNV sites either creating or destroying a pTyr-motif (D) Genotype-to-Phenotype study using B16-F10-OVA and TCR transgenic CD8 T cells (E) Generation of pTyr-SNV-expressing OT-I CD8 T cells within 5-days via co-cultivation of primary OT-I CD3 T cells and GP-E-OVA retrovirus producing packaging cell lines (F) Percentage of retroviral transduction (G & H) Selective activation of OT-I CD8 T cells (Figure taken from Ulaganathan VK et al., J. Genet Genomics, 2023, doi: 10.1016/j.jgg.2023.01.001).