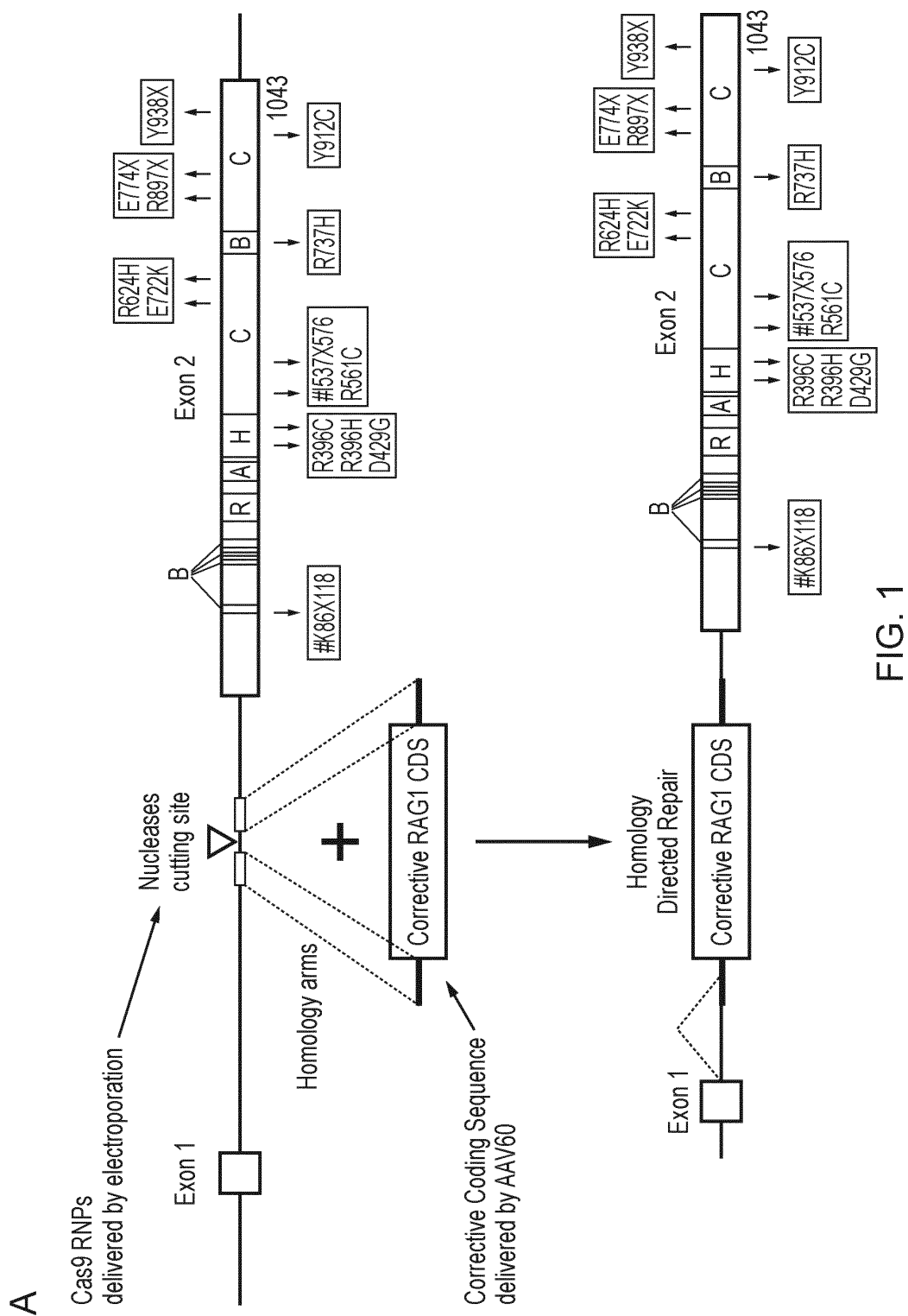


1/29



2/29

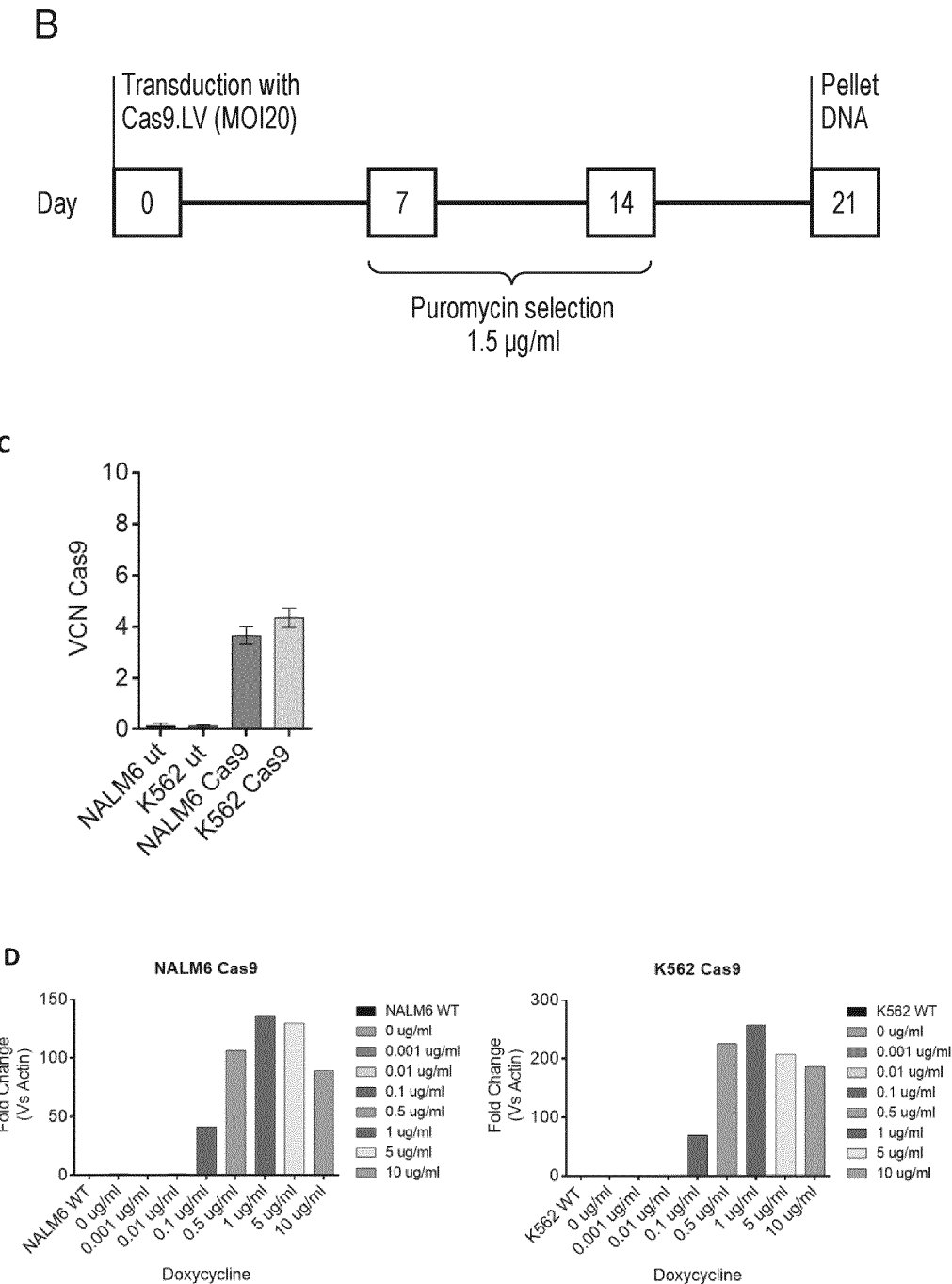


FIG. 1 (Continued)

3/29

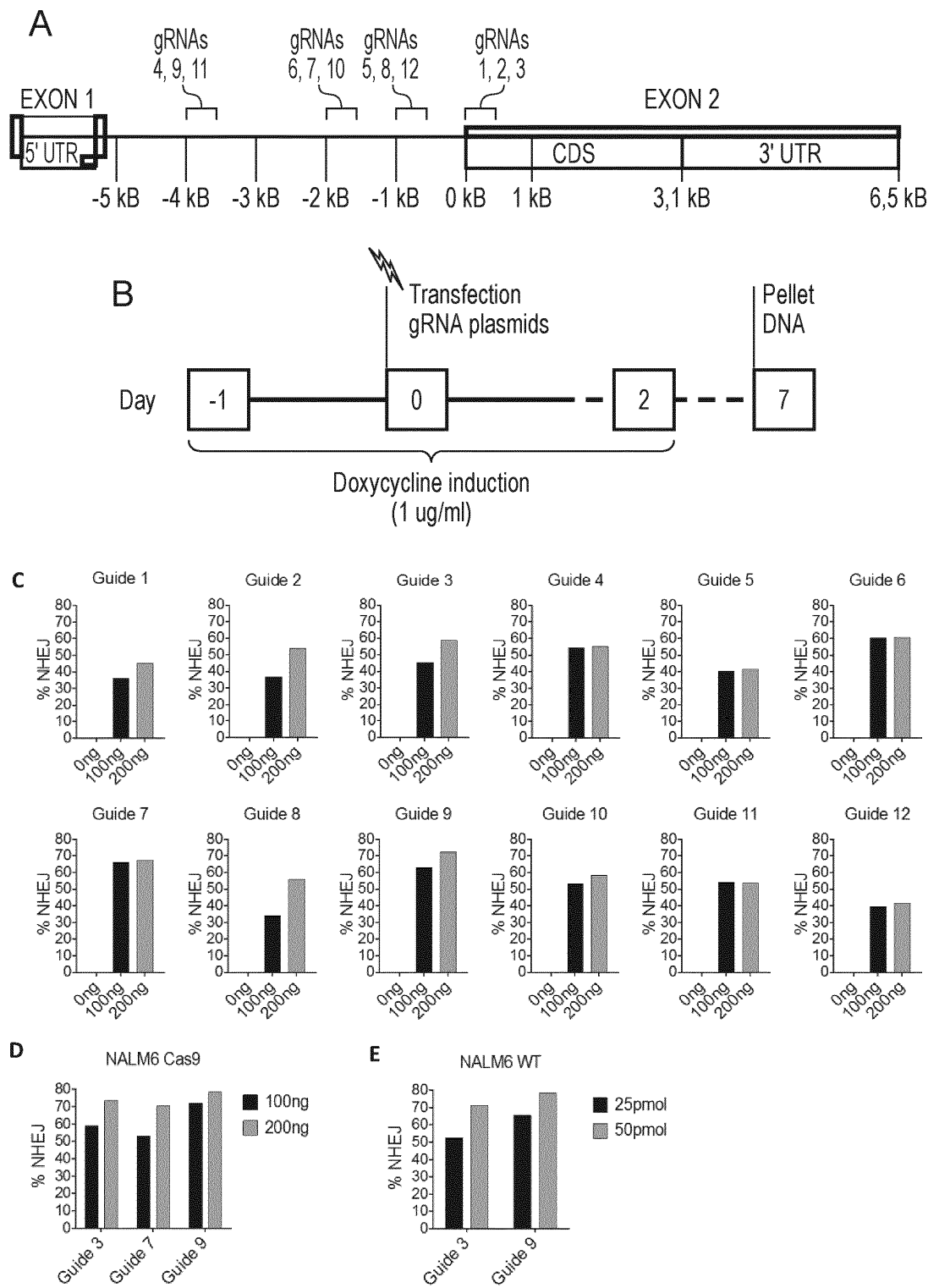


FIG. 2

SUBSTITUTE SHEET (RULE 26)

