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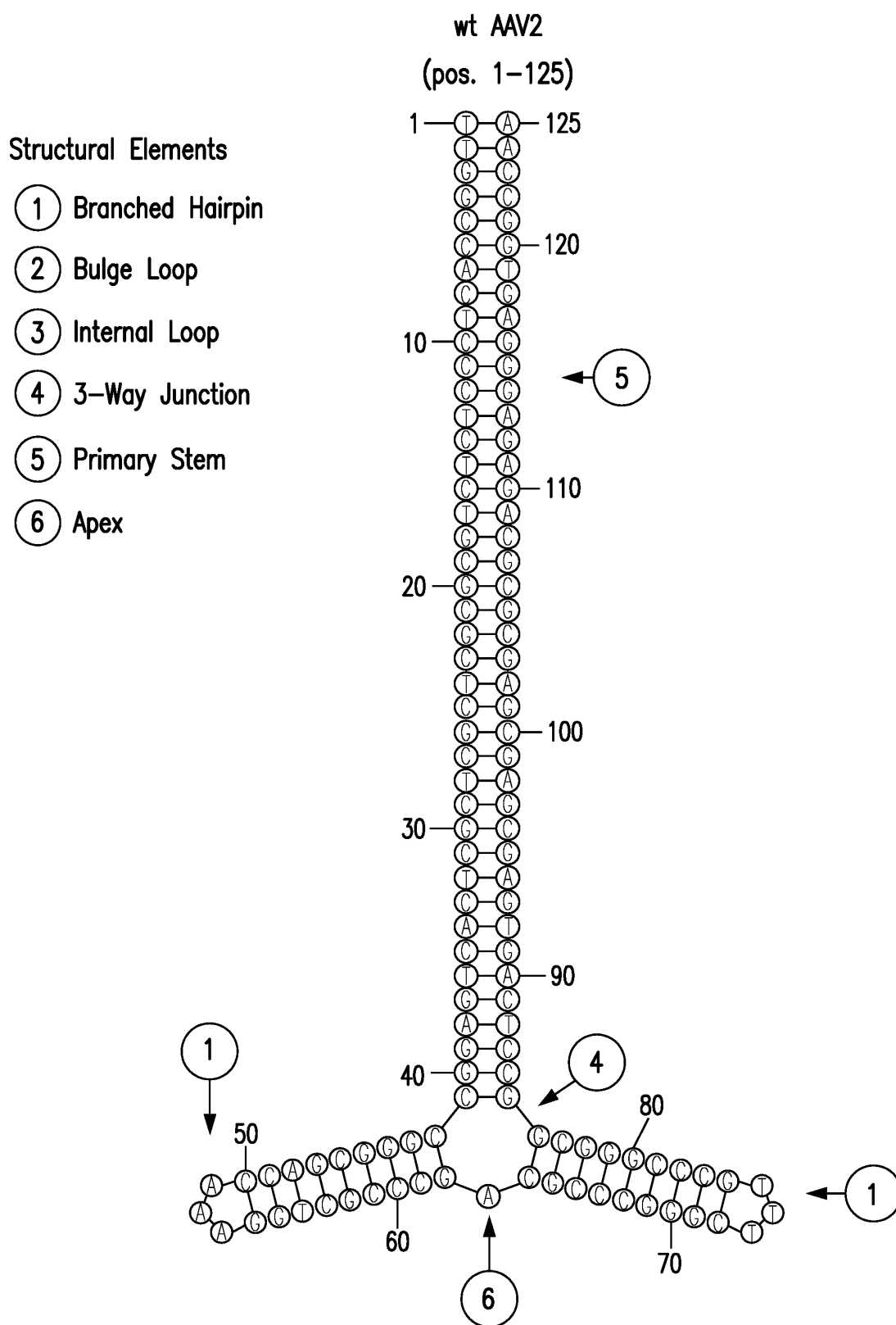


FIG. 1
SUBSTITUTE SHEET (RULE 26)

(pos. 129-237)



