

FIG. 1

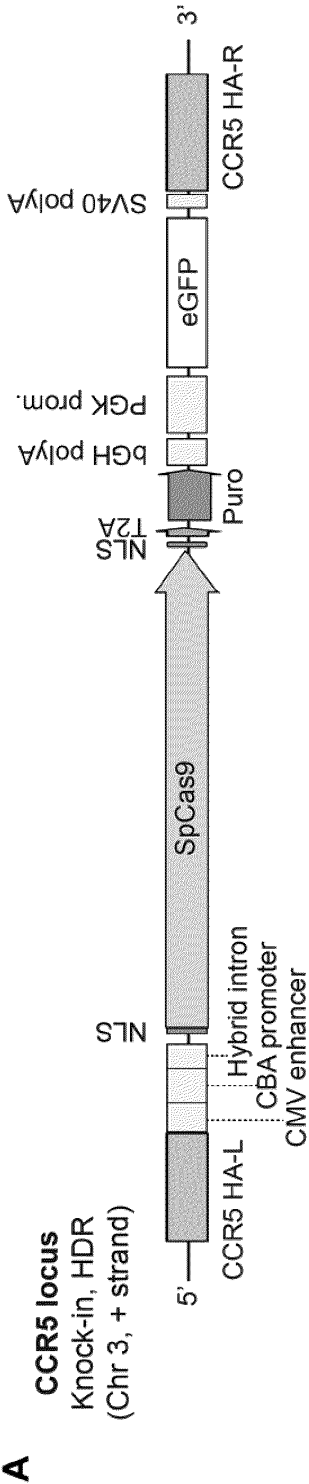
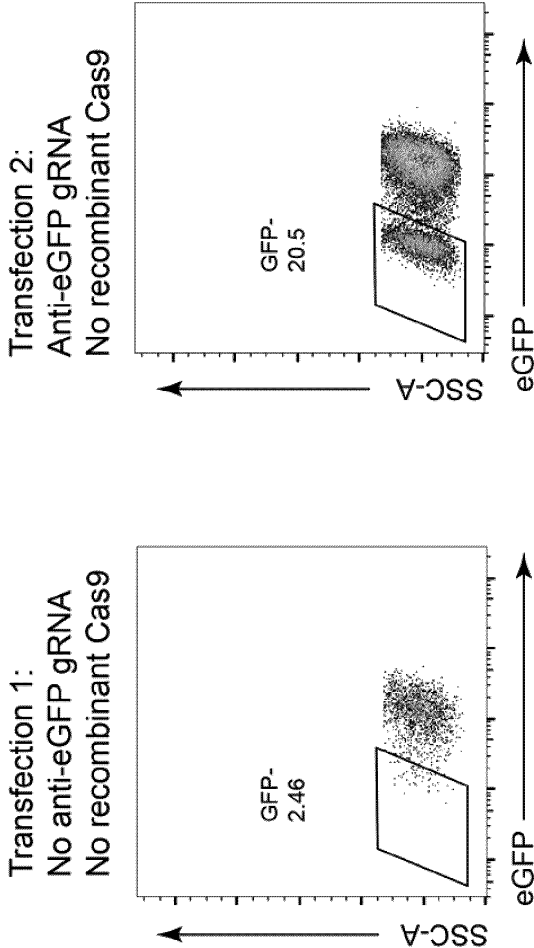