4/29

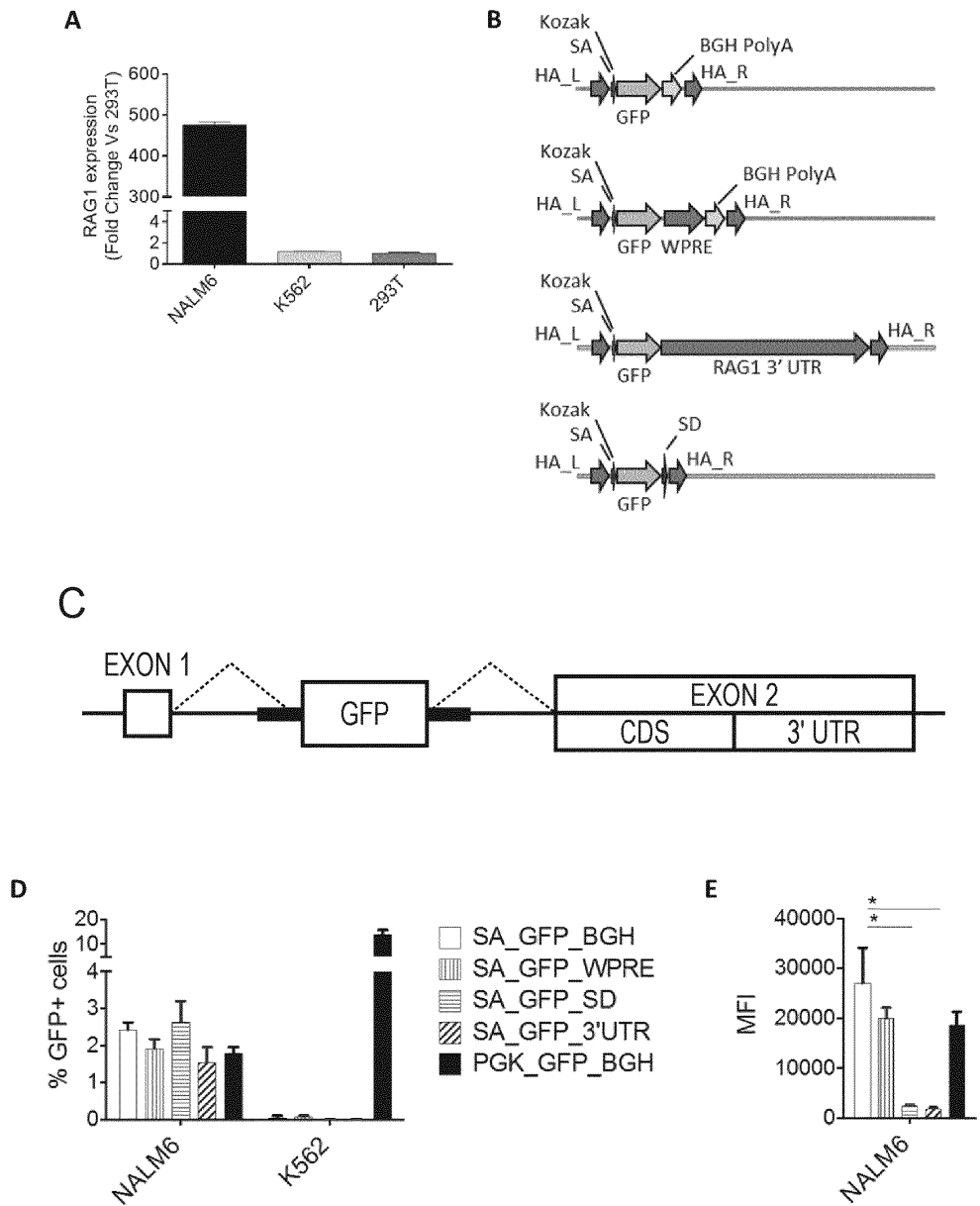


FIG. 3

5/29

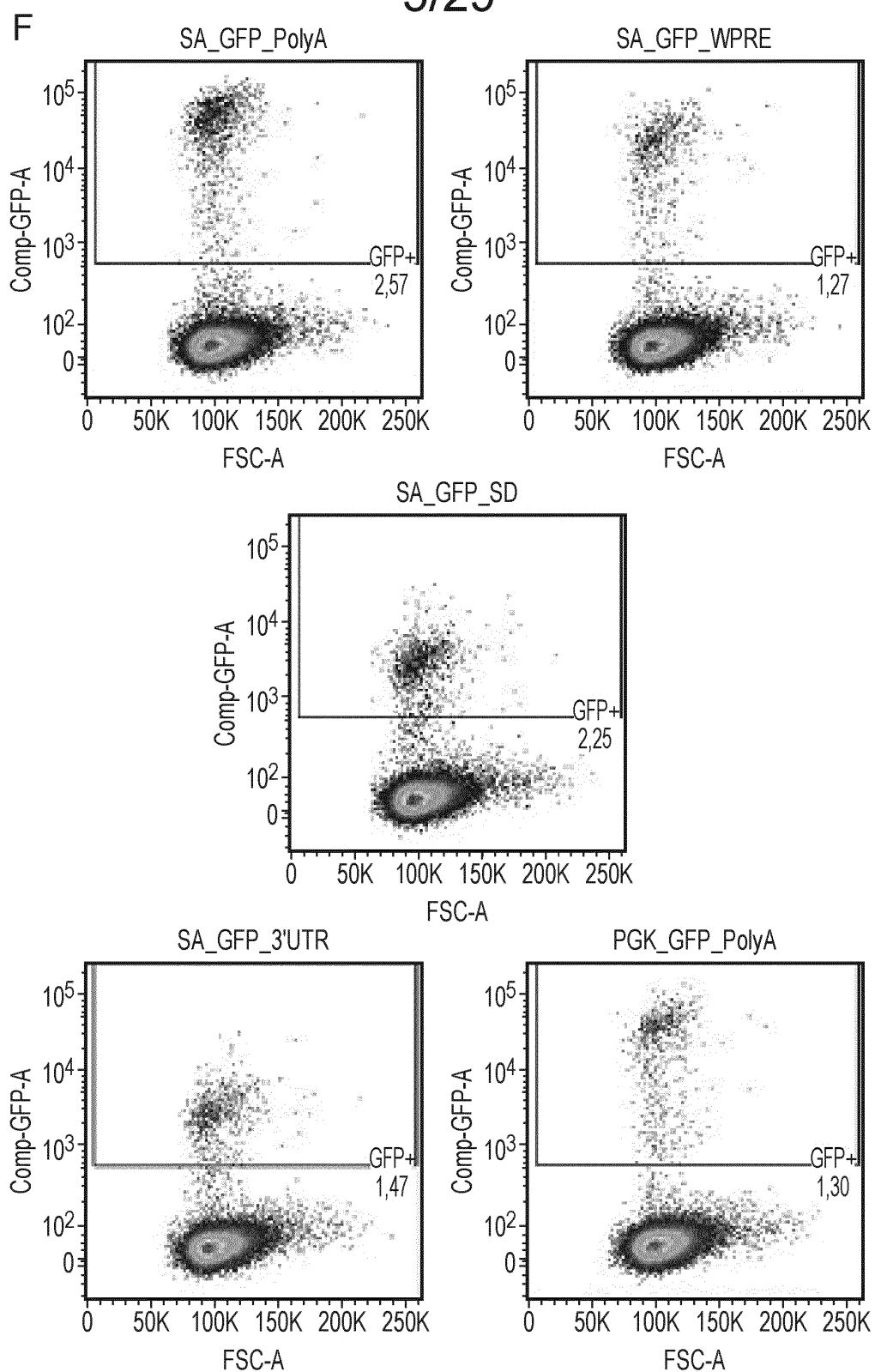


FIG. 3 (Continued)
SUBSTITUTE SHEET (RULE 26)

6/29

A

	Sequence	PAM	Score	#MM	Gene	Locus
	GTCAGATGGCAATGTCGAGA	TGG	N/A			chr11:+36569279
1	GCCA-ATGTCAATGTCGAGA	GGG	23	3		chr3:-26123249
2	GTCA-AAGTCAATGTCGAGA	AGG	28	3		chr18:+53676614
3	ATCAGATGGCAA-GTCGAGA	AGG	33	2		chr2:+1937736
4	TTCAGATGGAAATGTCAAGA	AAG	33	3		chr6:+130332344
5	GTAAGAT-GCAATGTAGAGA	AAG	34	3		chr9:+212232
6	GTCAGATGGCAATGTC-AGA	TGG	35	1		chr10:+1807106
7	ATCAGATCGCAA-GTCGAGA	AGG	35	3		chr2:+1937580
8	GCCAGA-GCCAATGTCGAGA	AGG	37	3		chr2:+103826298
9	GACAGCTGGCAATGTTAAGA	AAG	41	4		chr1:+234003953
10	TTCACATGGCAATGTCAAGT	TGG	42	4		chr13:+7840848

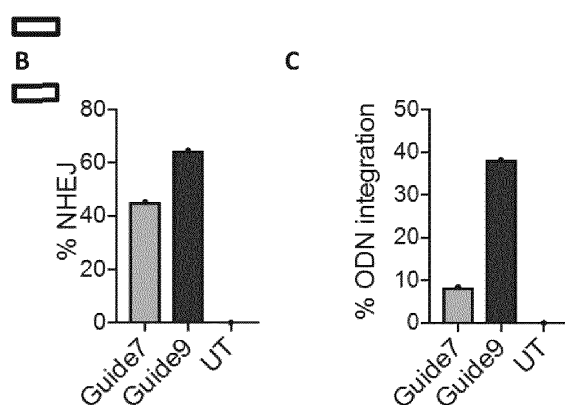


FIG. 4

7/29

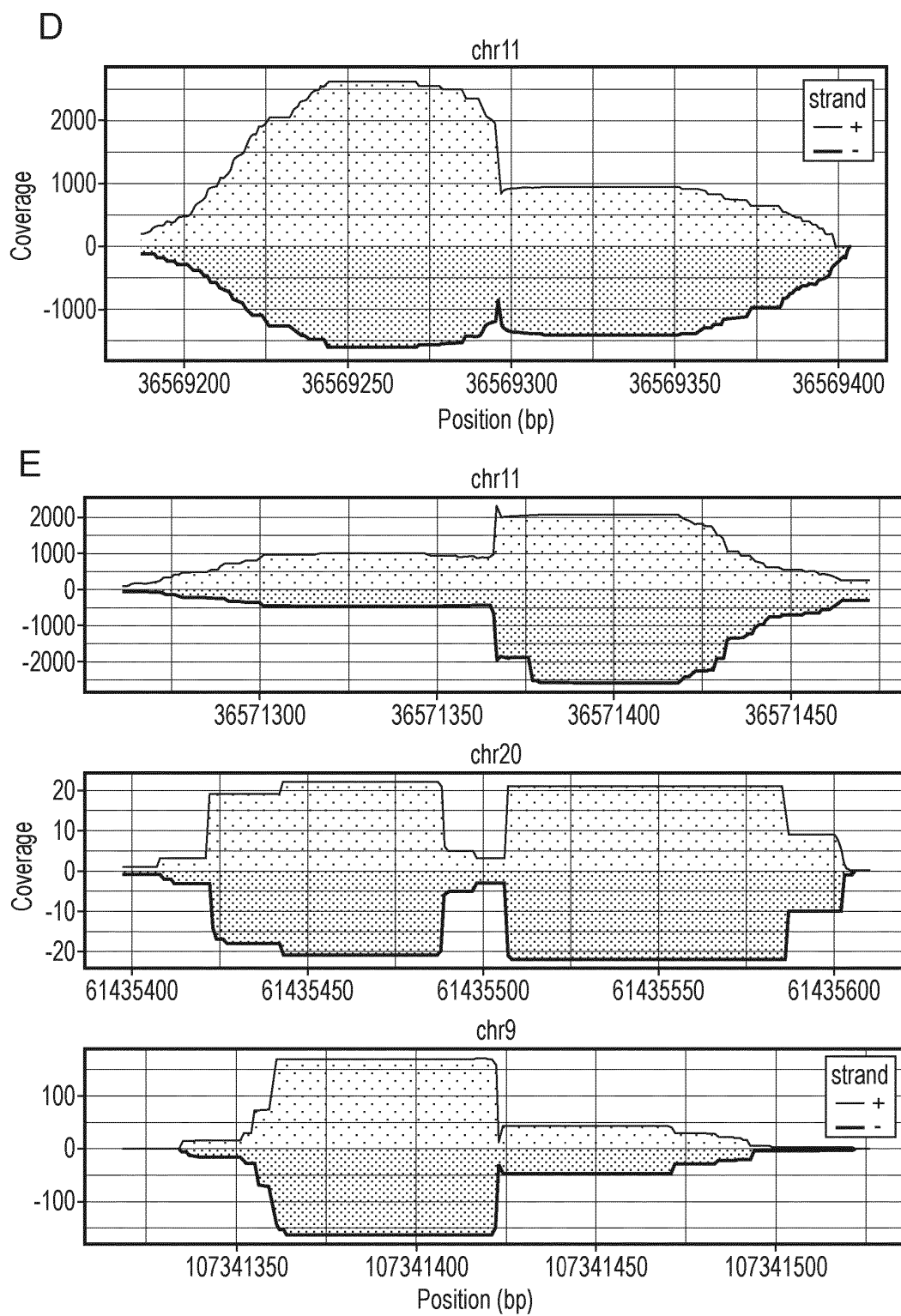


FIG. 4 (Continued)