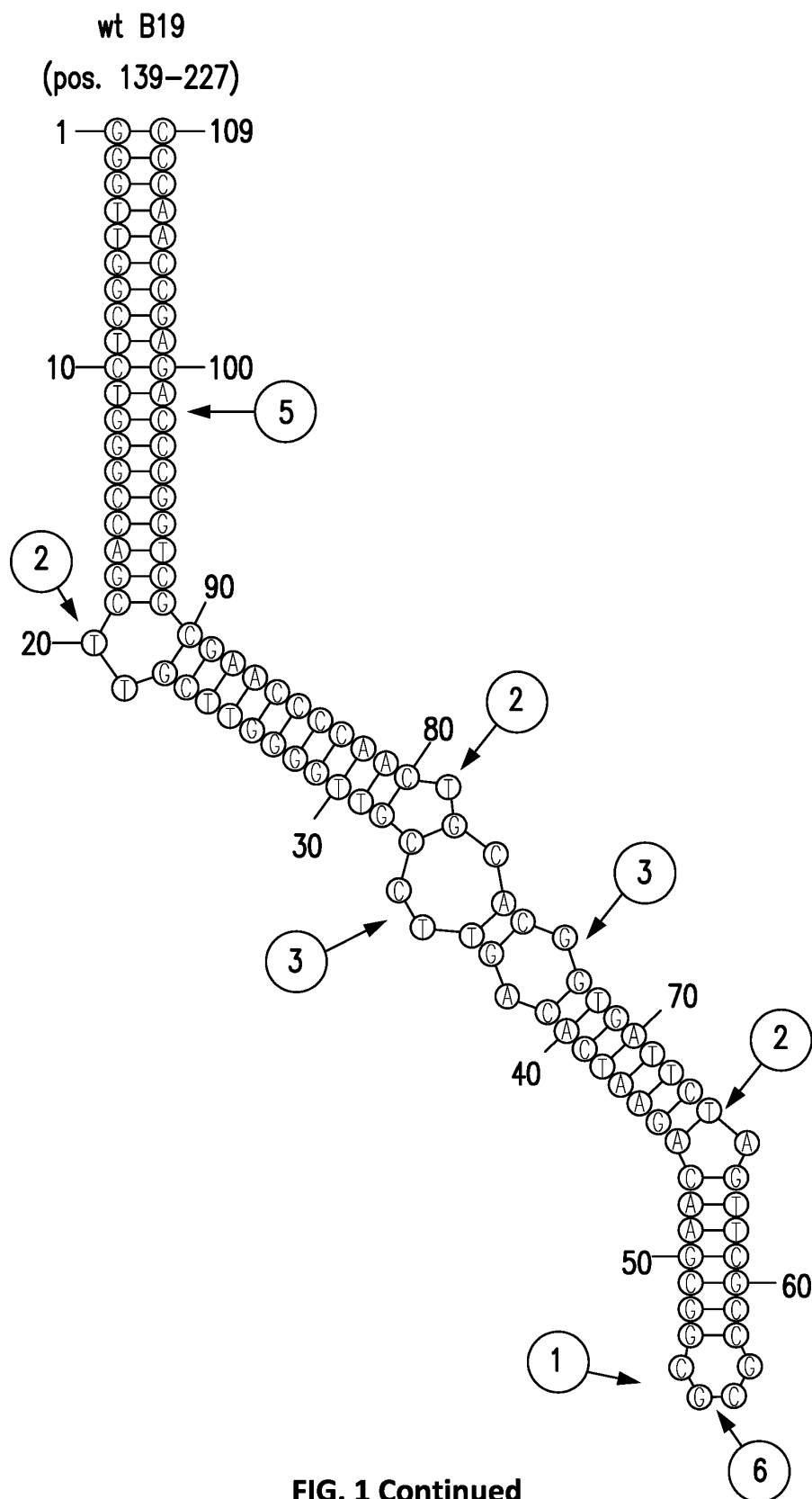
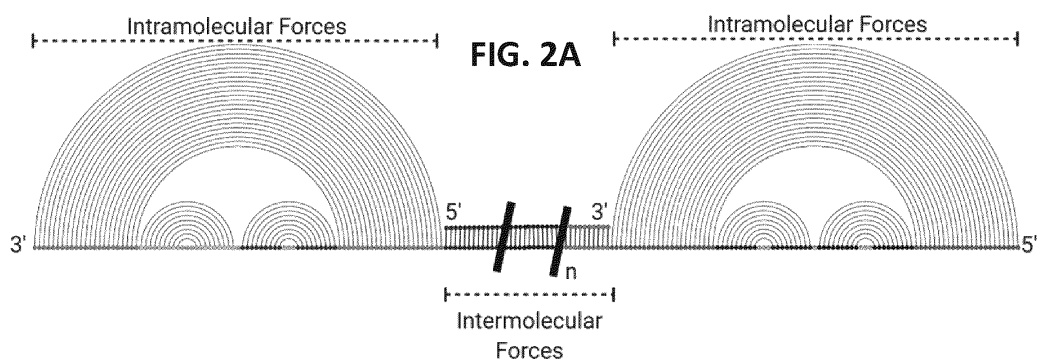
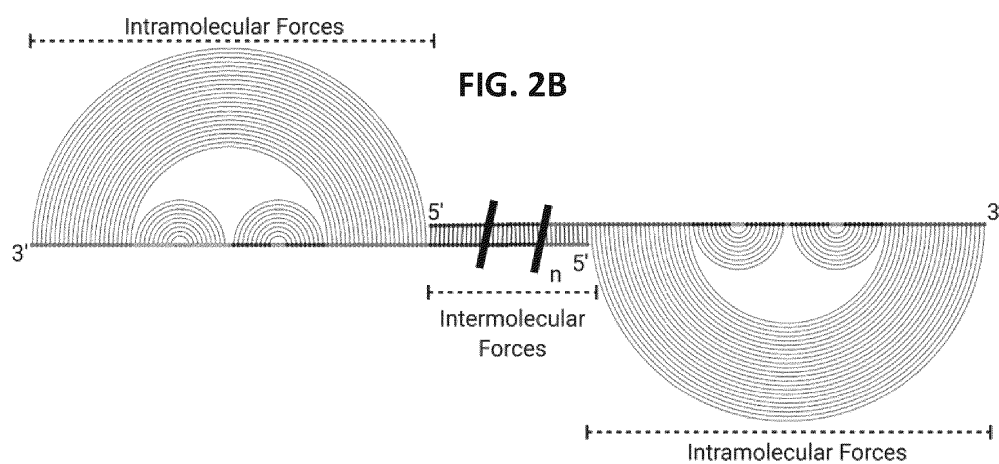
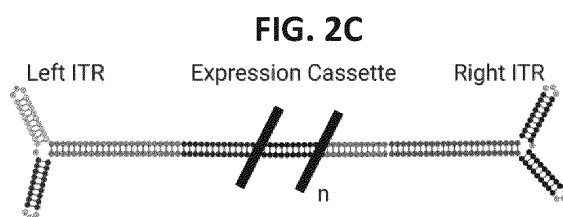