FIG. 2A&B

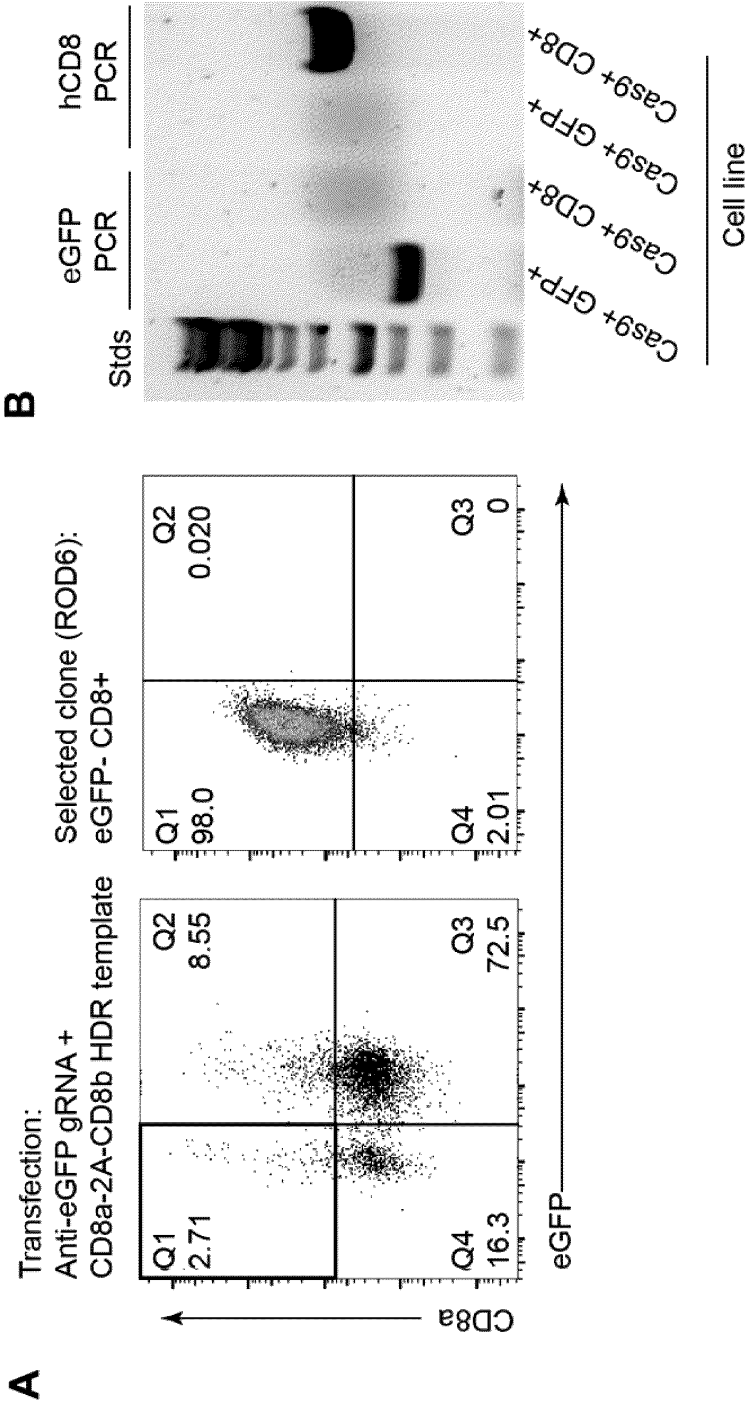


FIG. 2 C

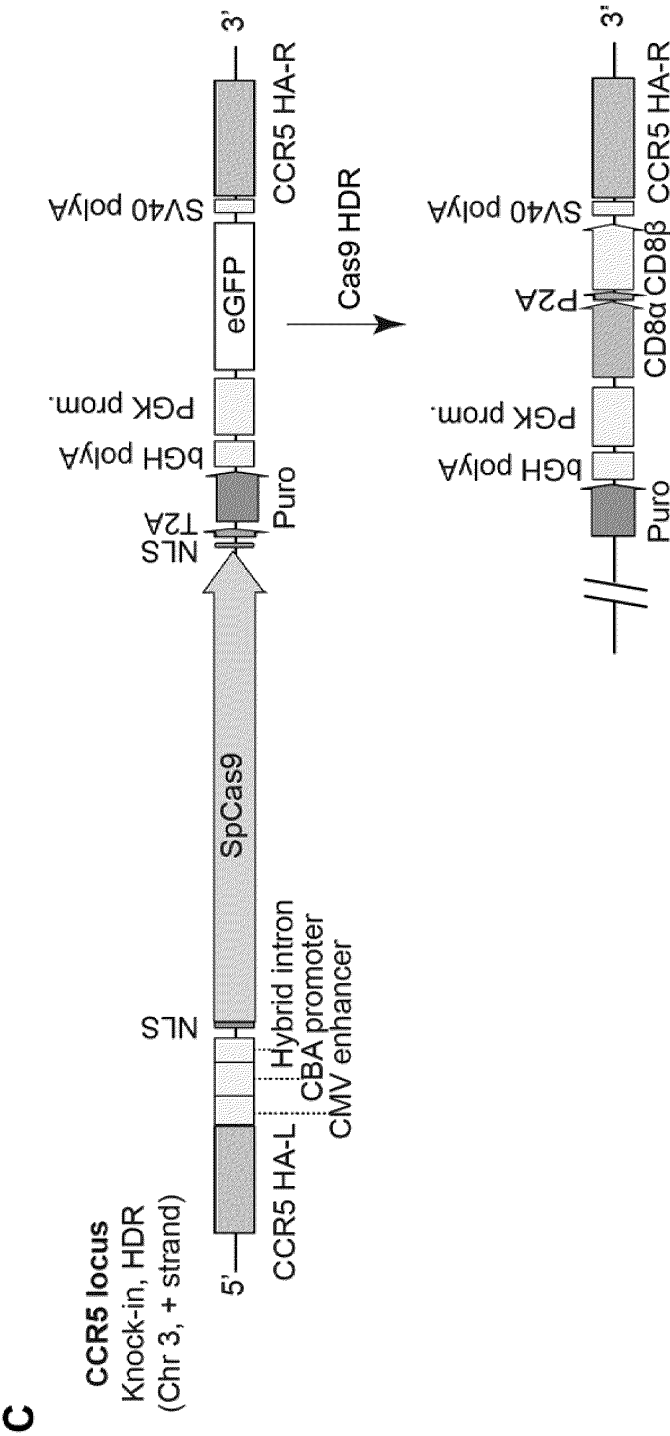


FIG. 3

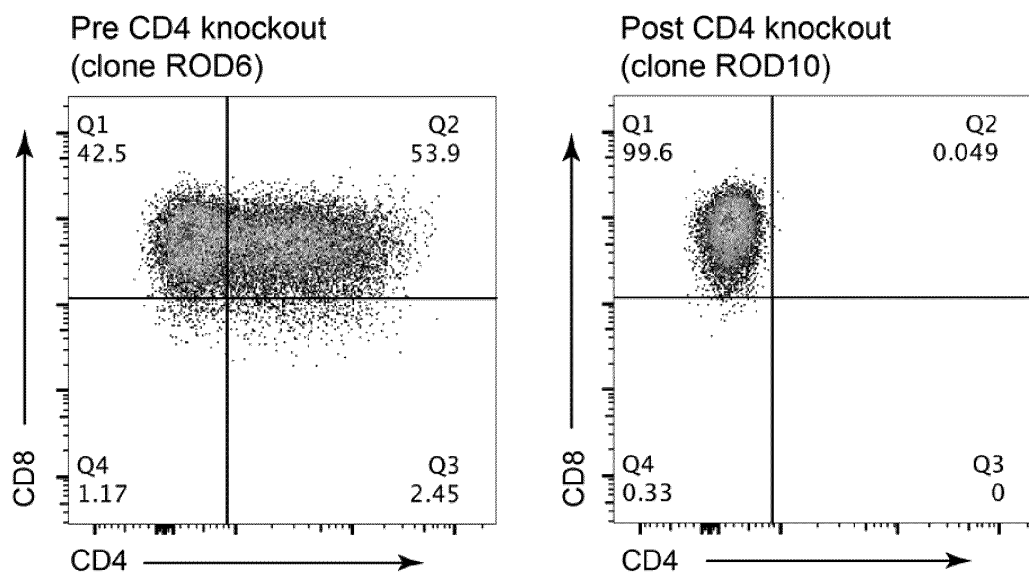


FIG. 4 A

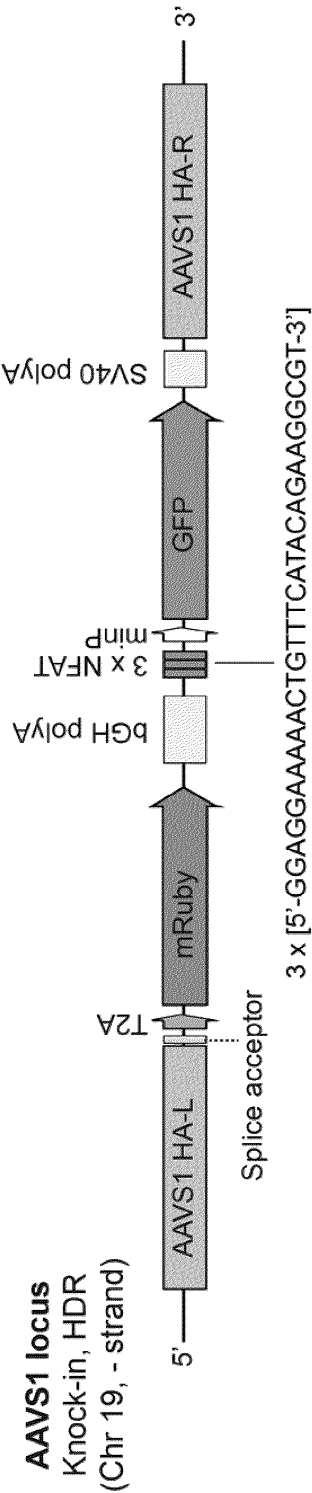


FIG. 4 B

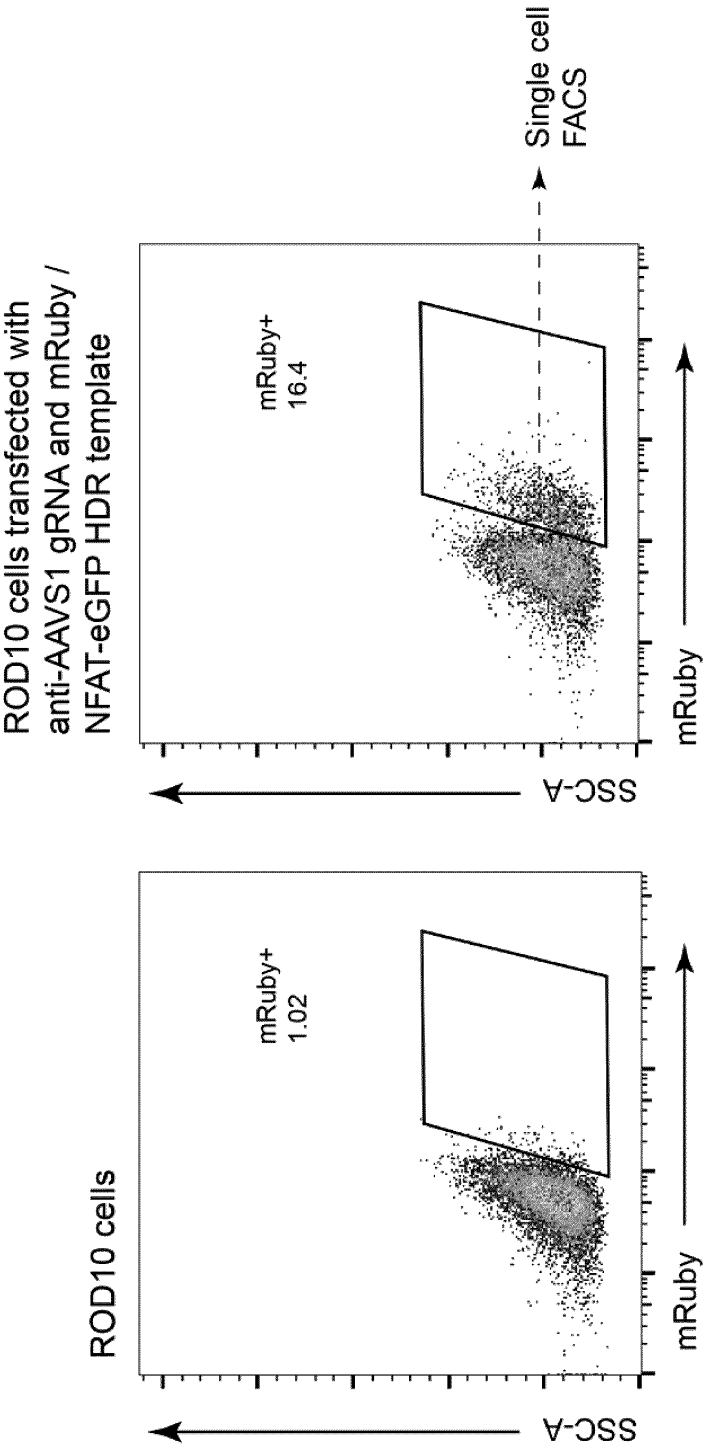


FIG. 4 C