8/29

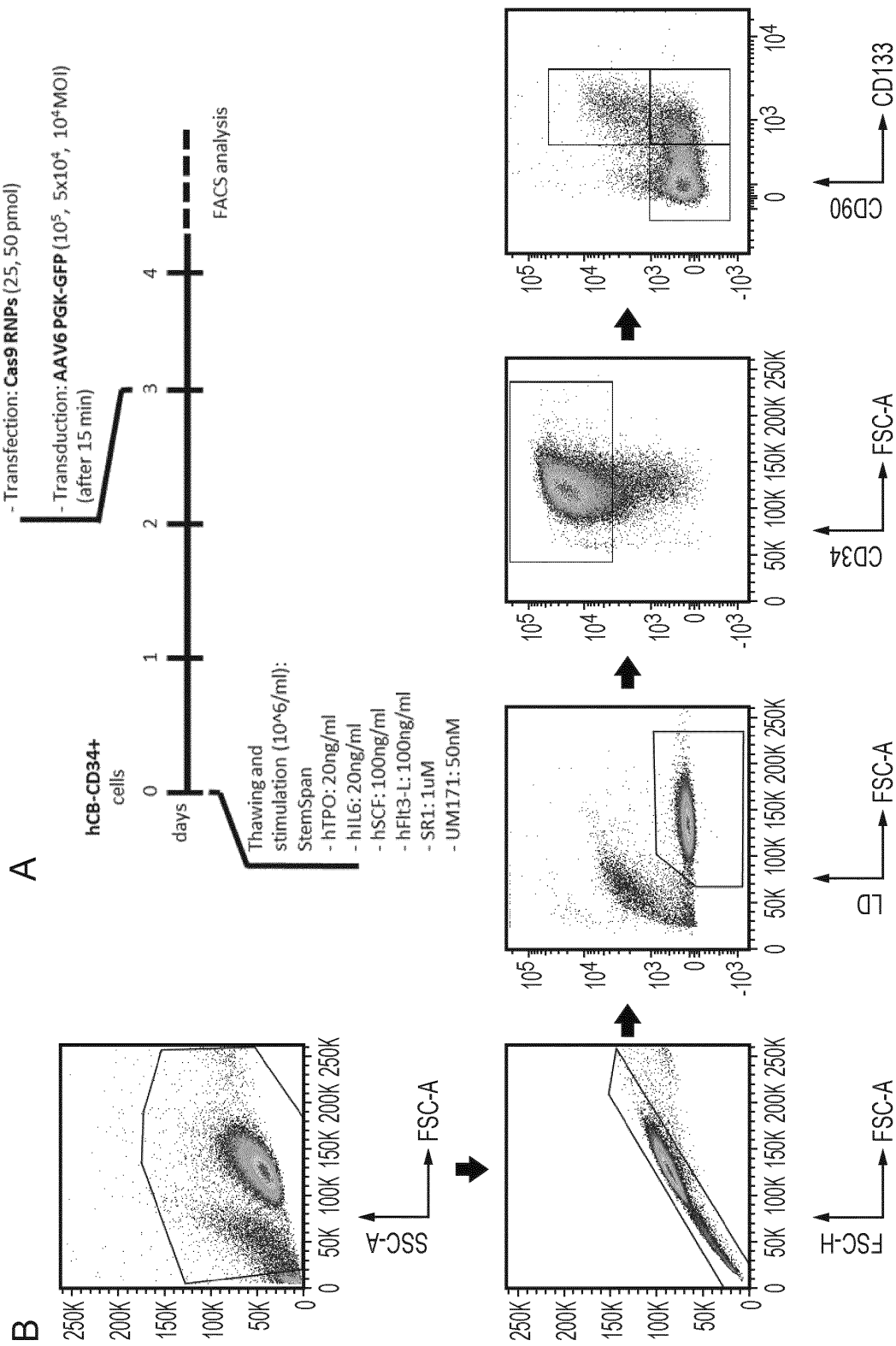


FIG. 5

9/29

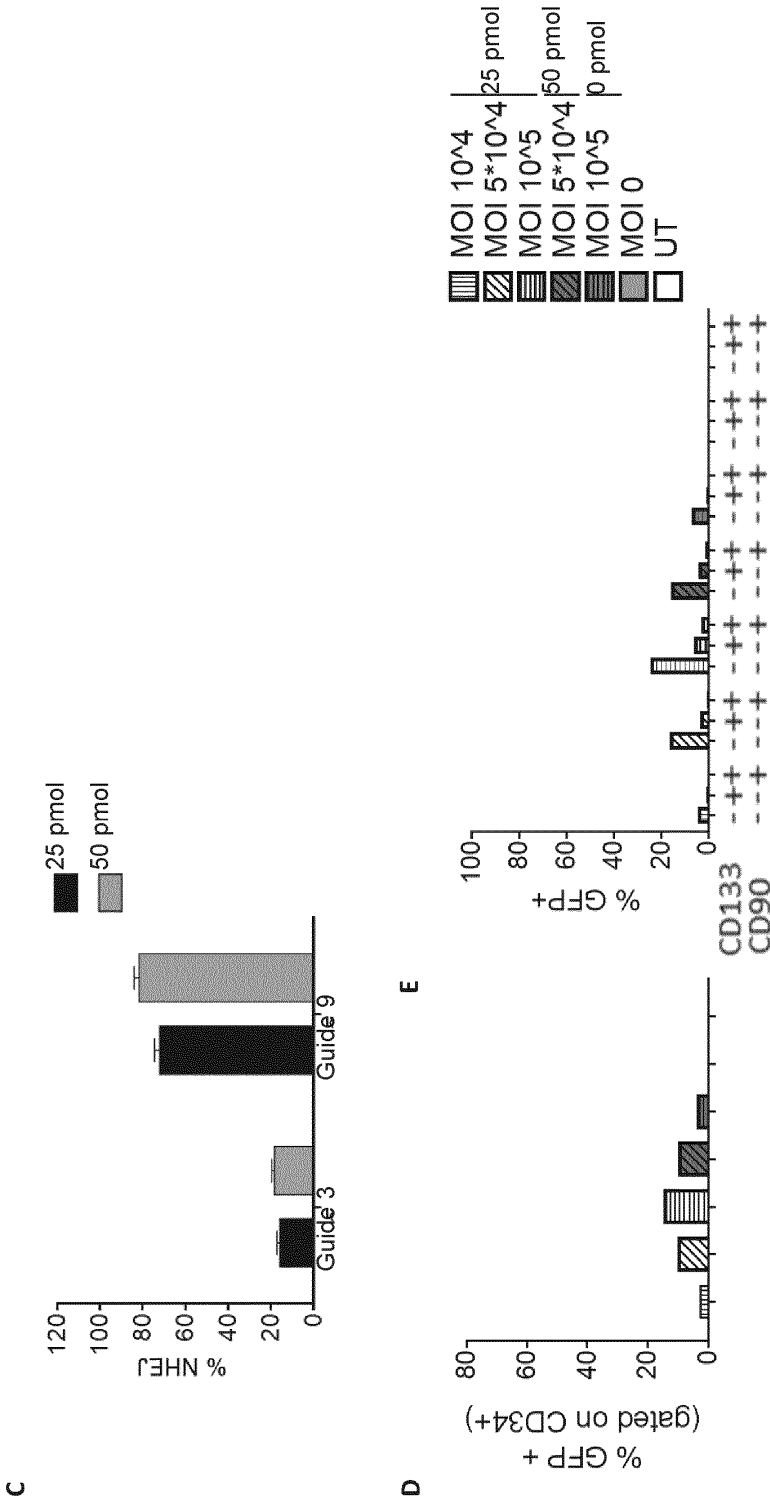


FIG. 5 (Continued)

10/29

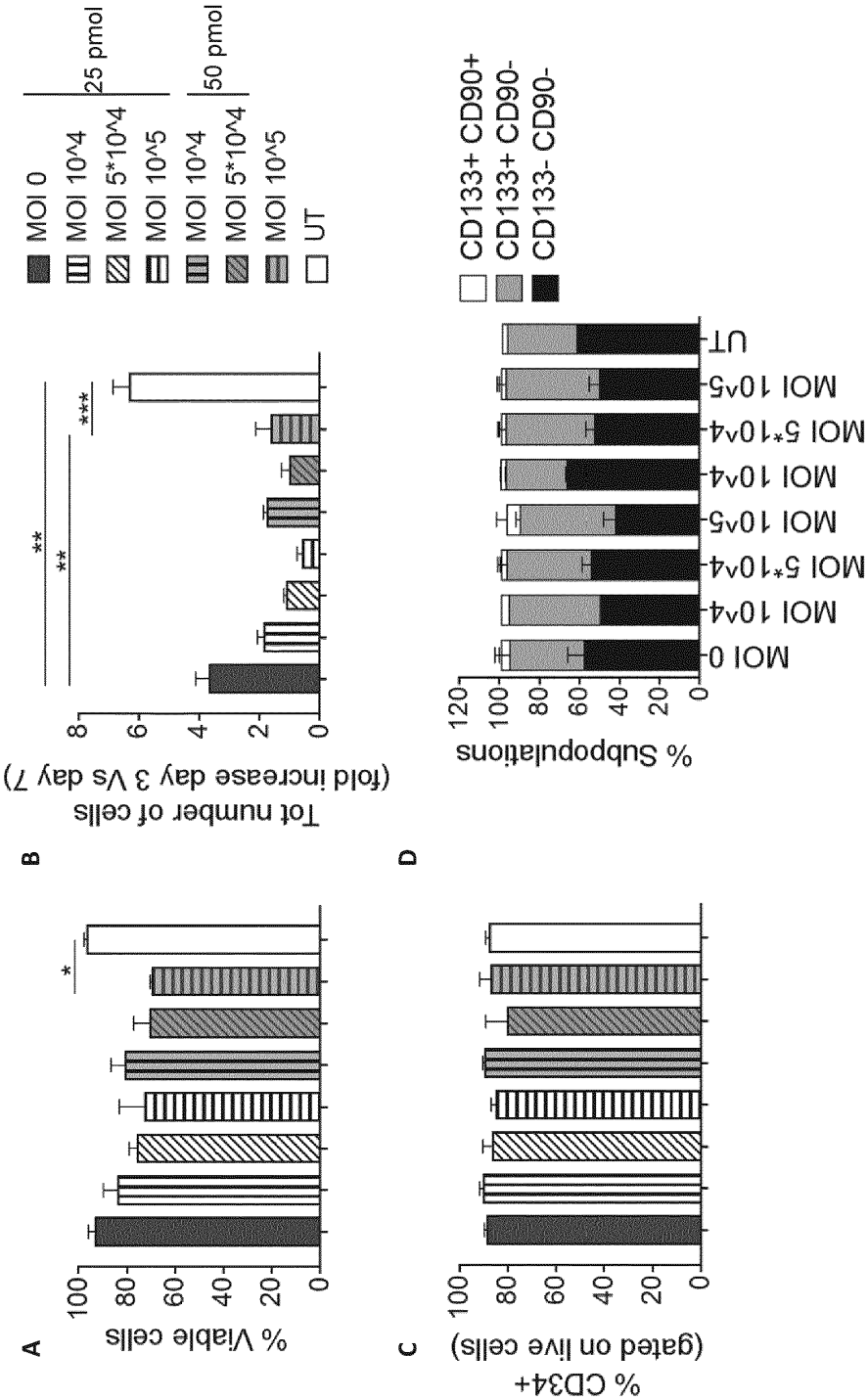
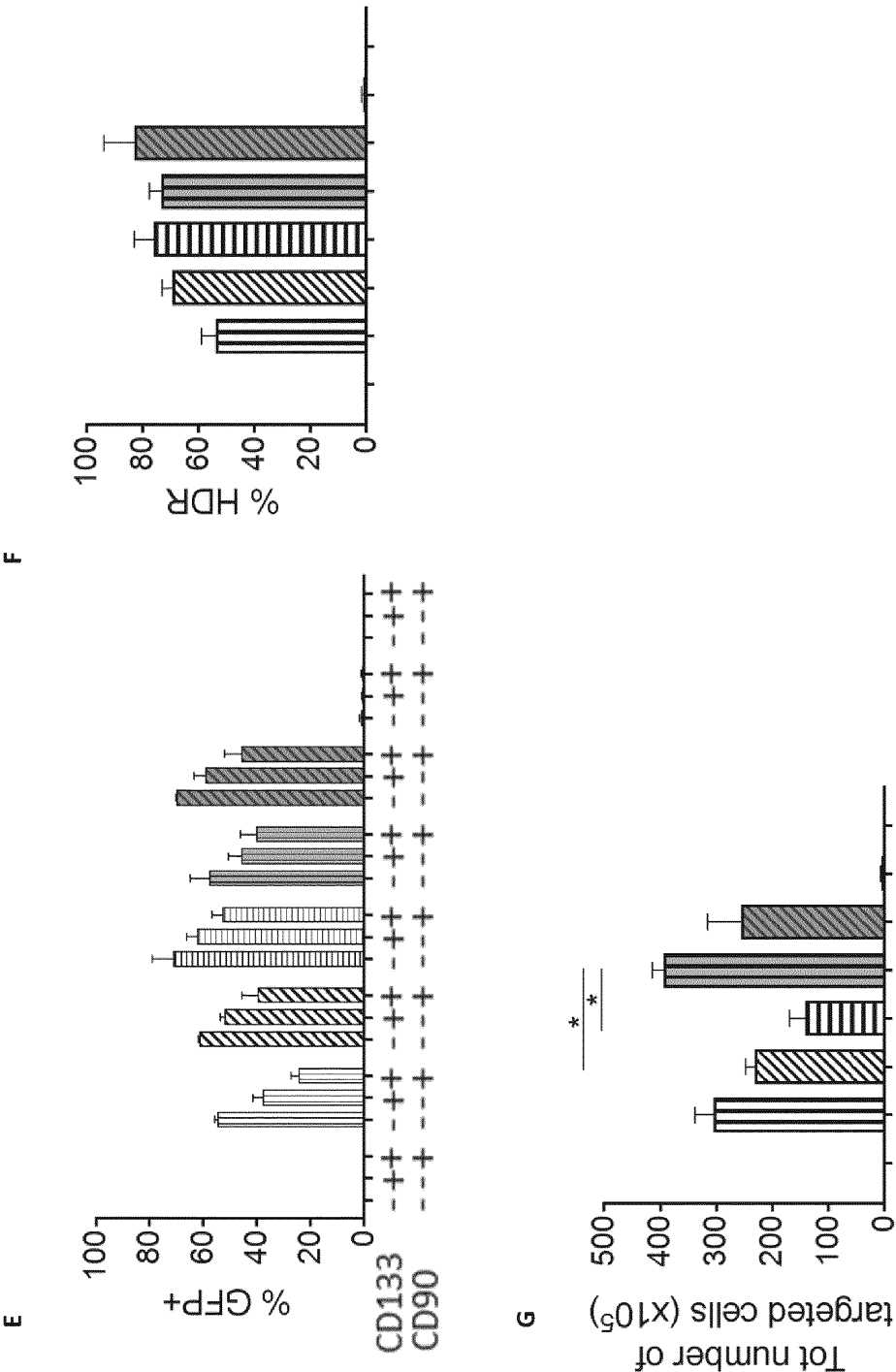