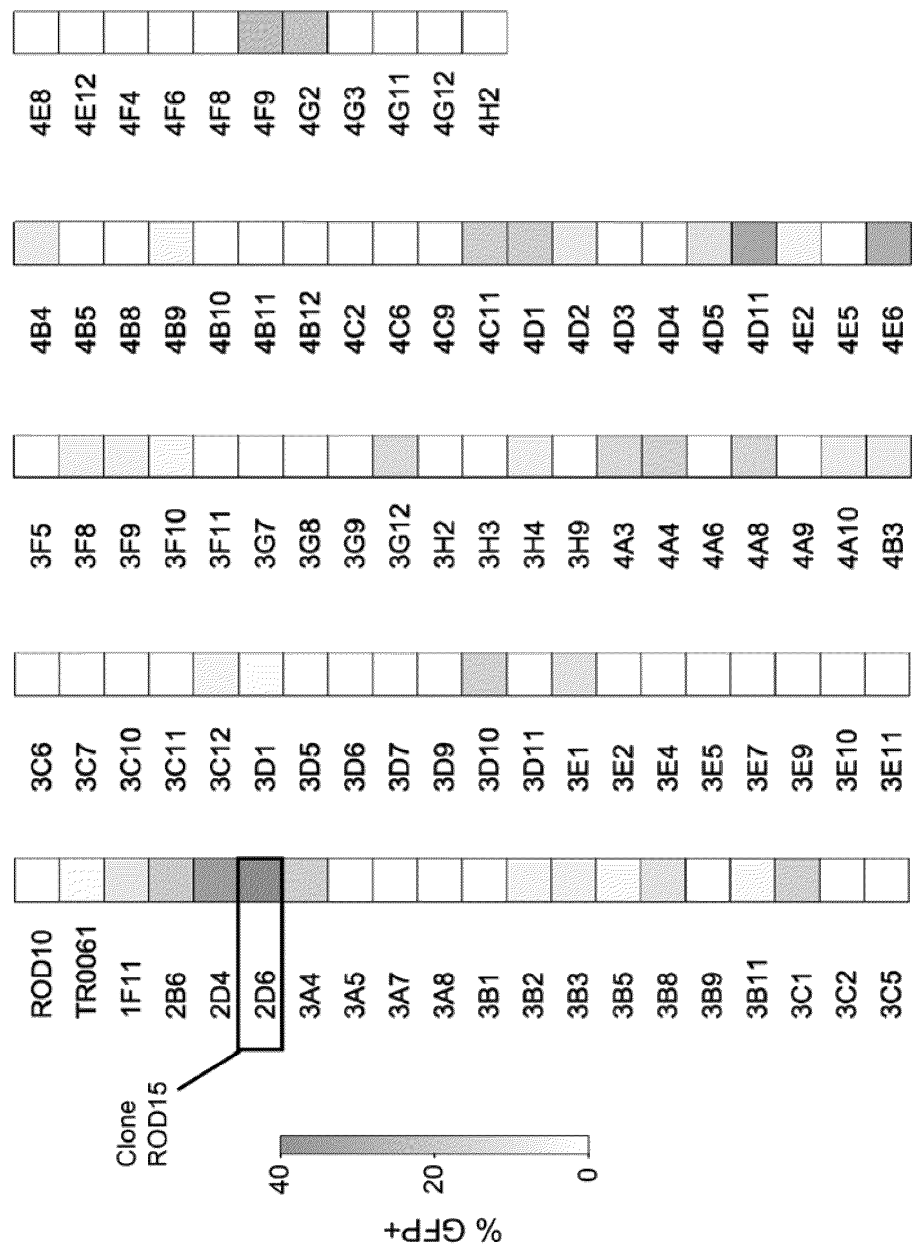


FIG. 4 D

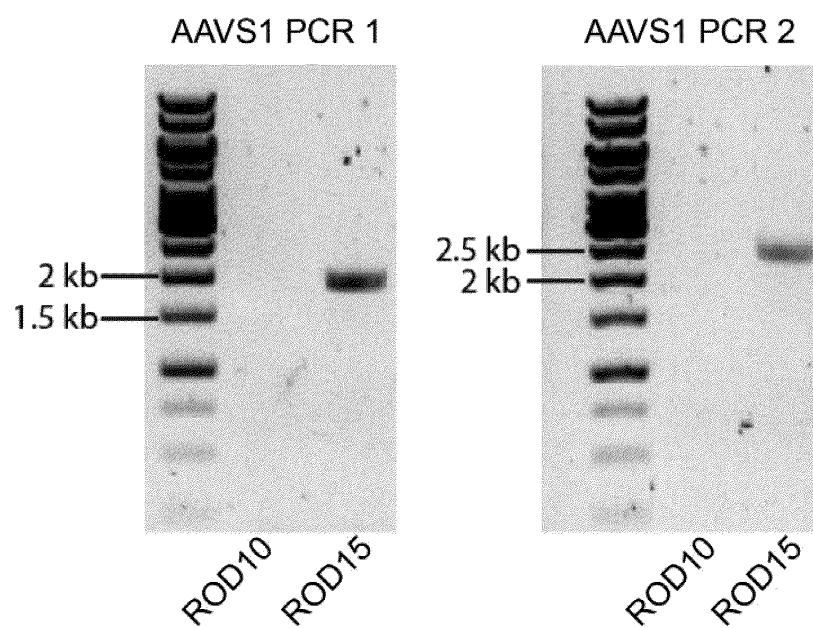


FIG. 5

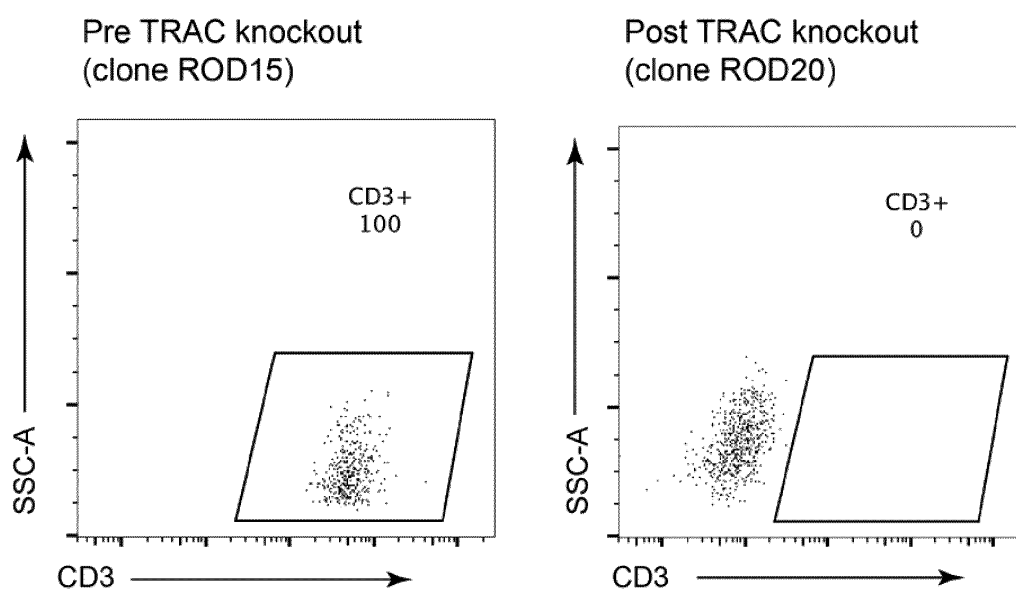


FIG. 6 A

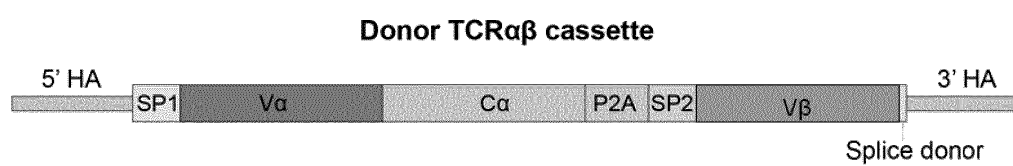


FIG. 6 B

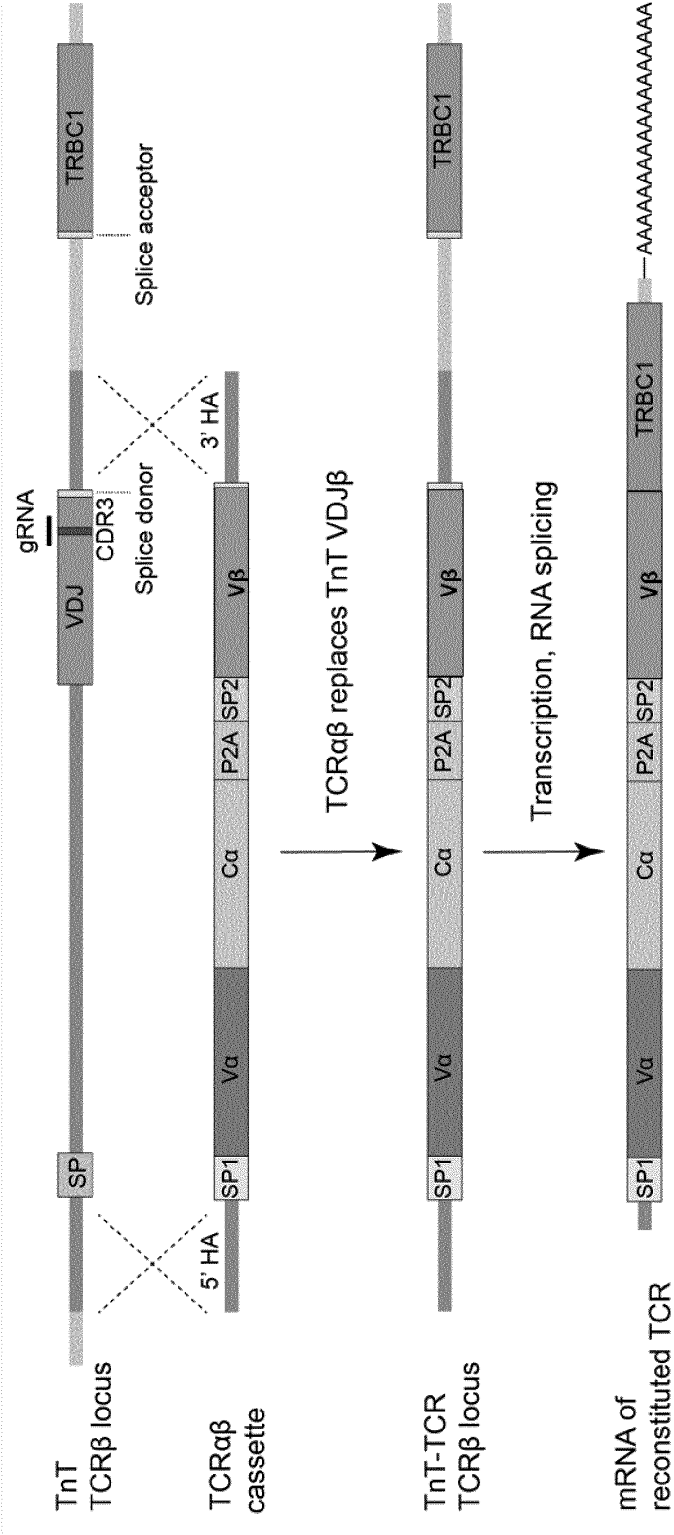


FIG. 6 C