FIG. 6

11/29



12/29

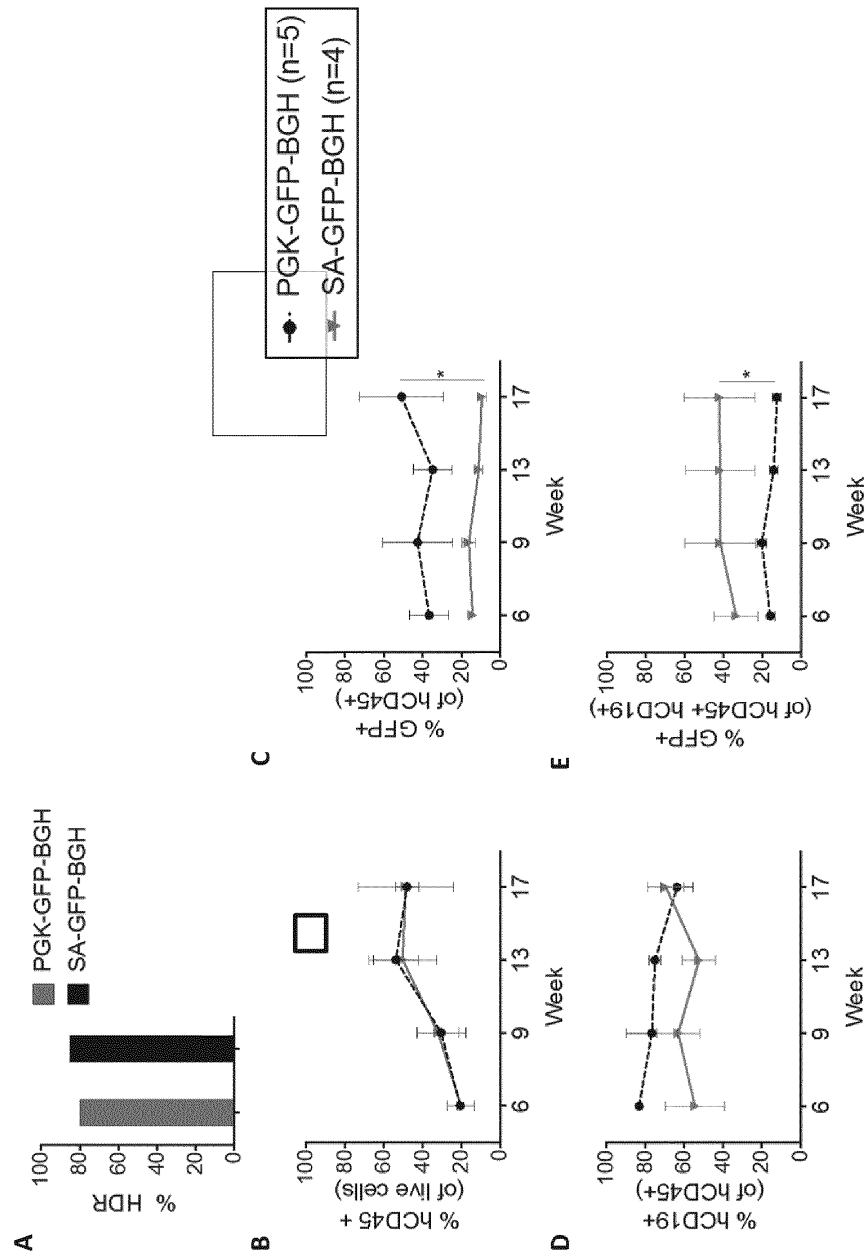


FIG. 7

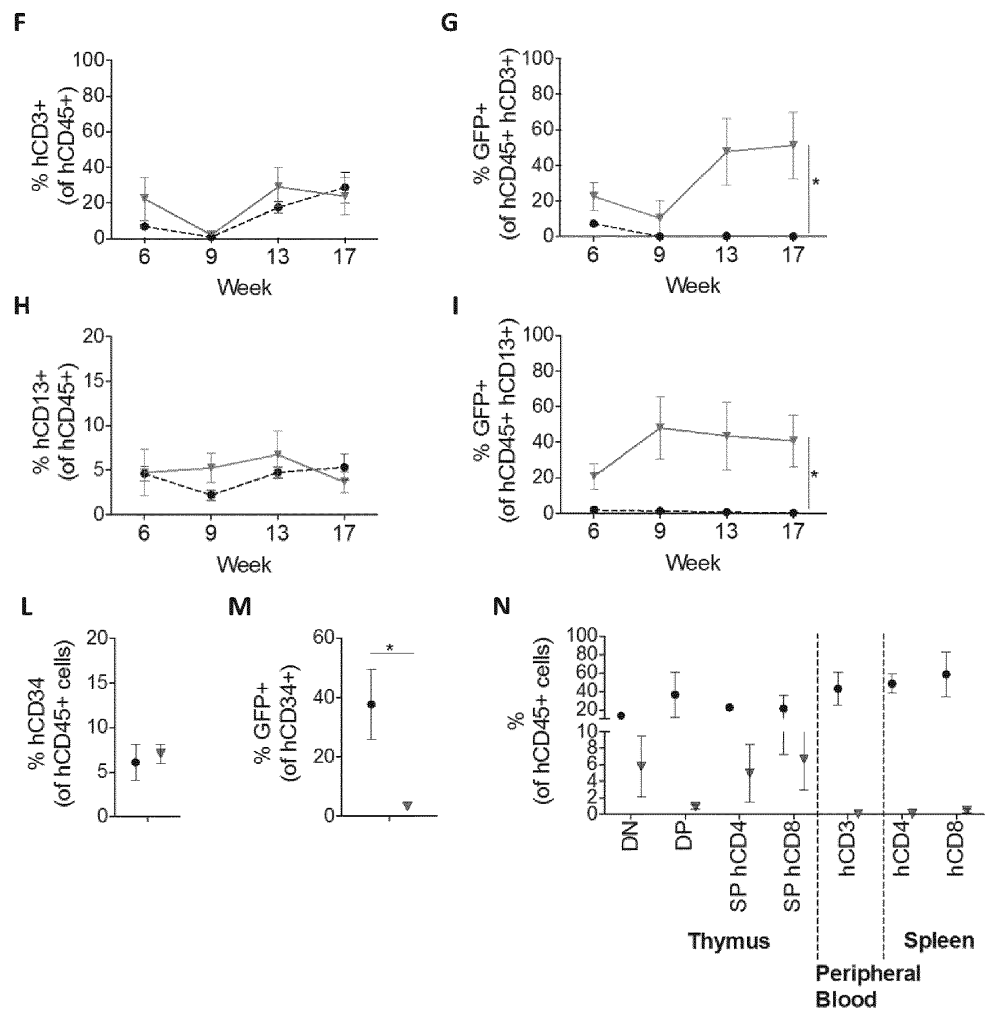


FIG. 7 (Continued)

14/29

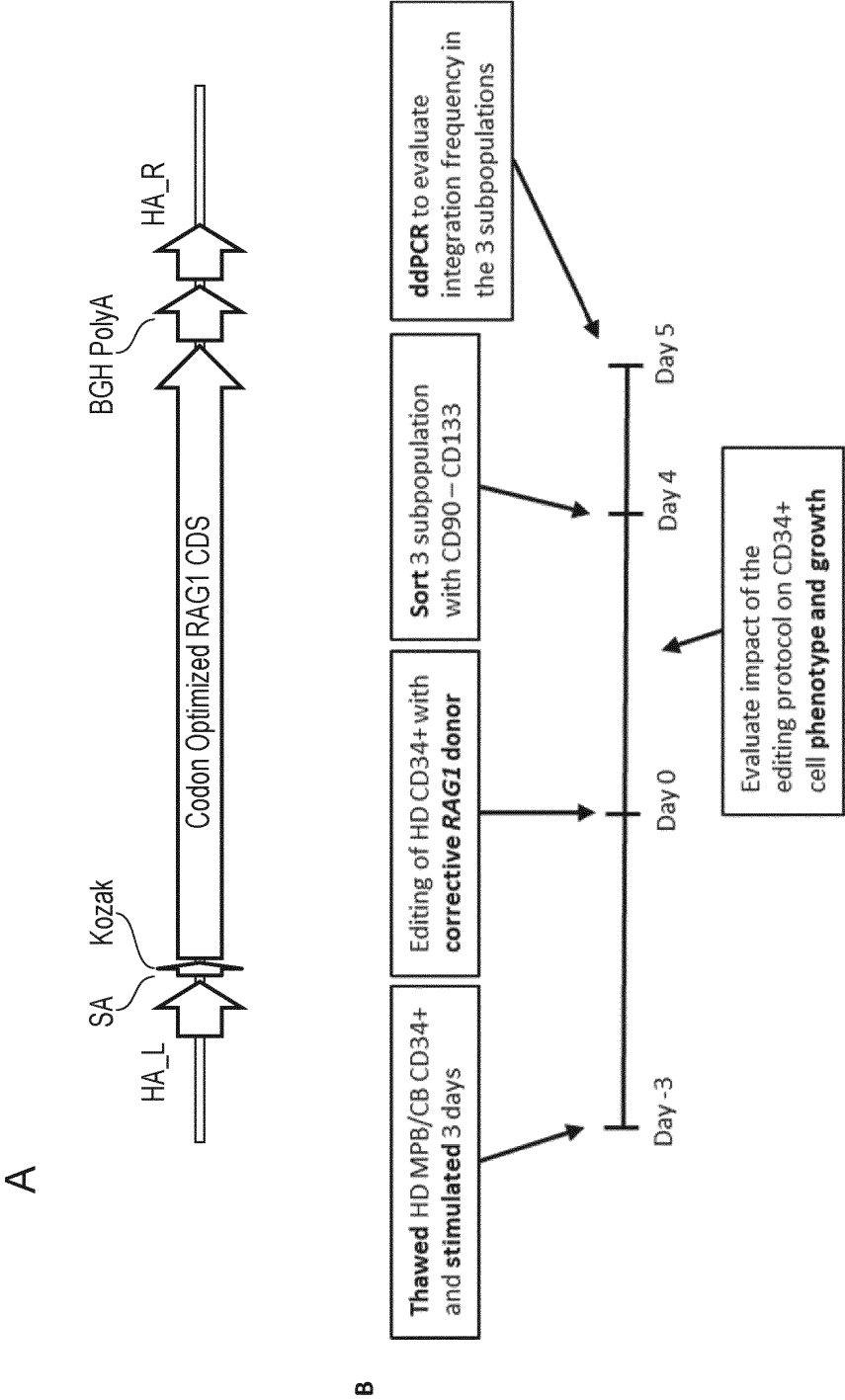


FIG. 8

15/29

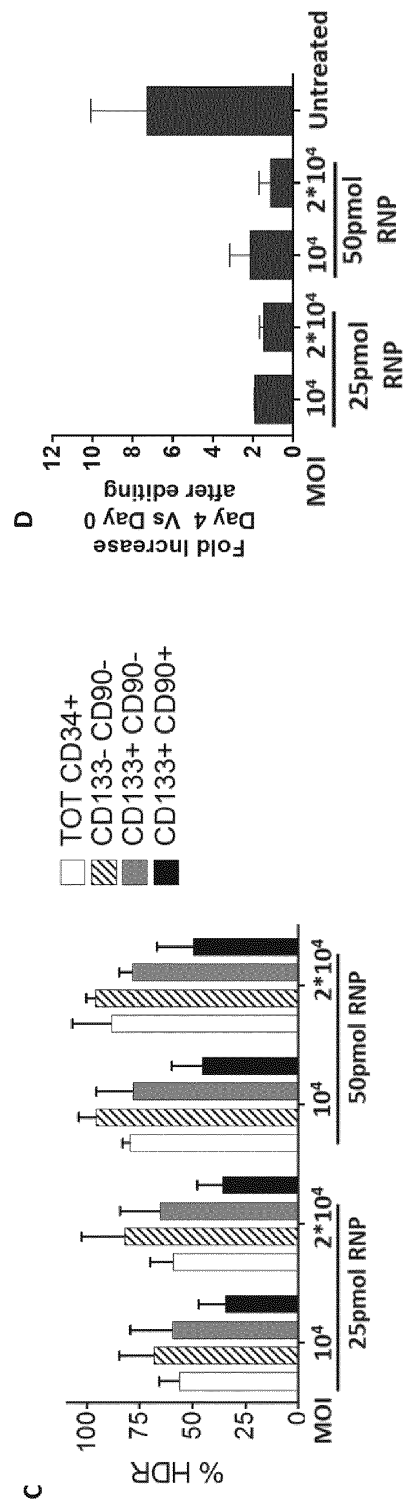


FIG. 8 (Continued)

16/29

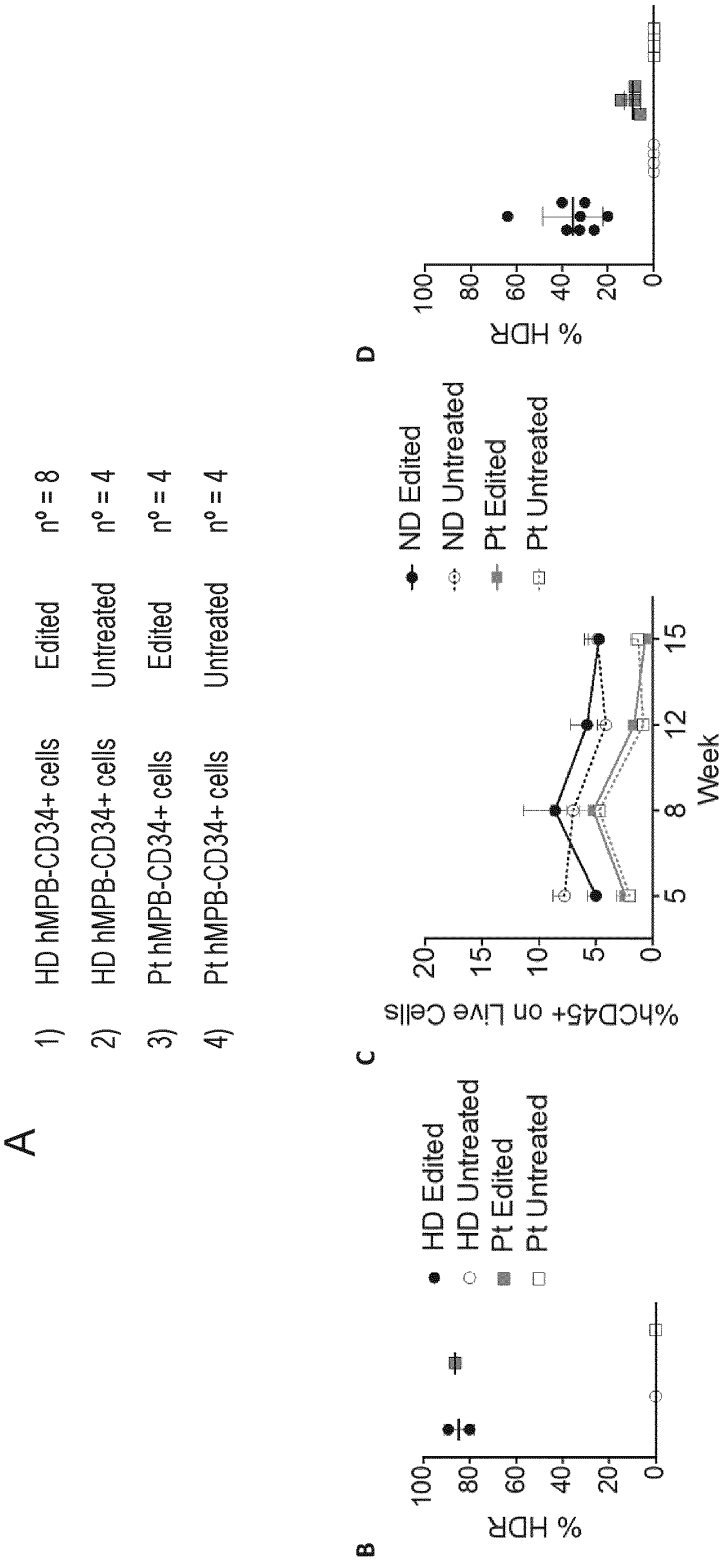


FIG. 9

17/29

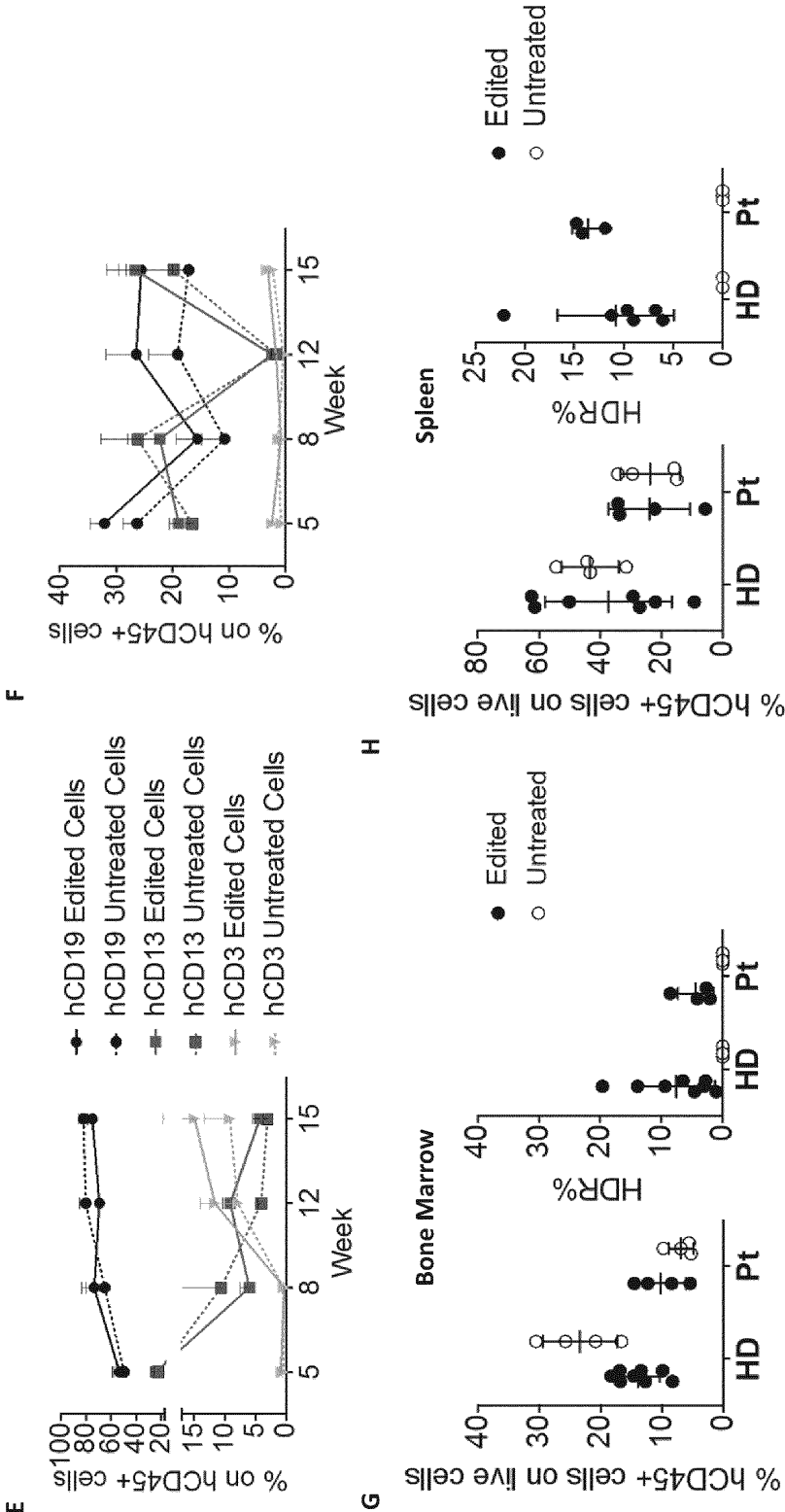


FIG. 9 (Continued)