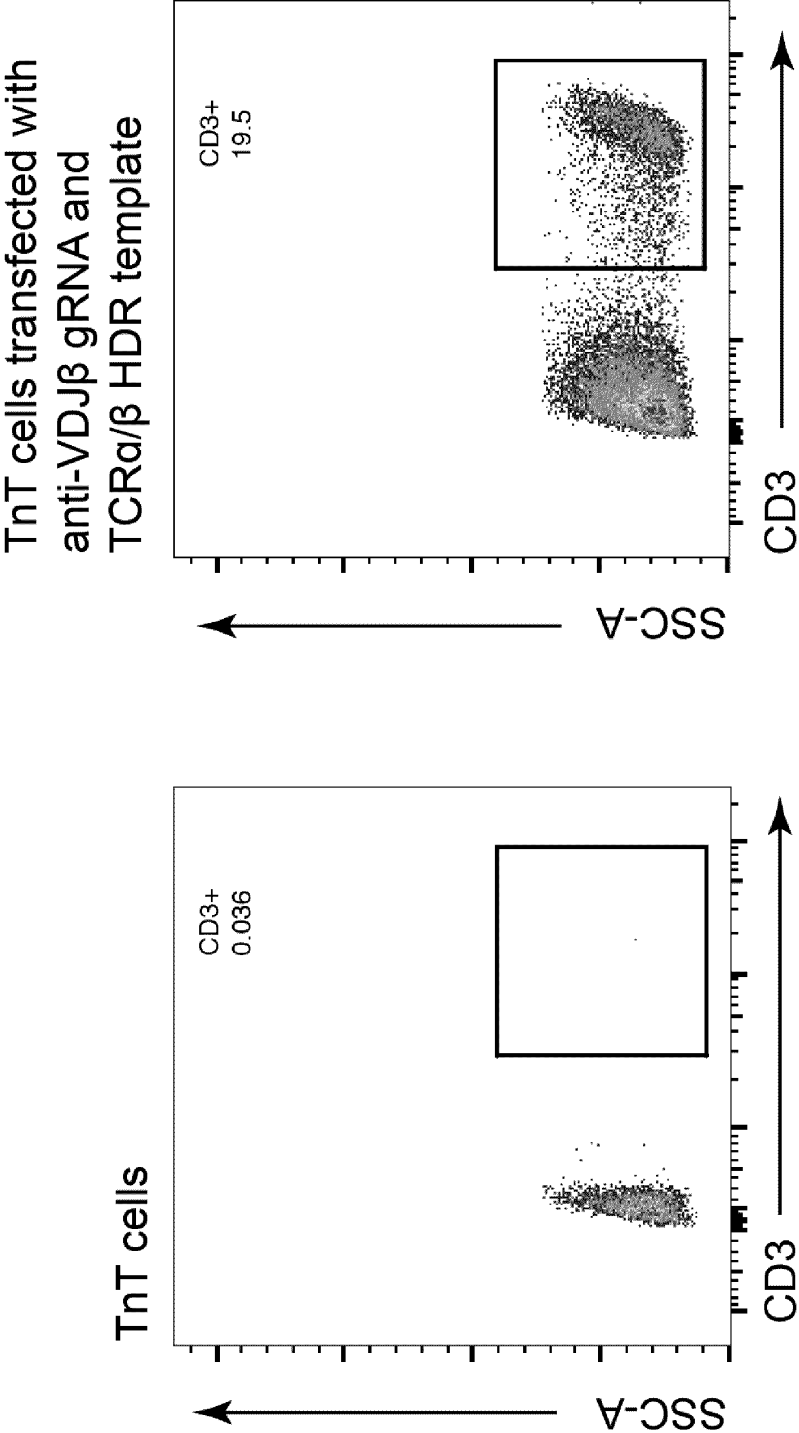


FIG. 6 D

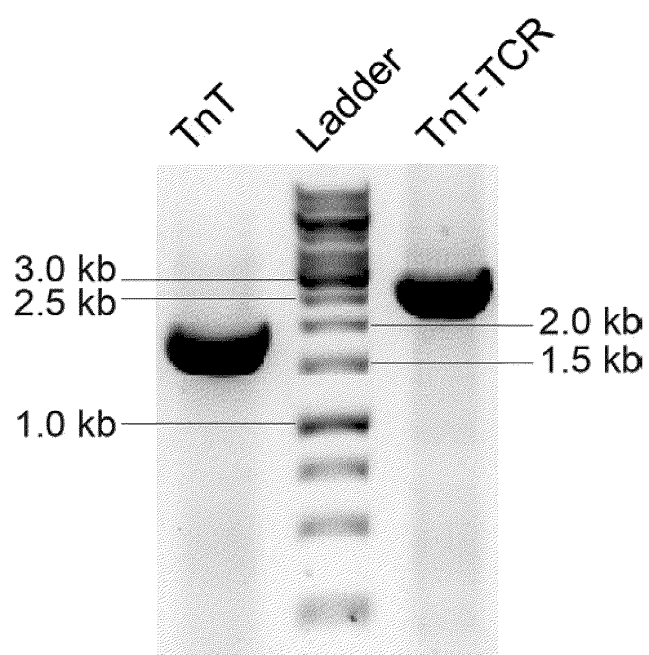


FIG. 7

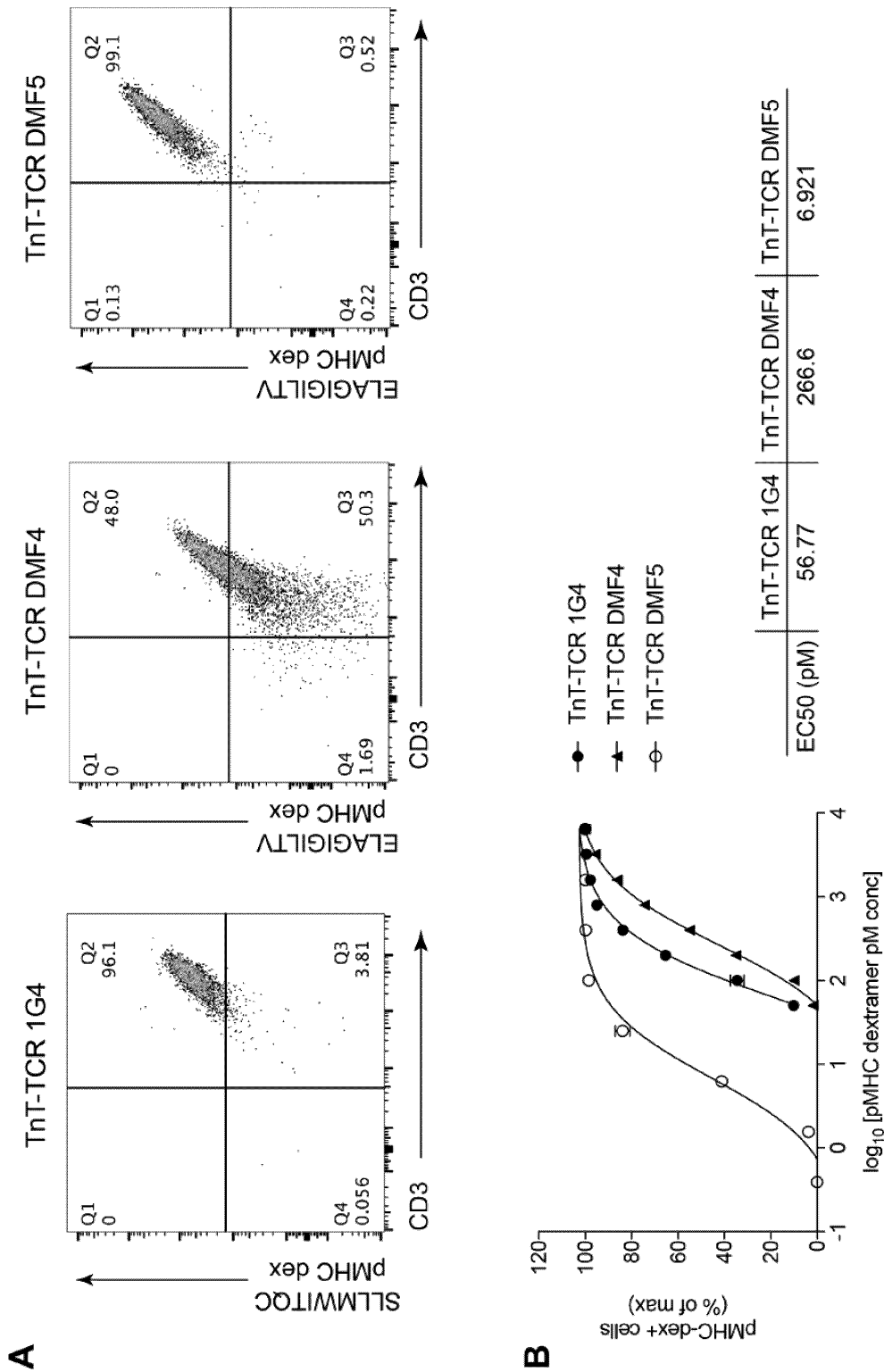


FIG. 8

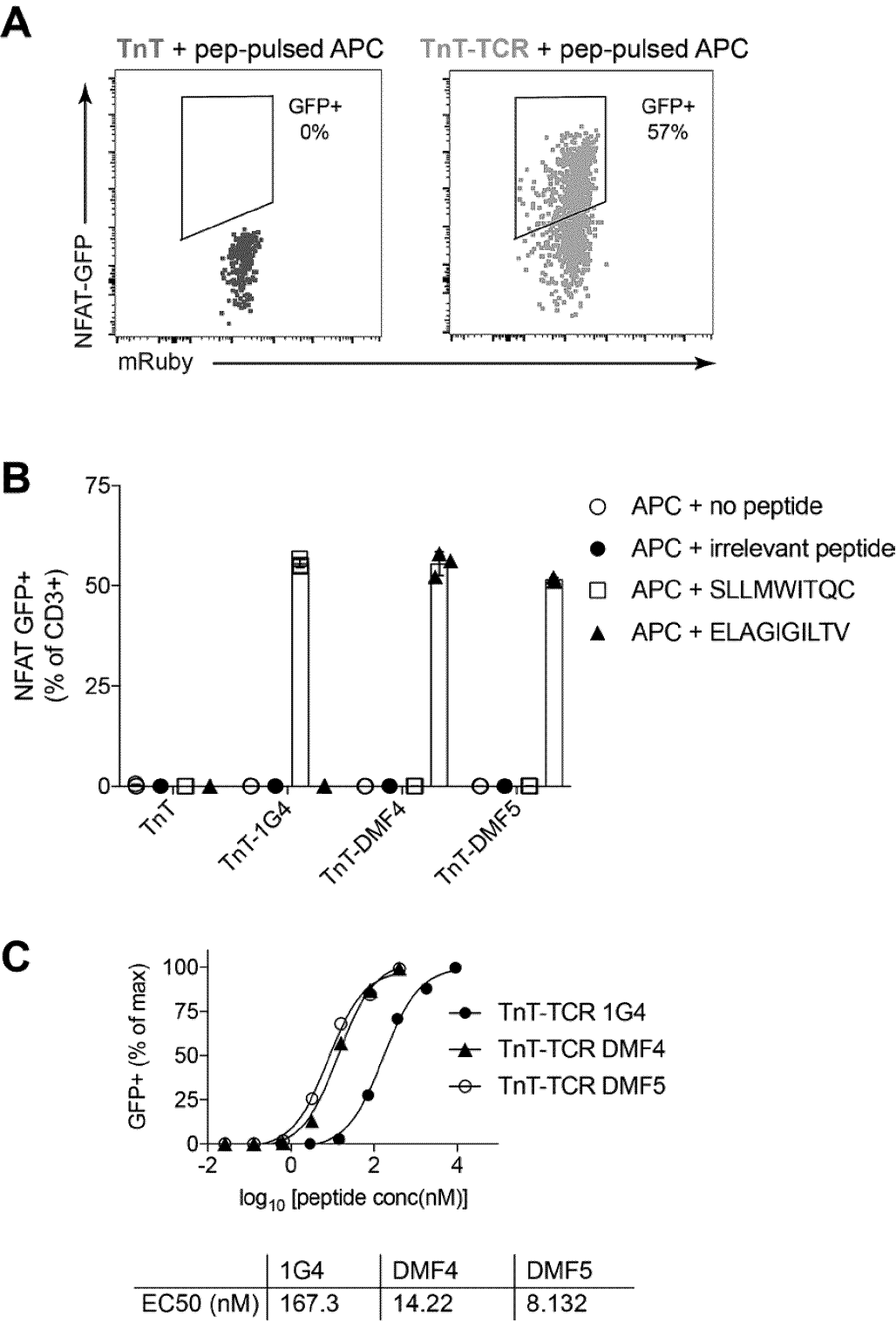


FIG. 9

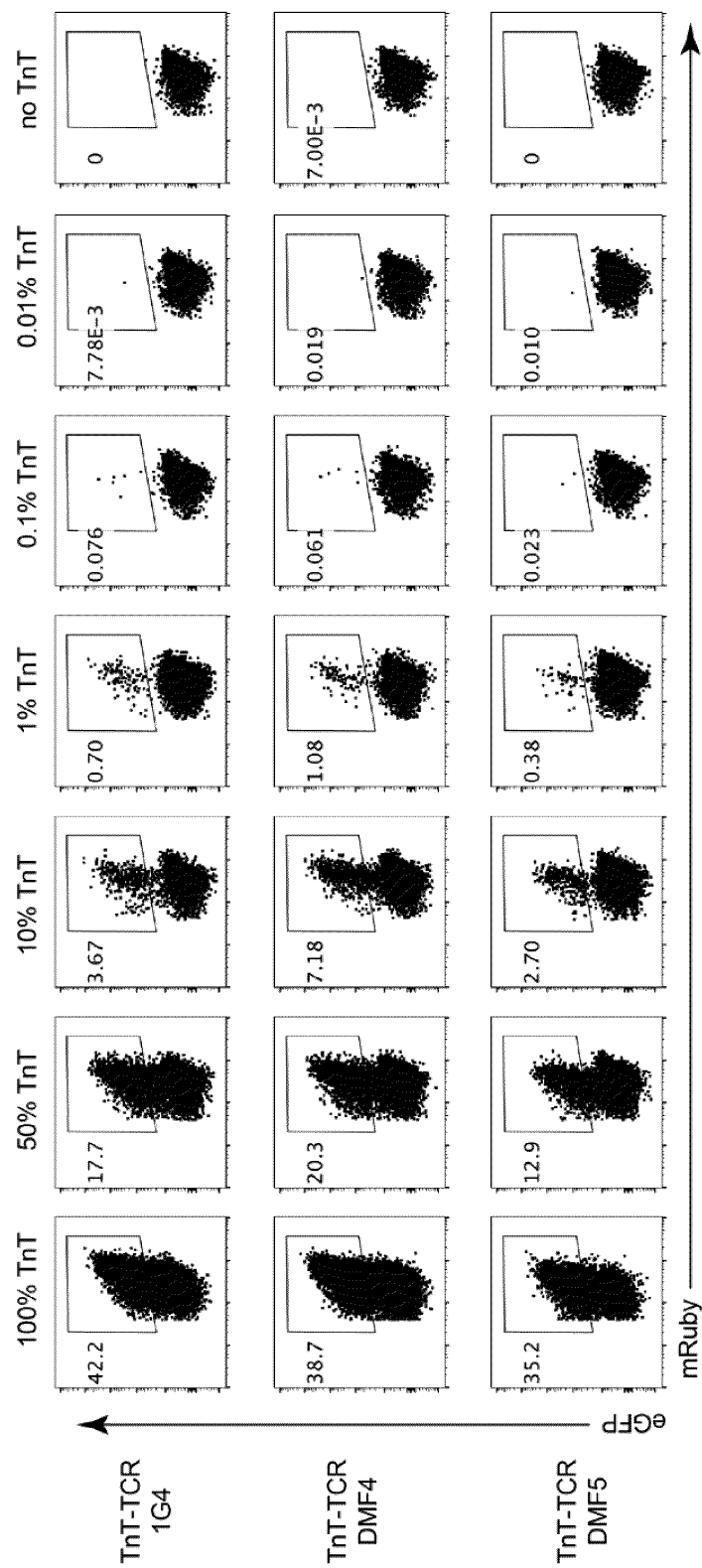
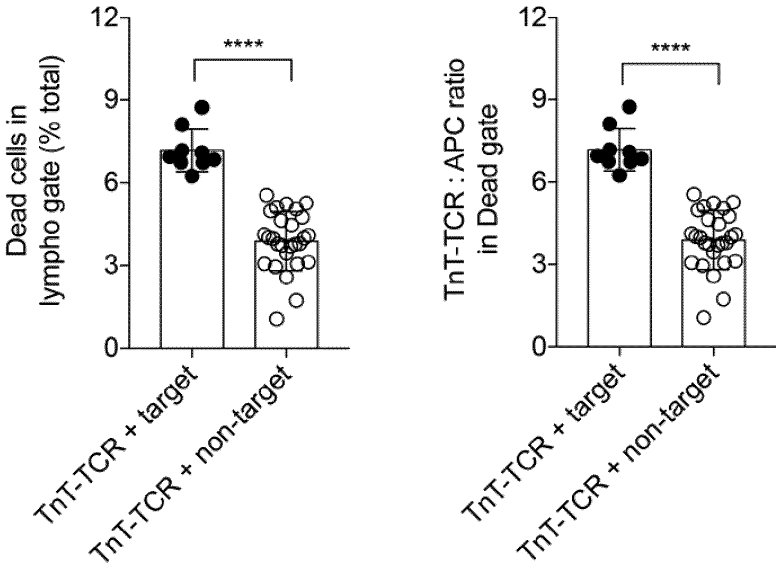


FIG. 10

A



B

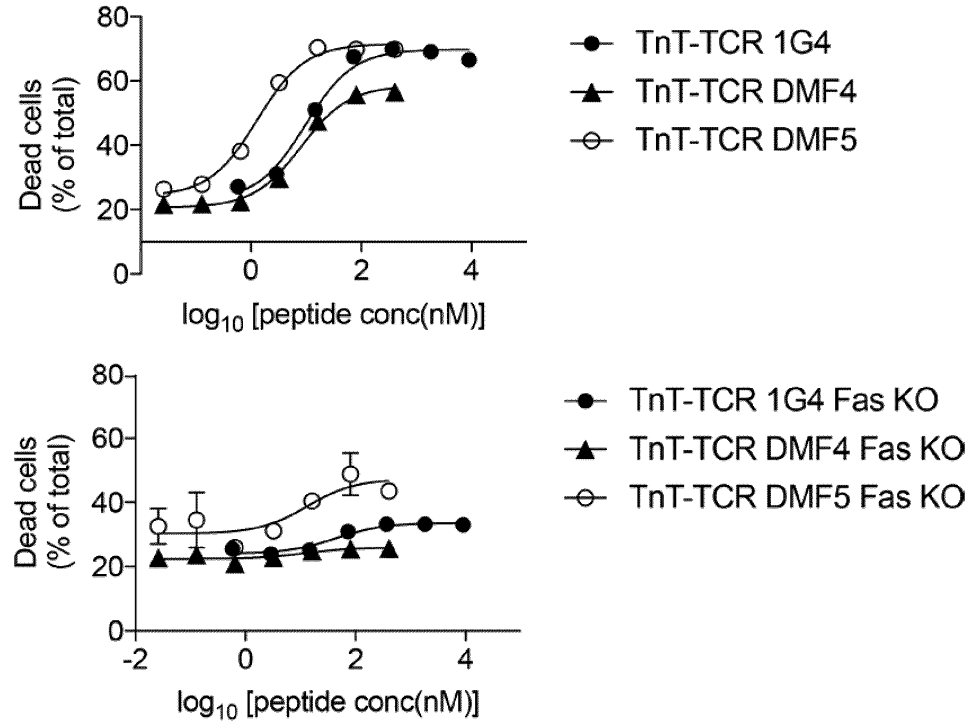


FIG. 10 C

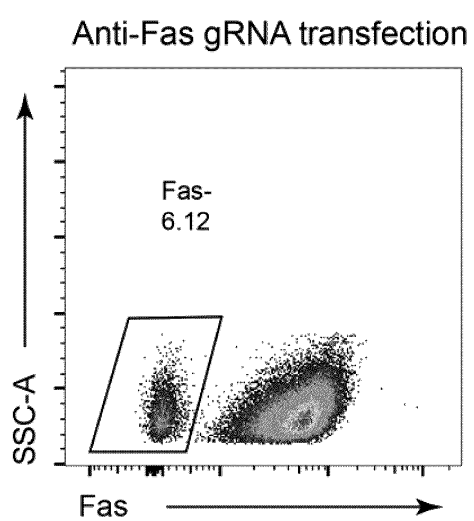


FIG. 10 D

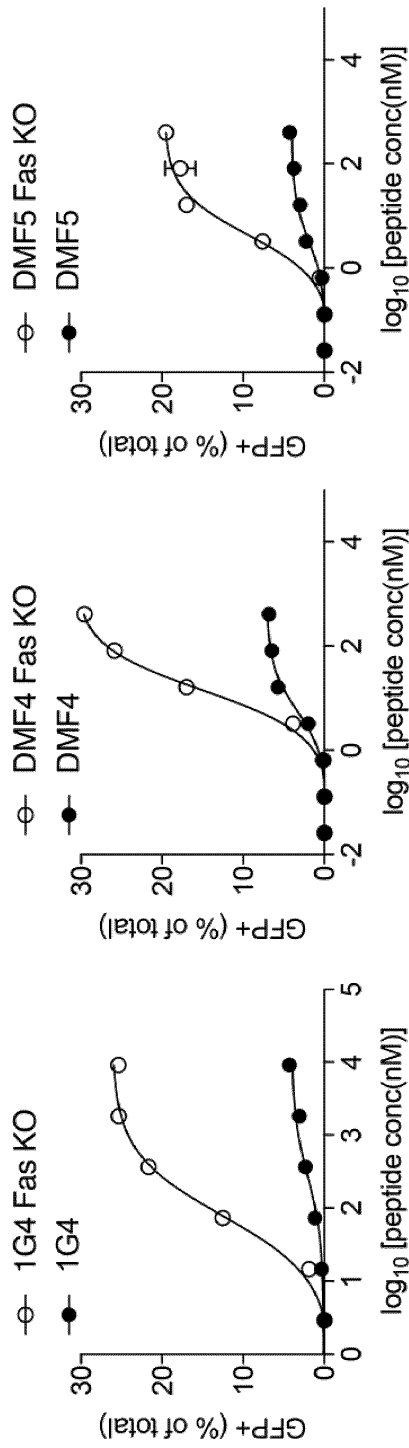


FIG. 11 A

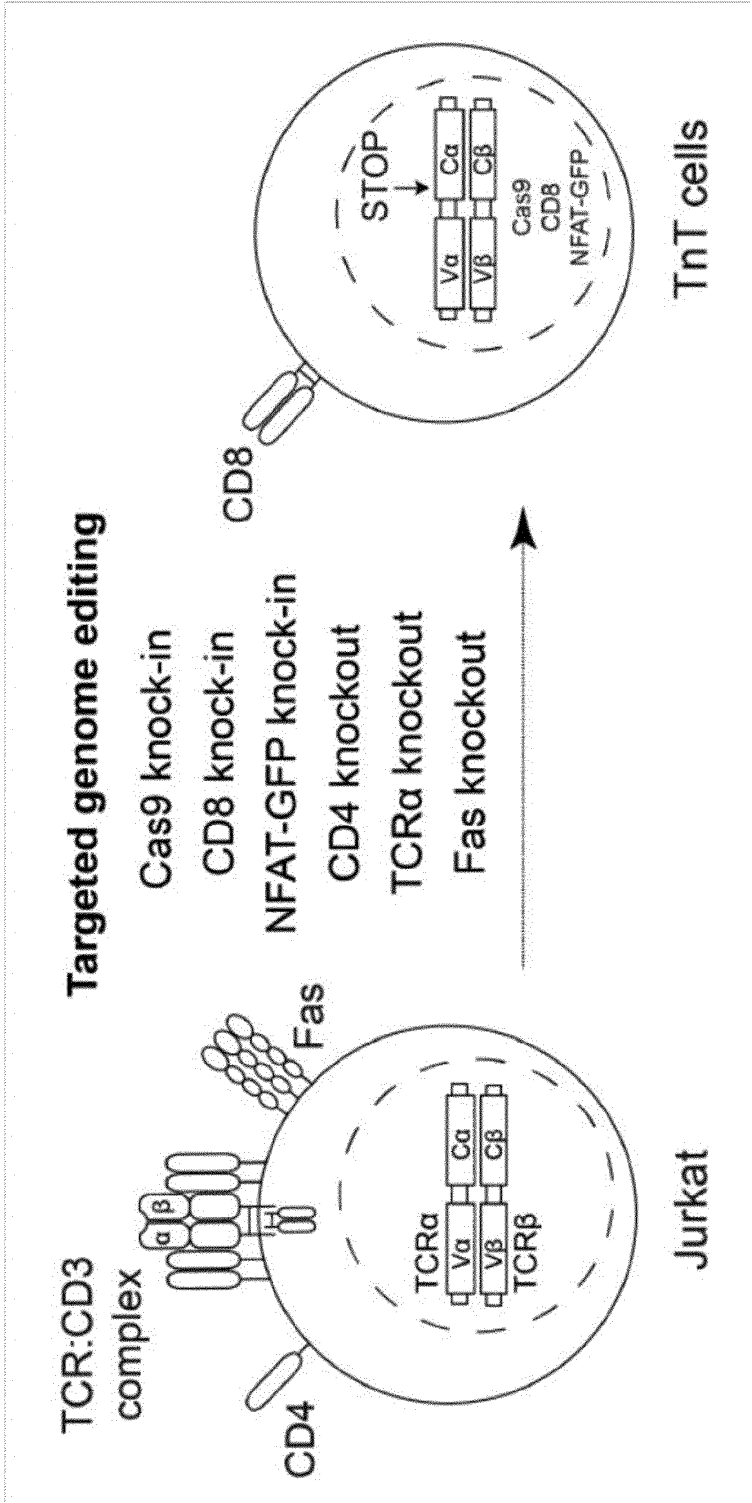


FIG. 11 B