18/29

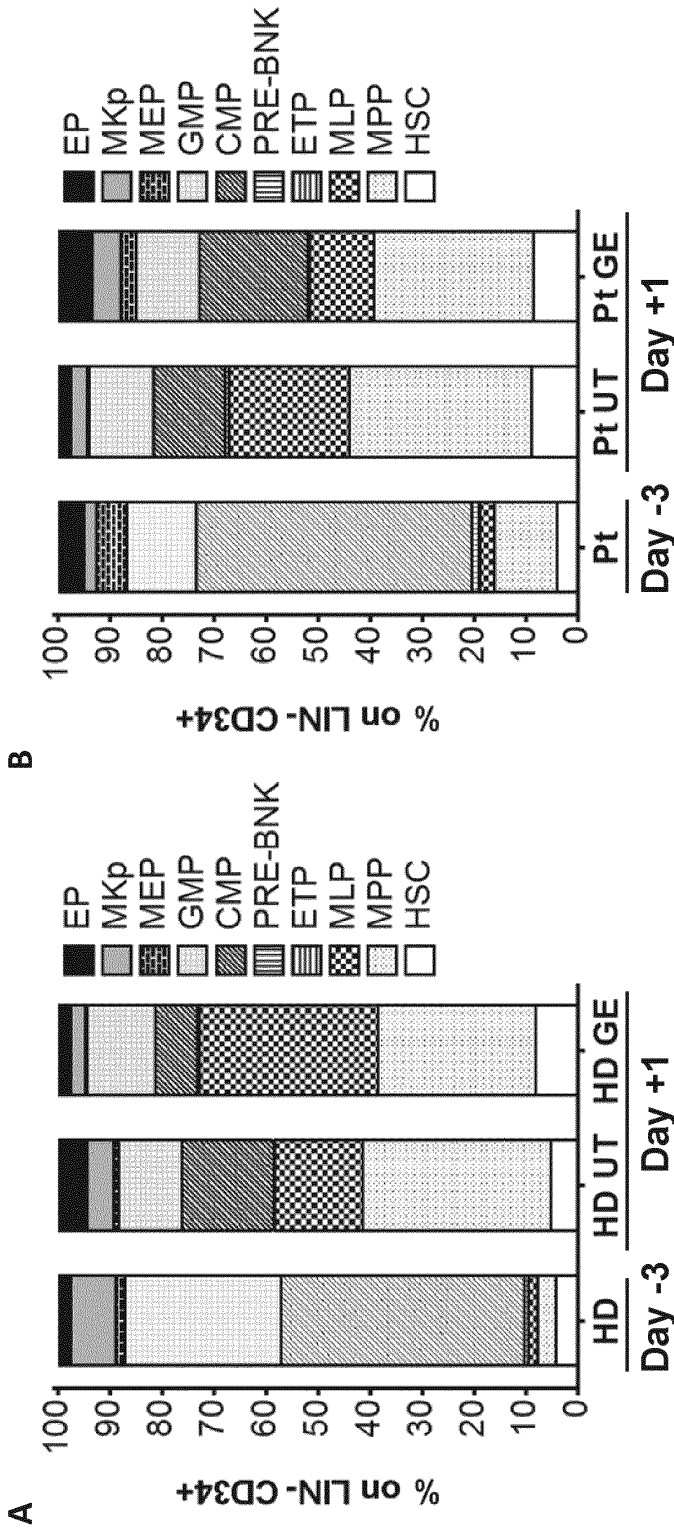


FIG. 10

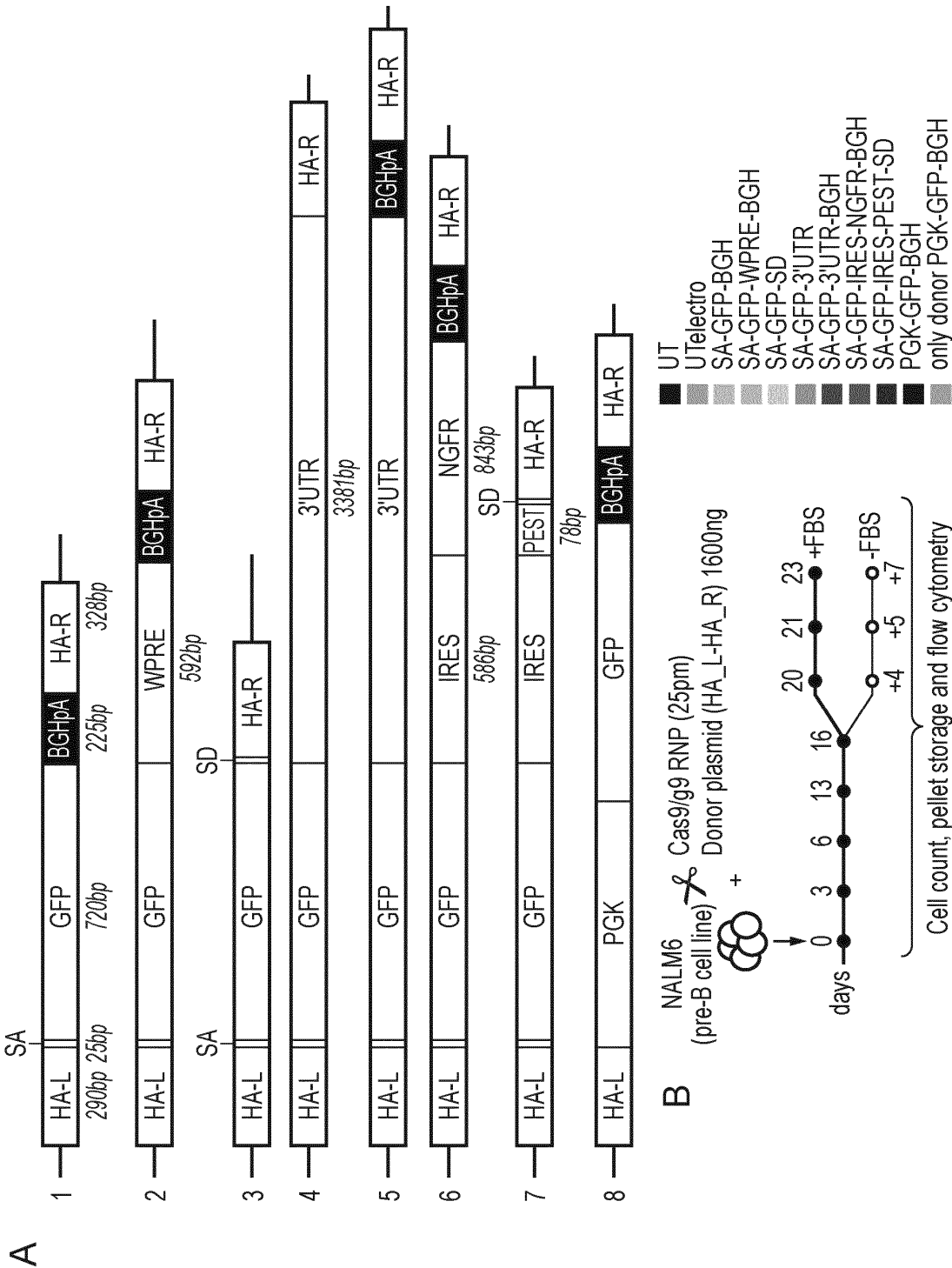
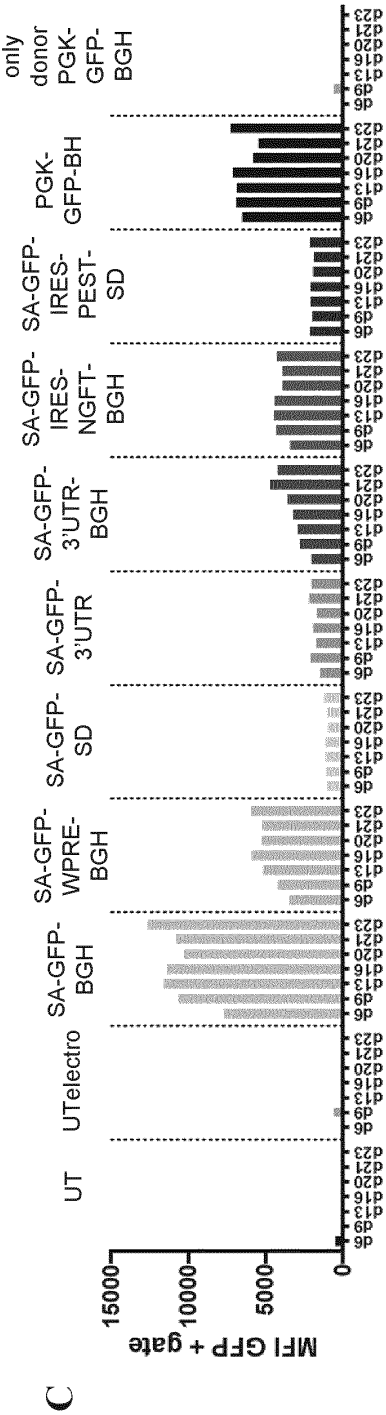