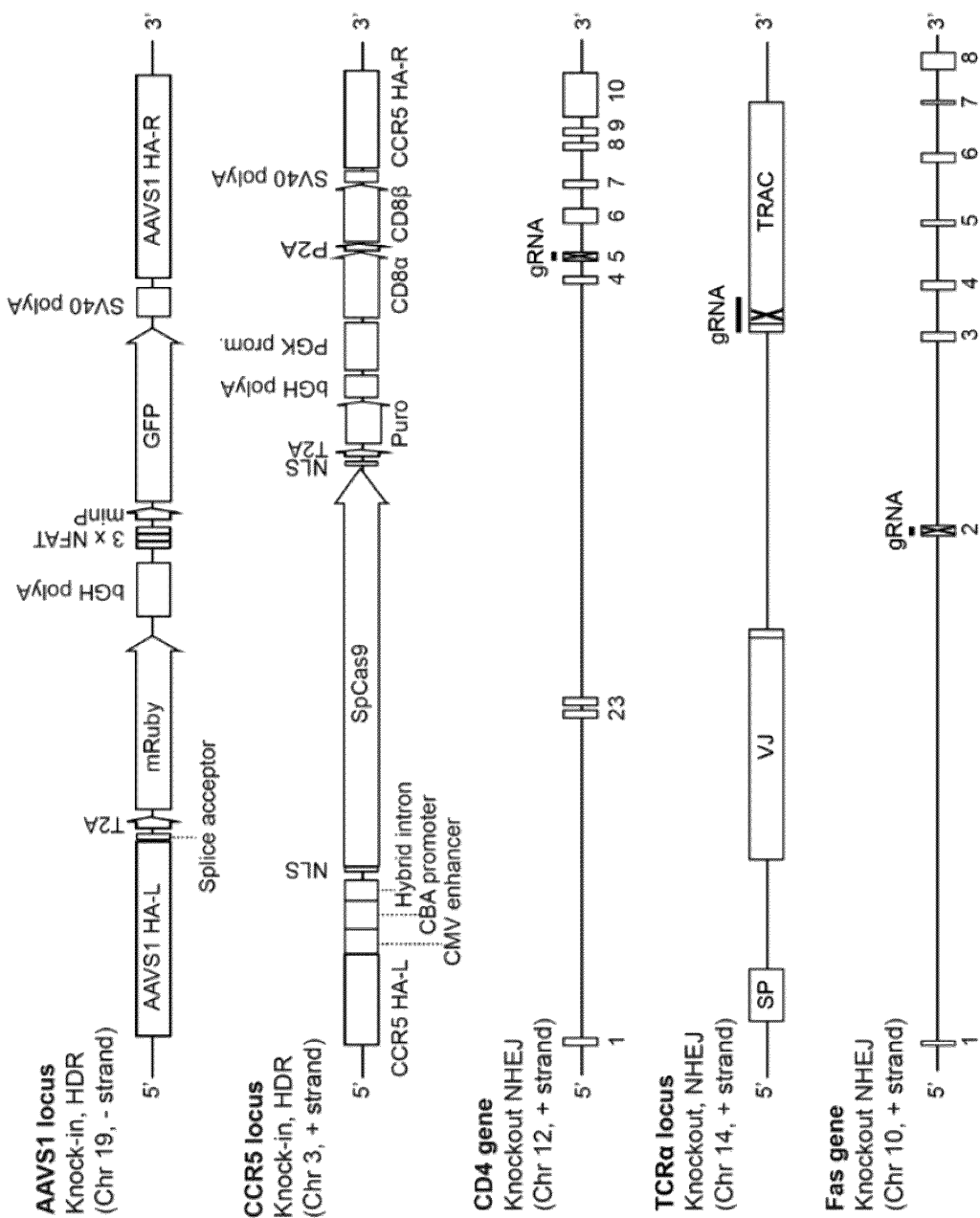


FIG. 12

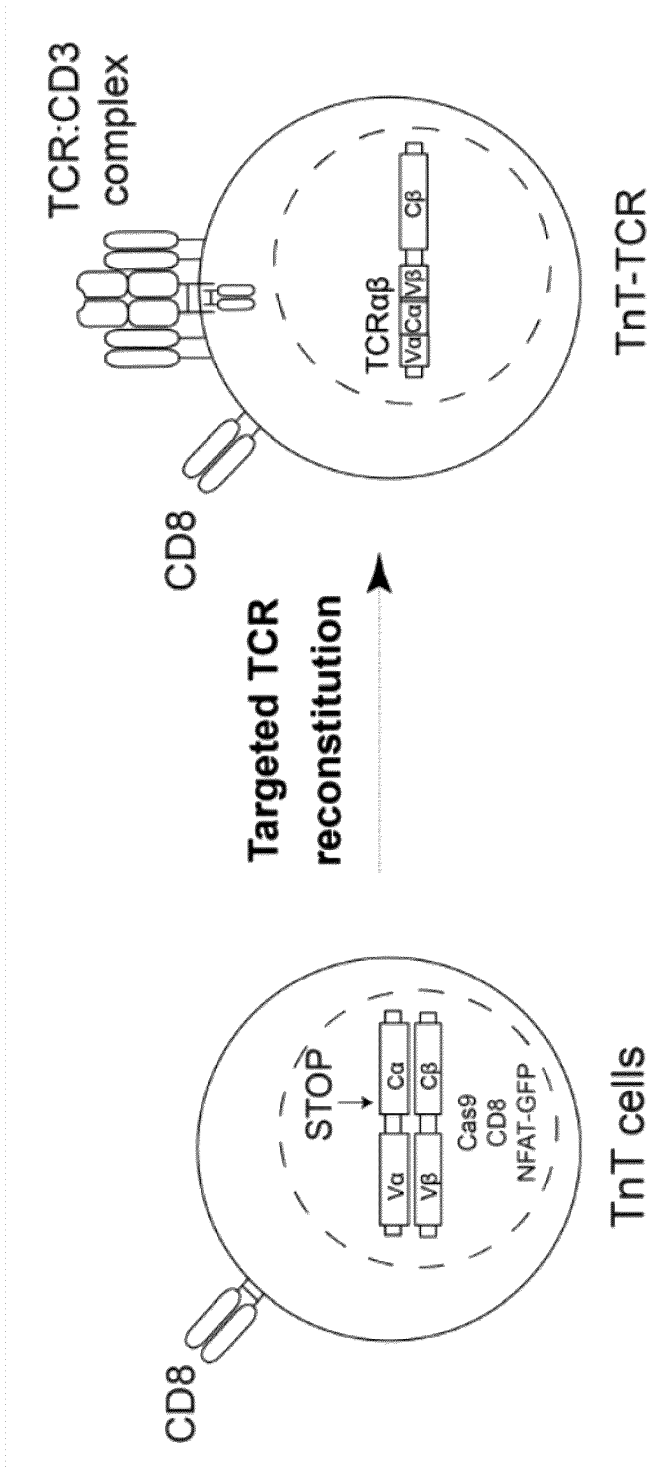


FIG. 13

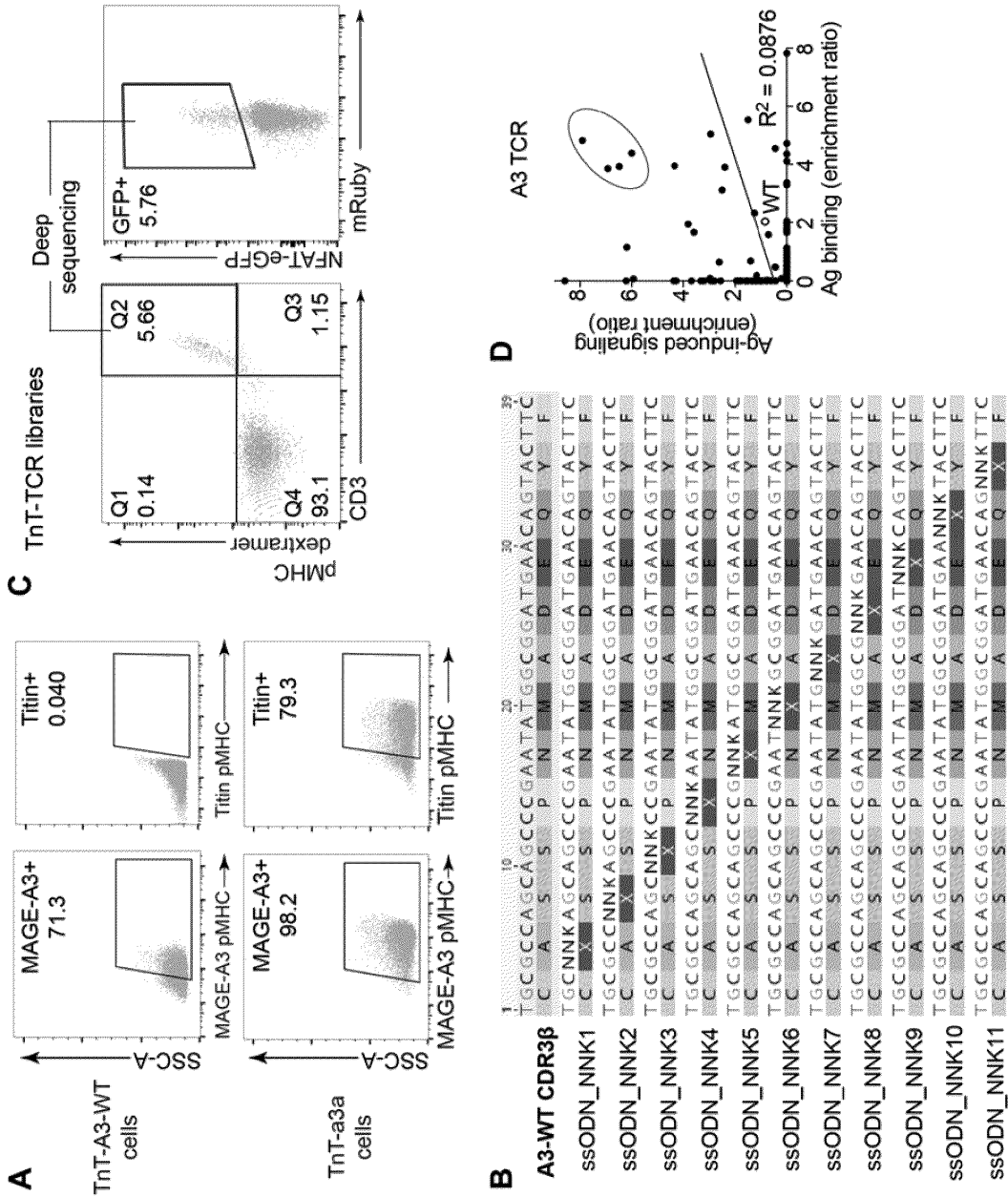


FIG. 14 A

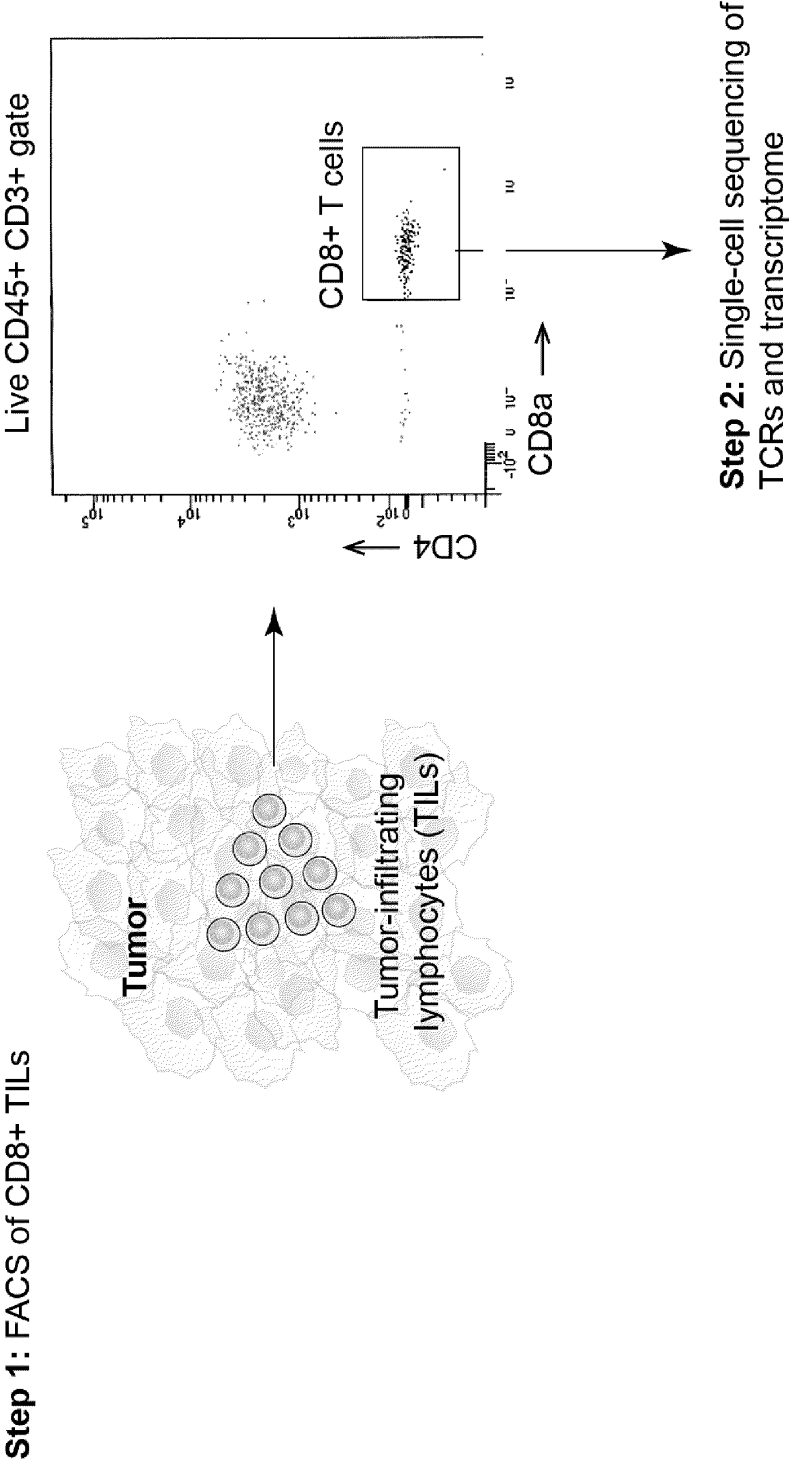


FIG. 14 B

Step 3: Analysis of sequencing data, identification of candidateTCRs

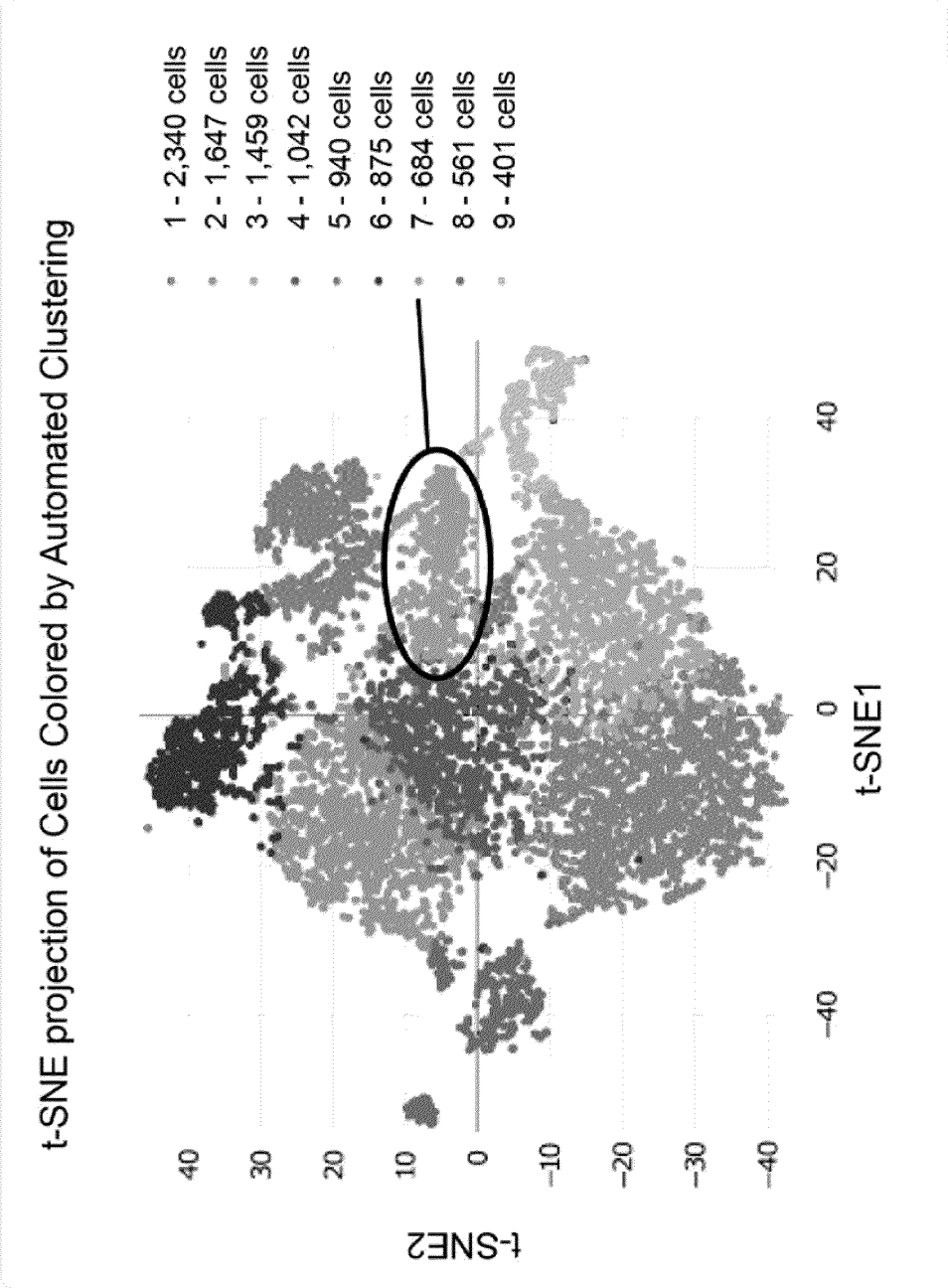


FIG. 14 C

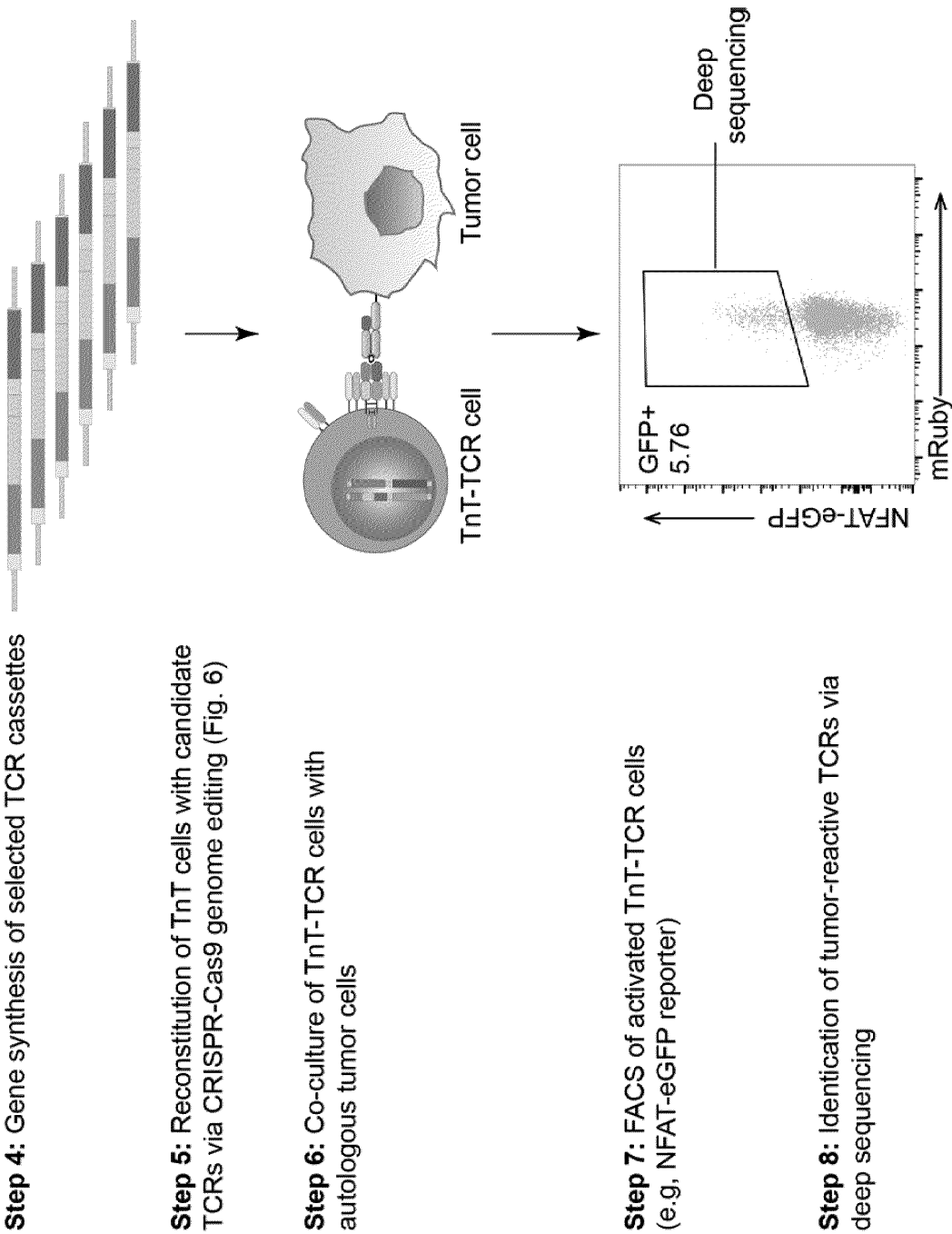


FIG.15

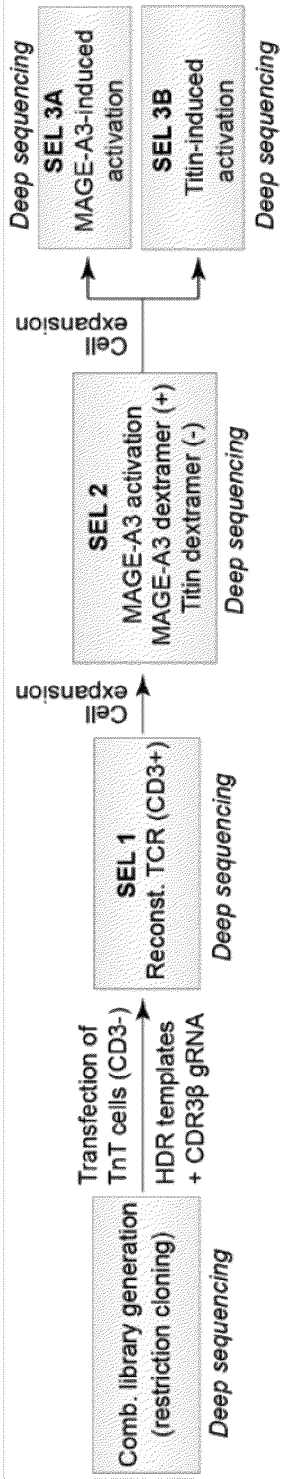


FIG.16

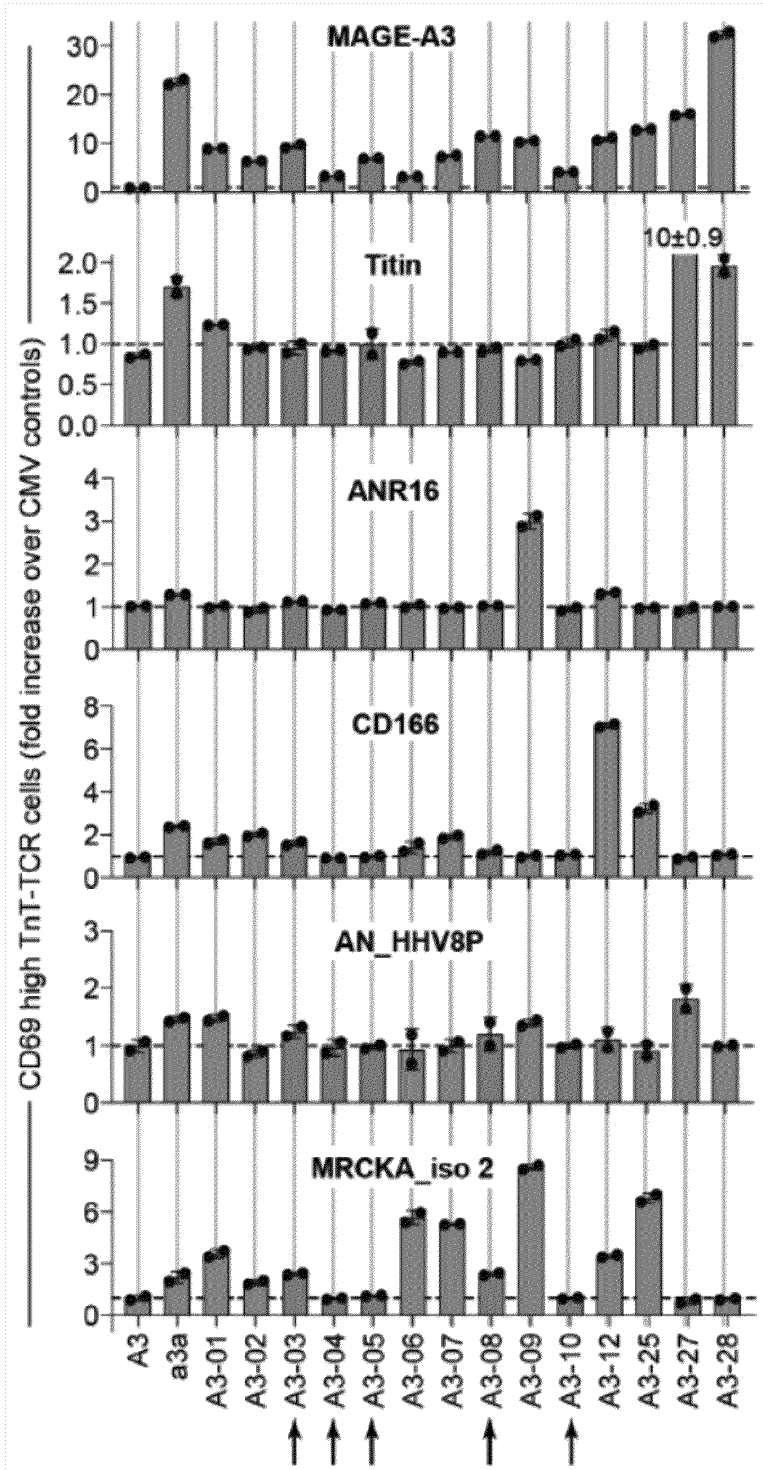