FIG. 11



21/29

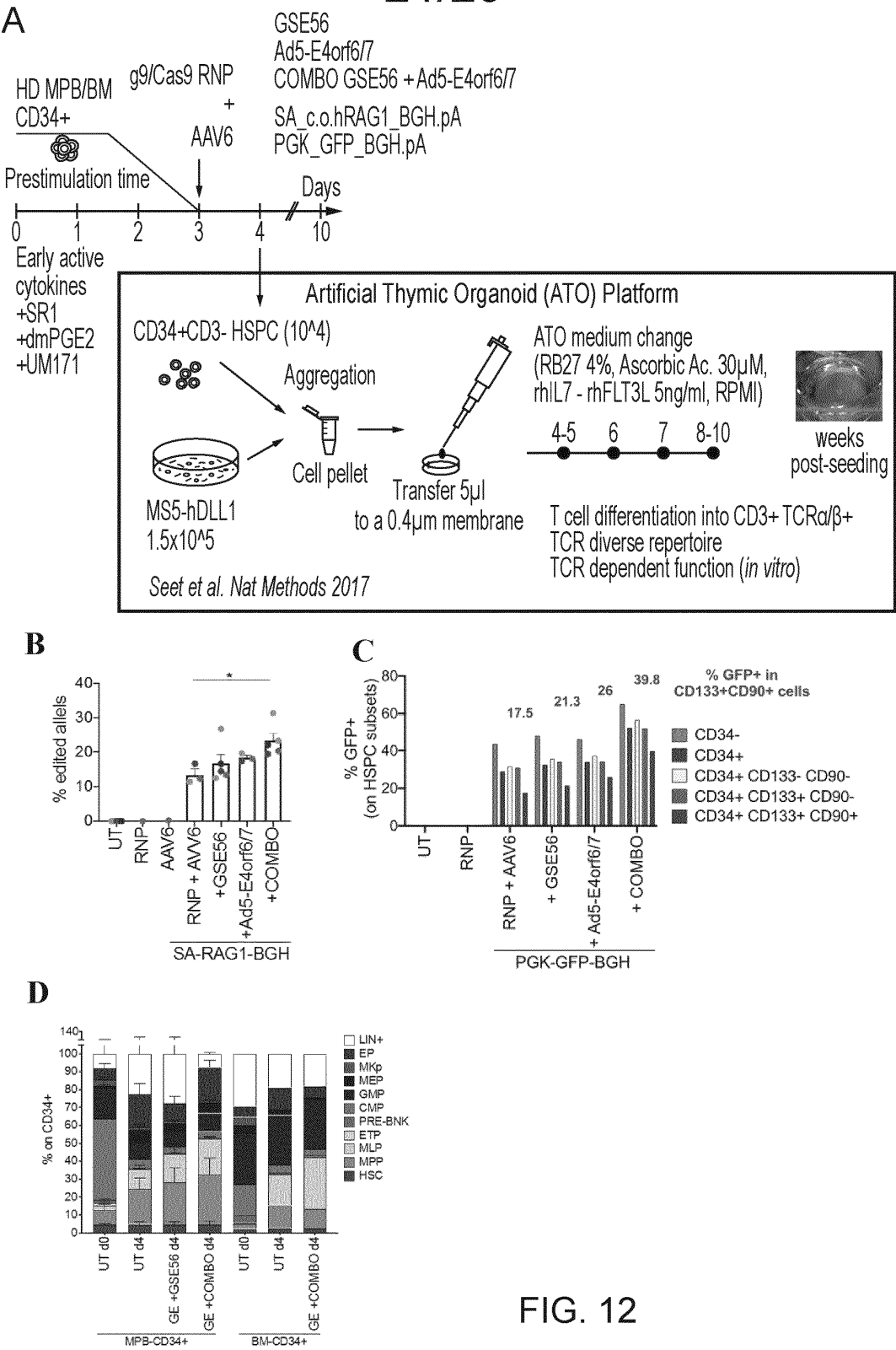


FIG. 12

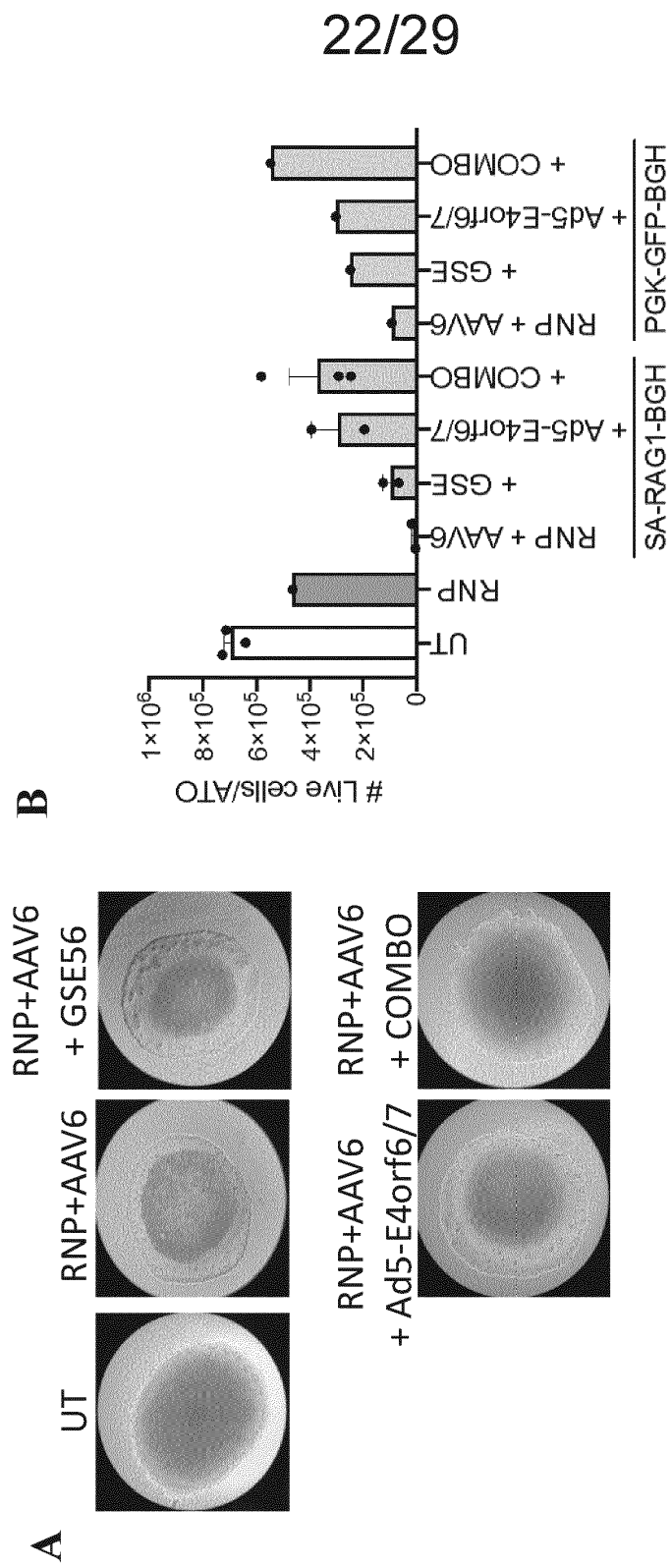


FIG. 13

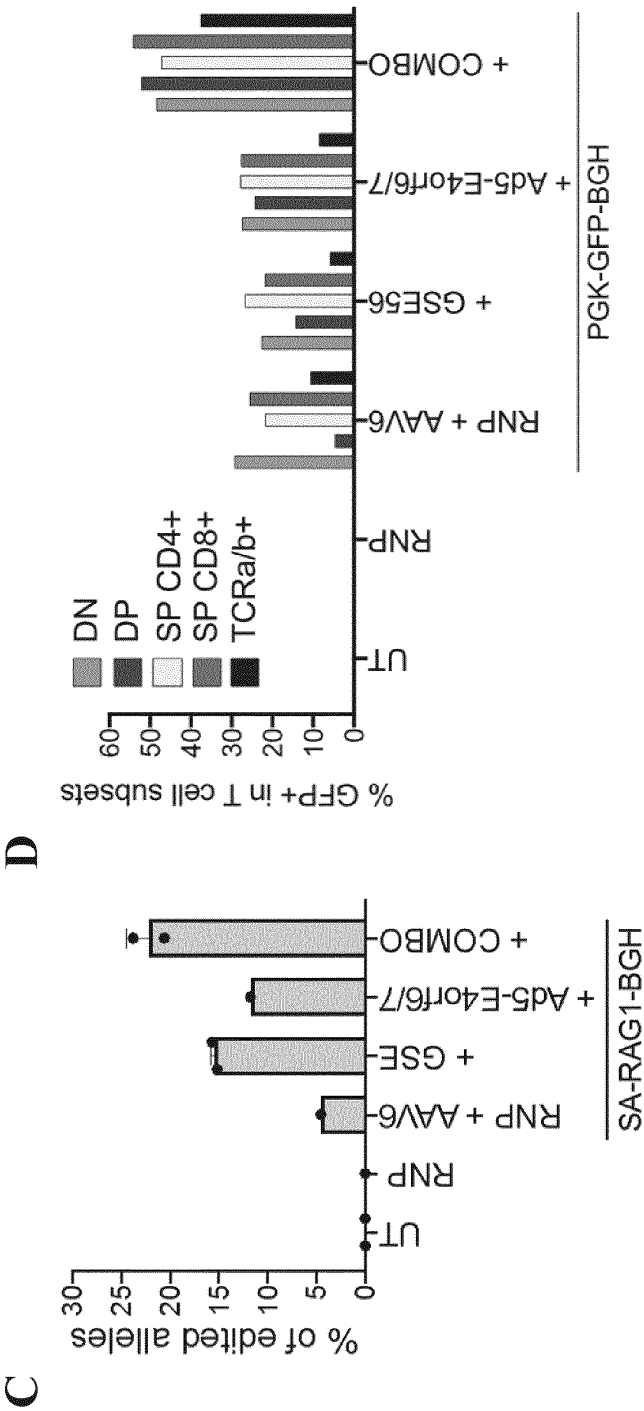


FIG. 13 (Continued)

24/29

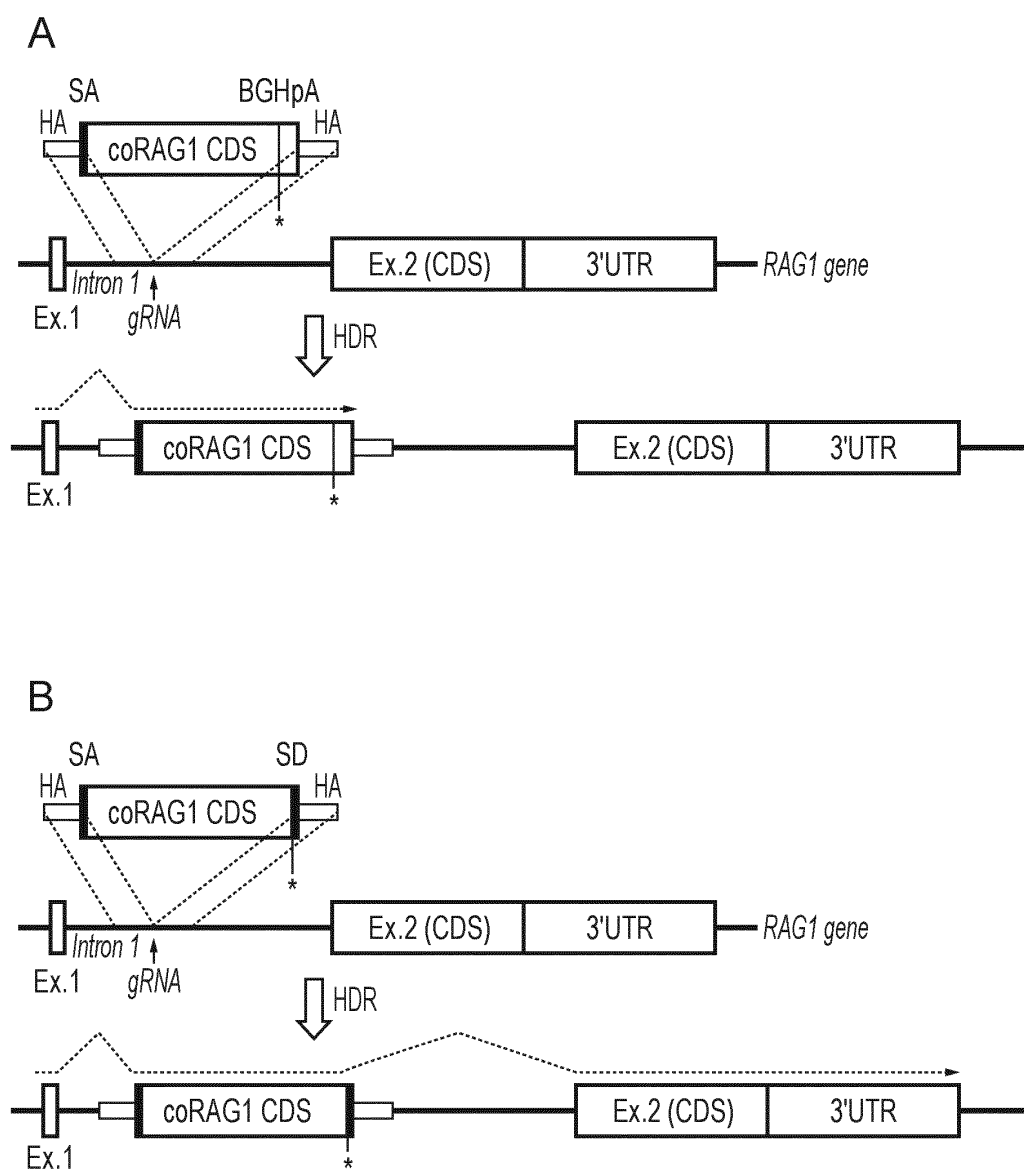


FIG. 14

25/29

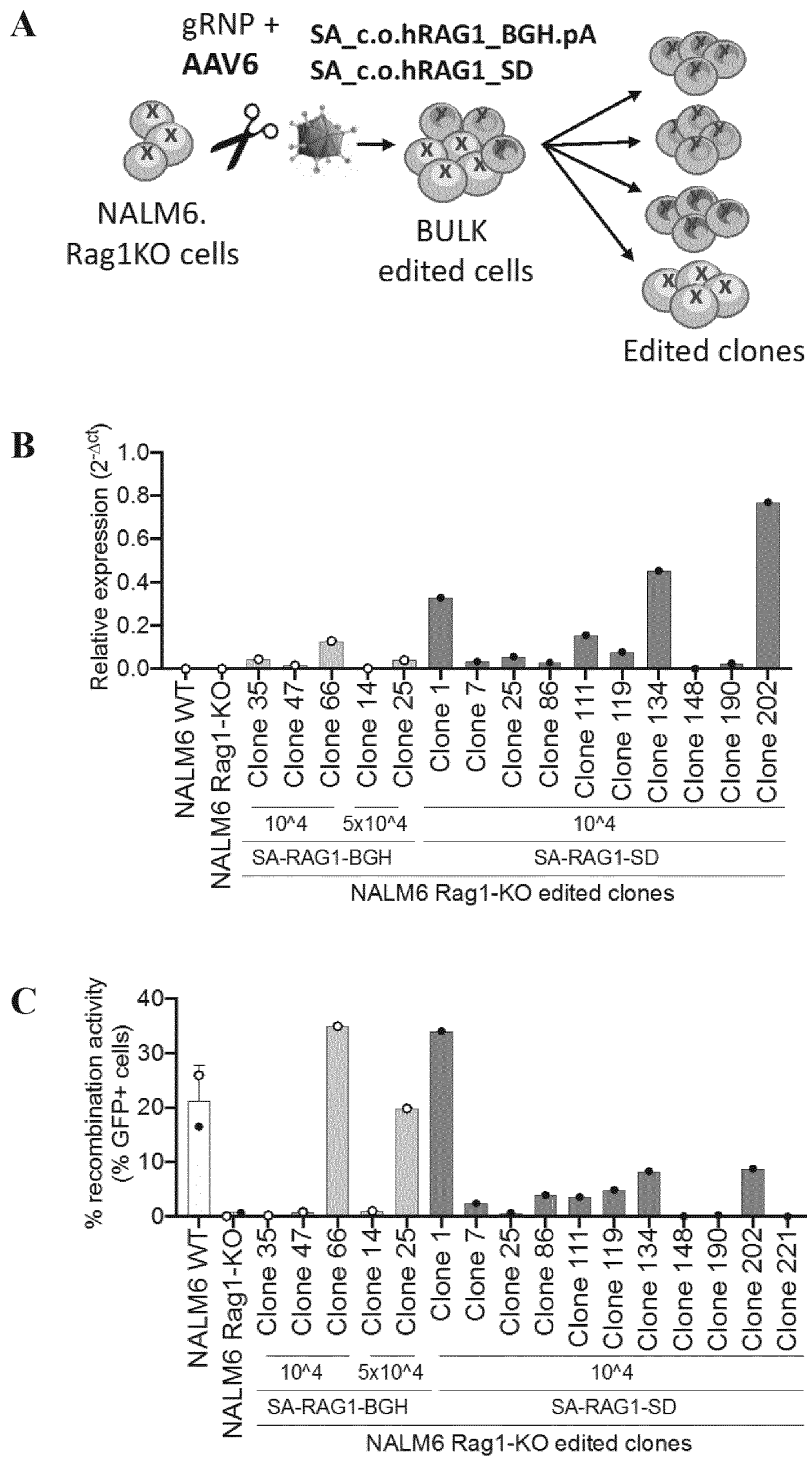


FIG. 15

SUBSTITUTE SHEET (RULE 26)