FIG.17

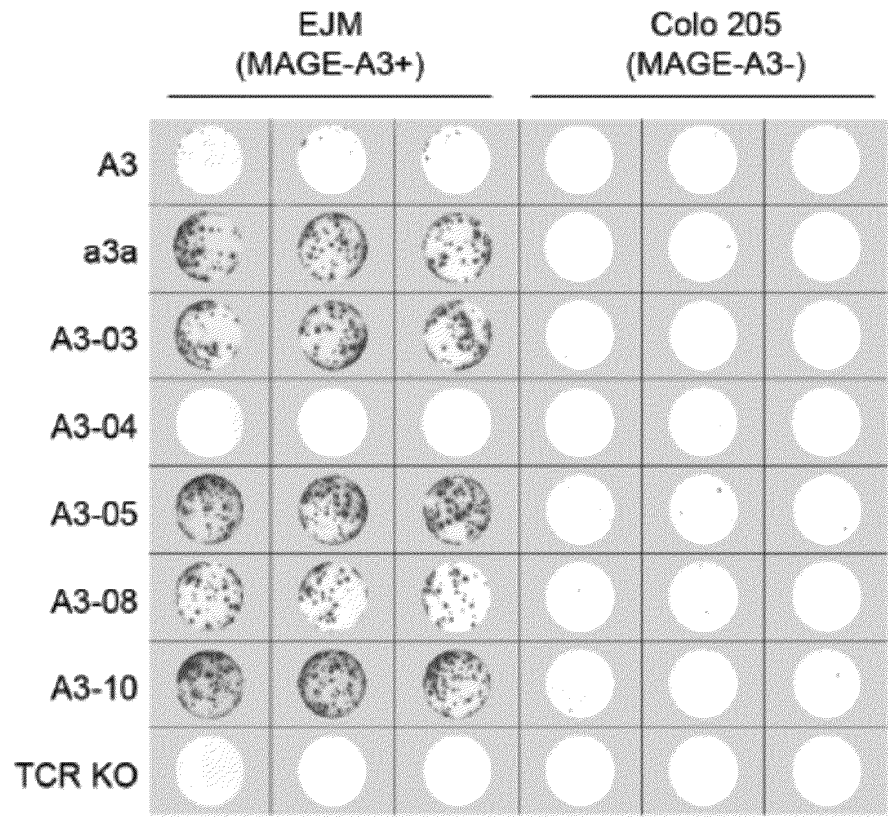


FIG.18

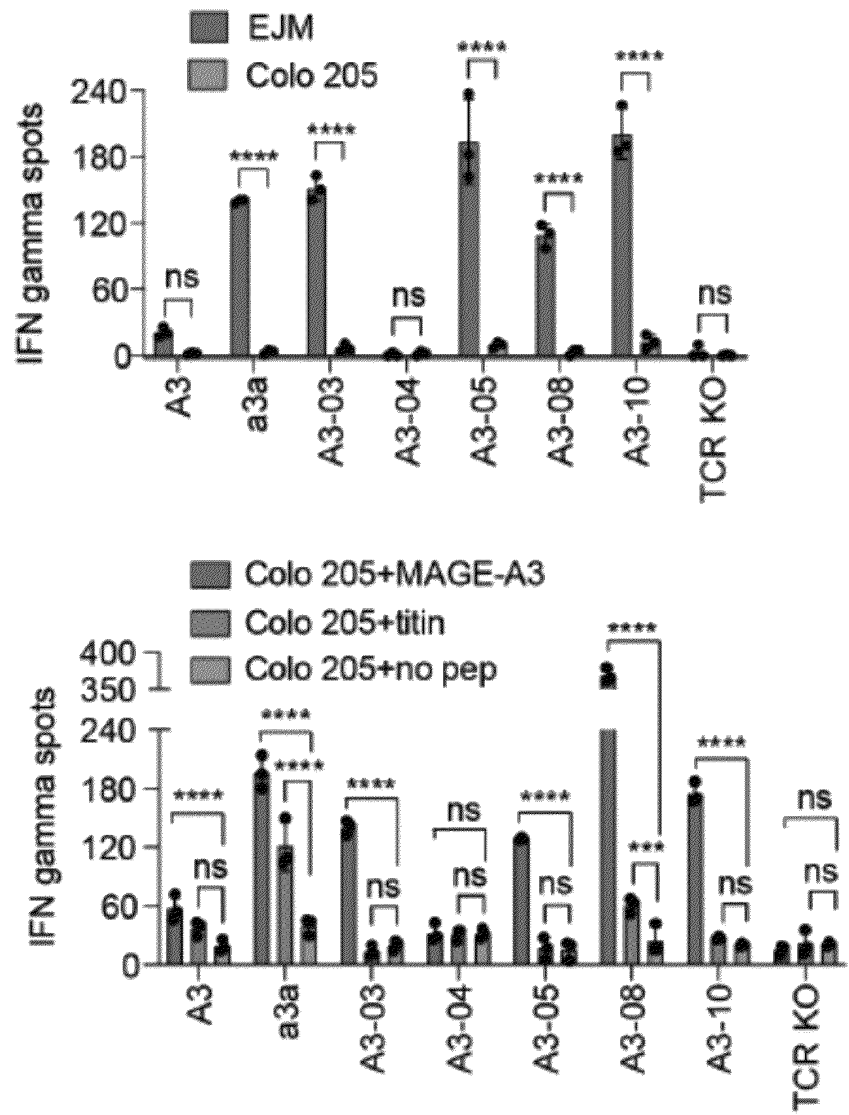


FIG.19

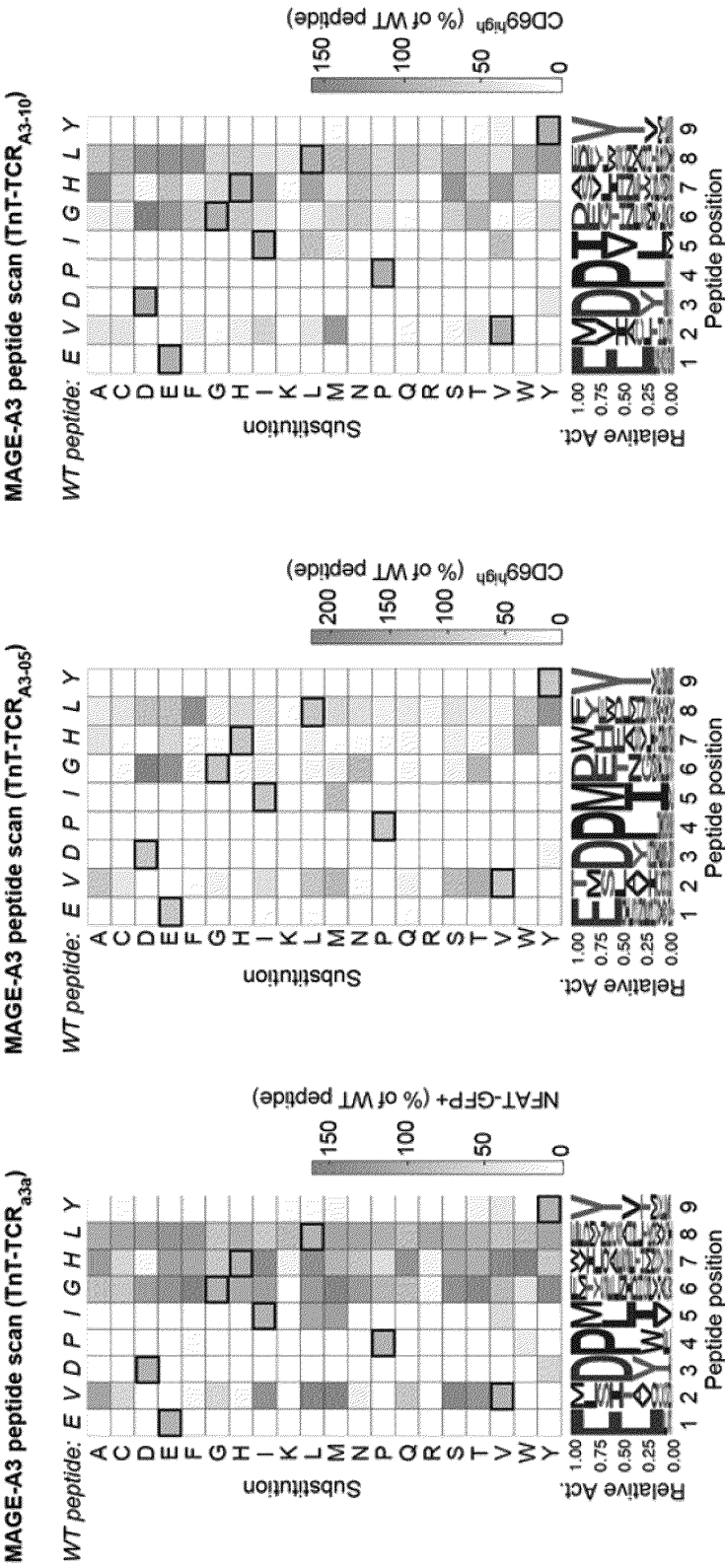


FIG.20

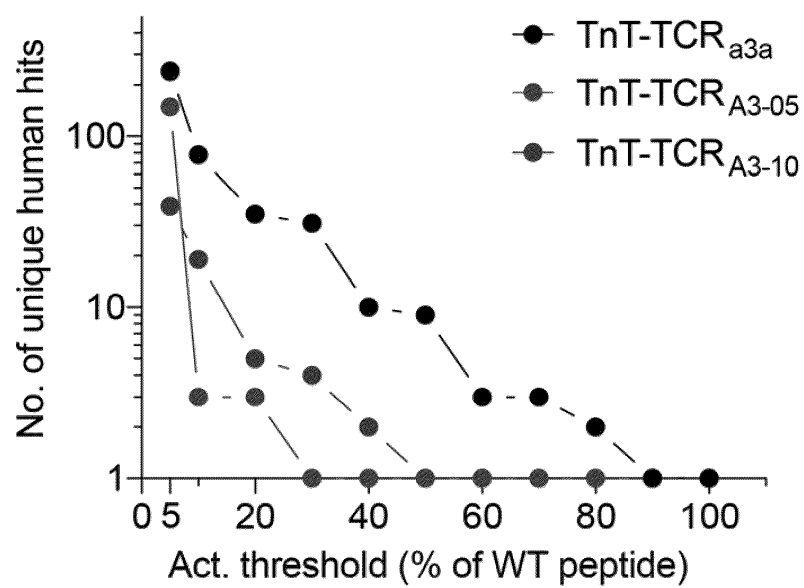


FIG.21

Target: EJM cells (MAGE-A3+)