26/29

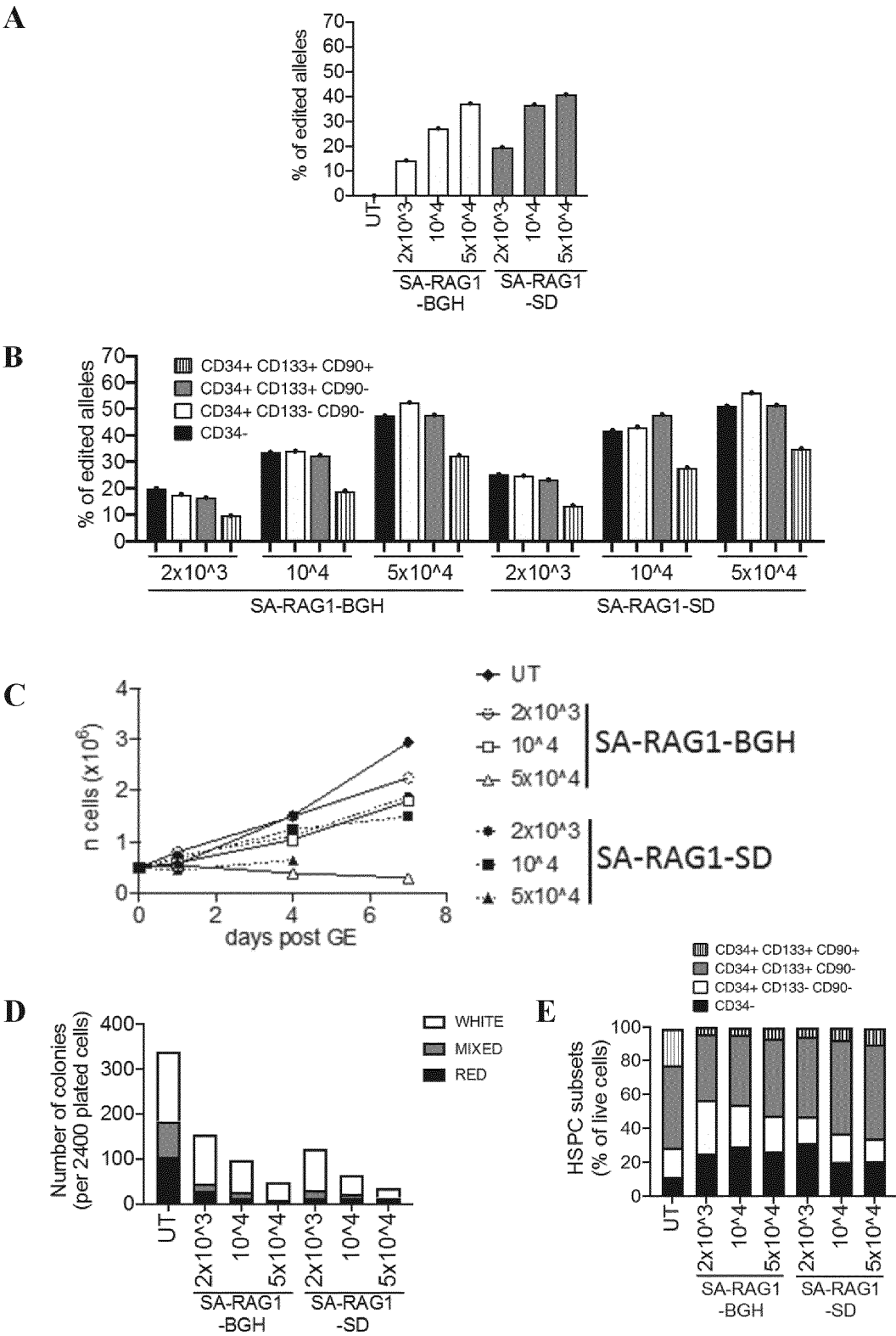


FIG. 16

SUBSTITUTE SHEET (RULE 26)

27/29

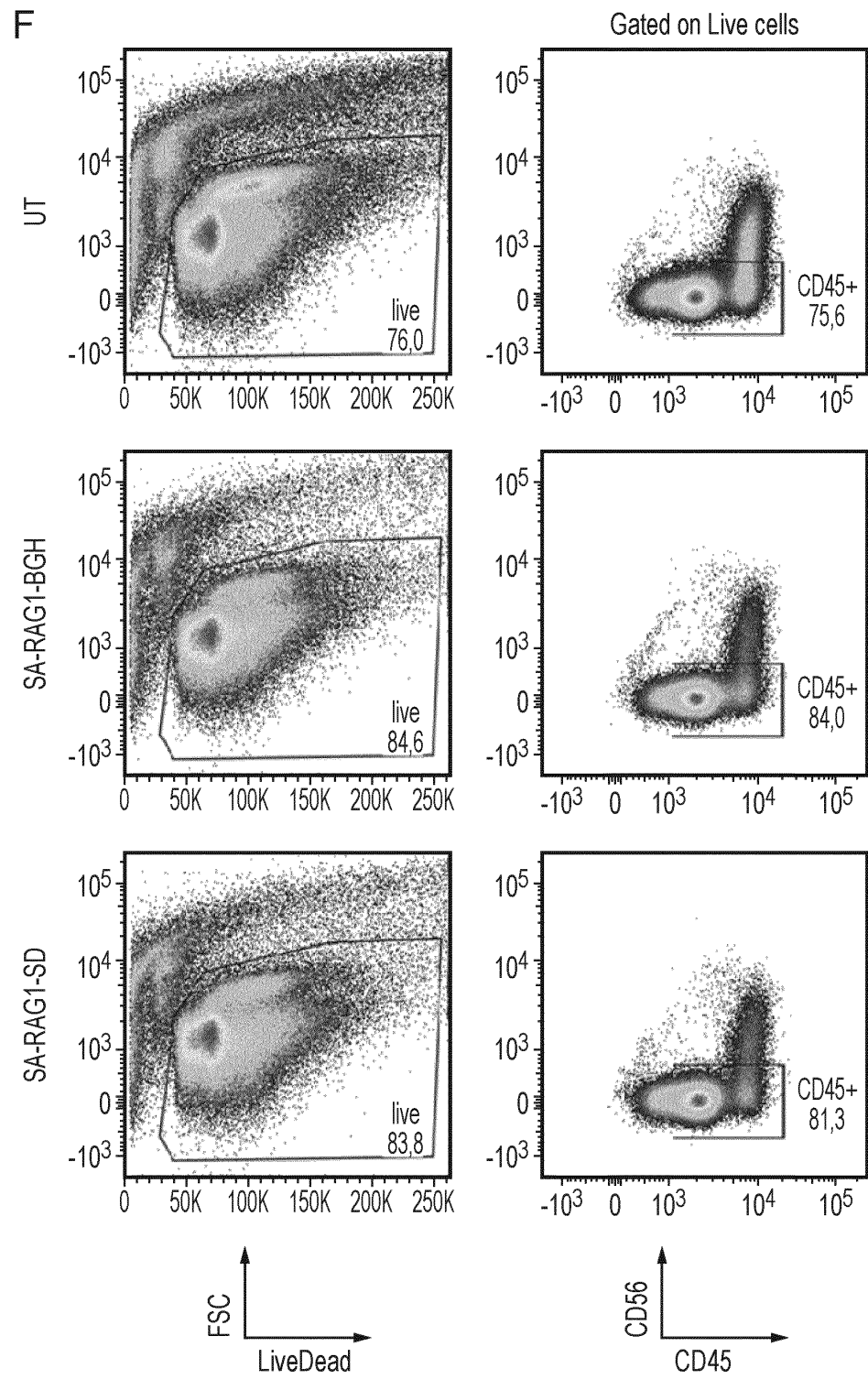


FIG. 16 (Continued)

SUBSTITUTE SHEET (RULE 26)

28/29

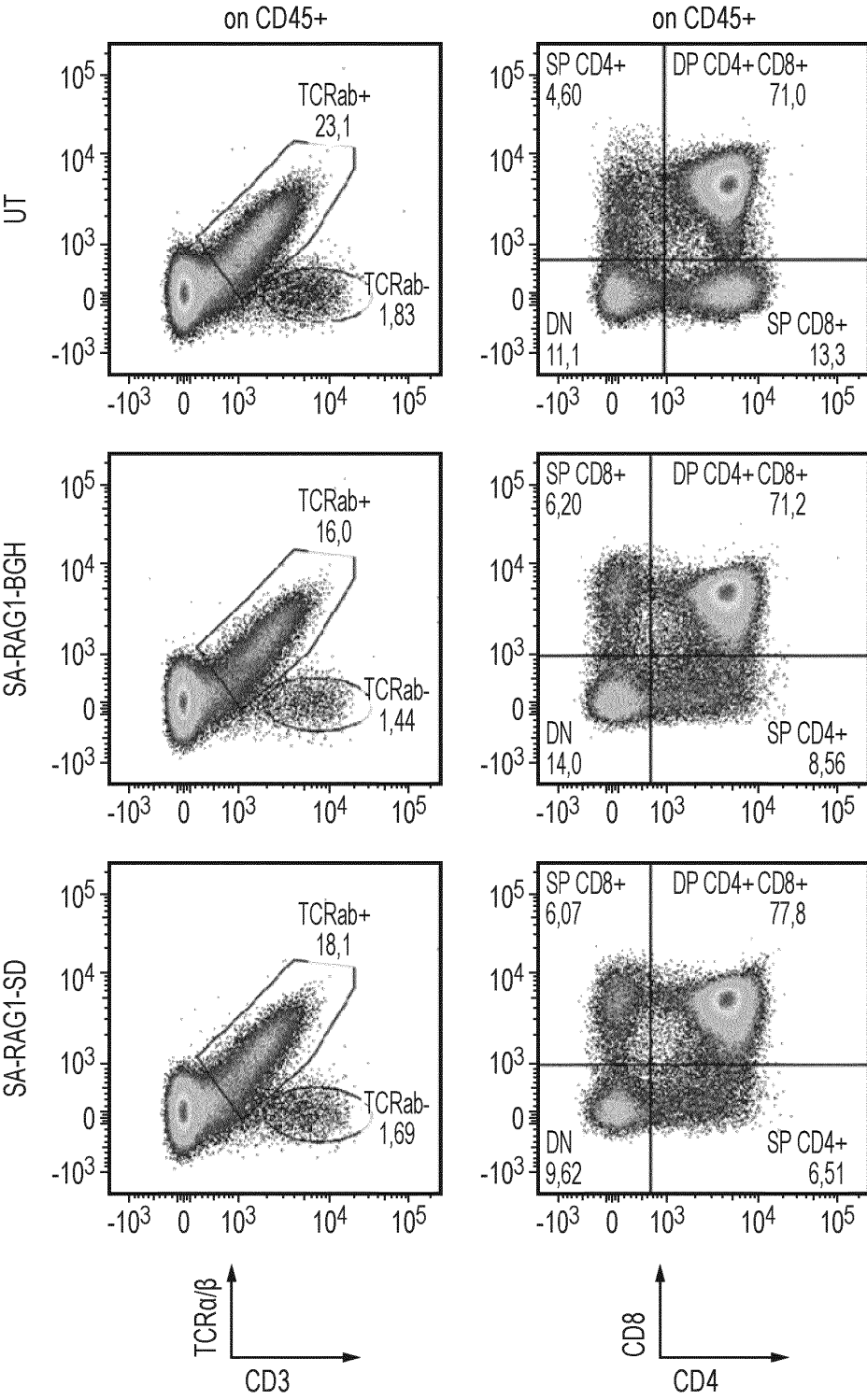


FIG. 16 (Continued)

SUBSTITUTE SHEET (RULE 26)

29/29

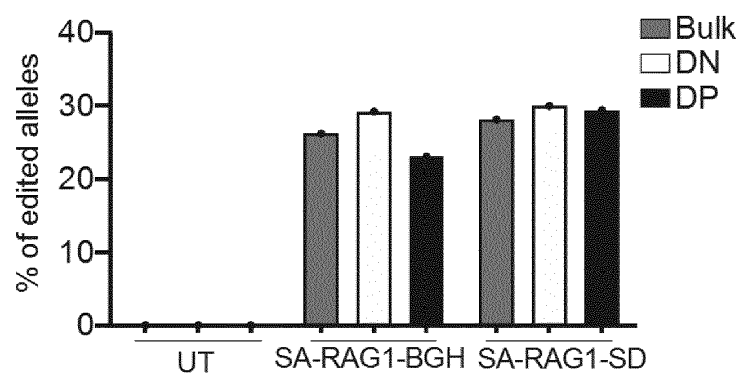
G

FIG. 16 (Continued)