FIG. 1 Continued
SUBSTITUTE SHEET (RULE 26)

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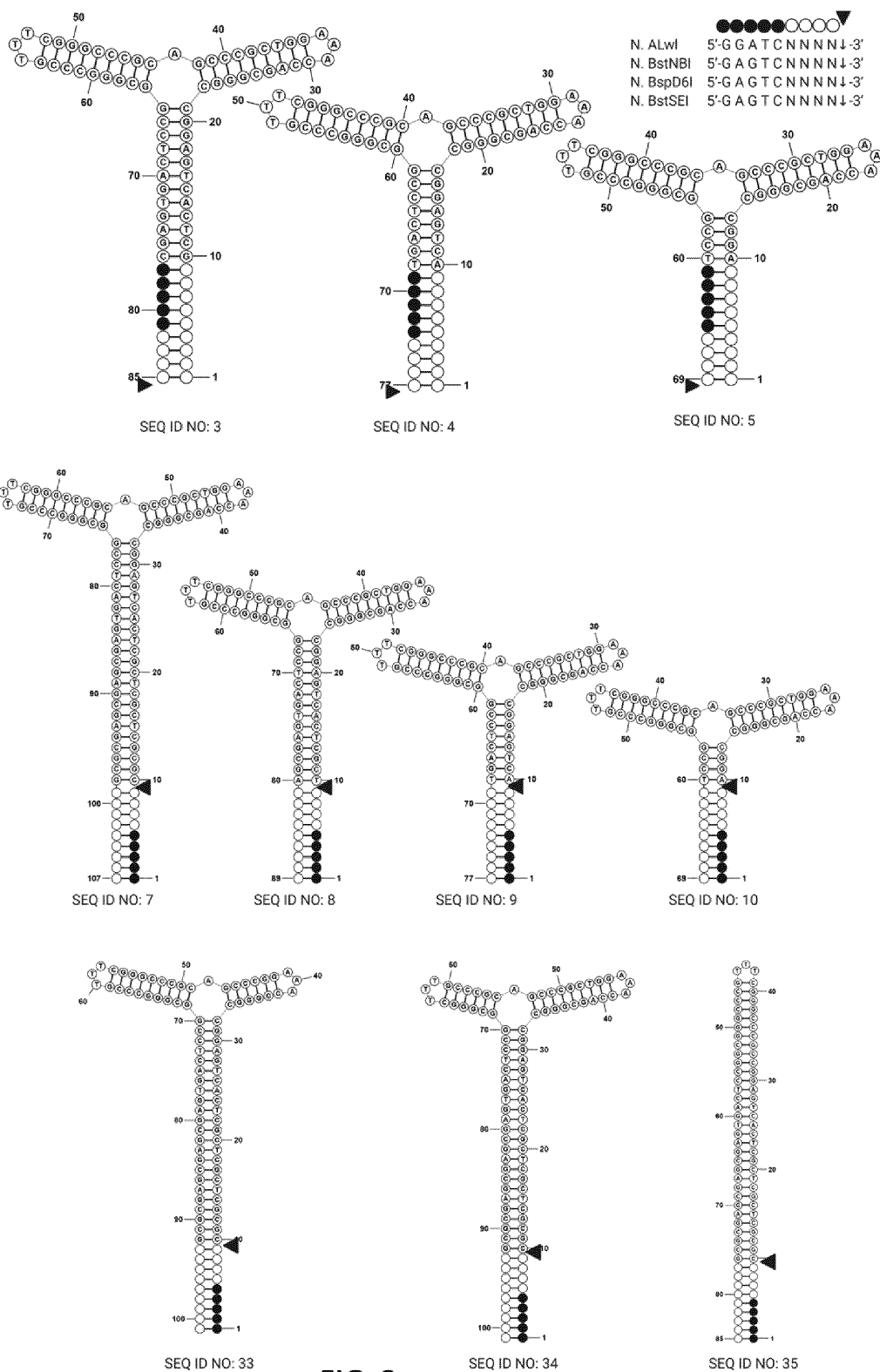
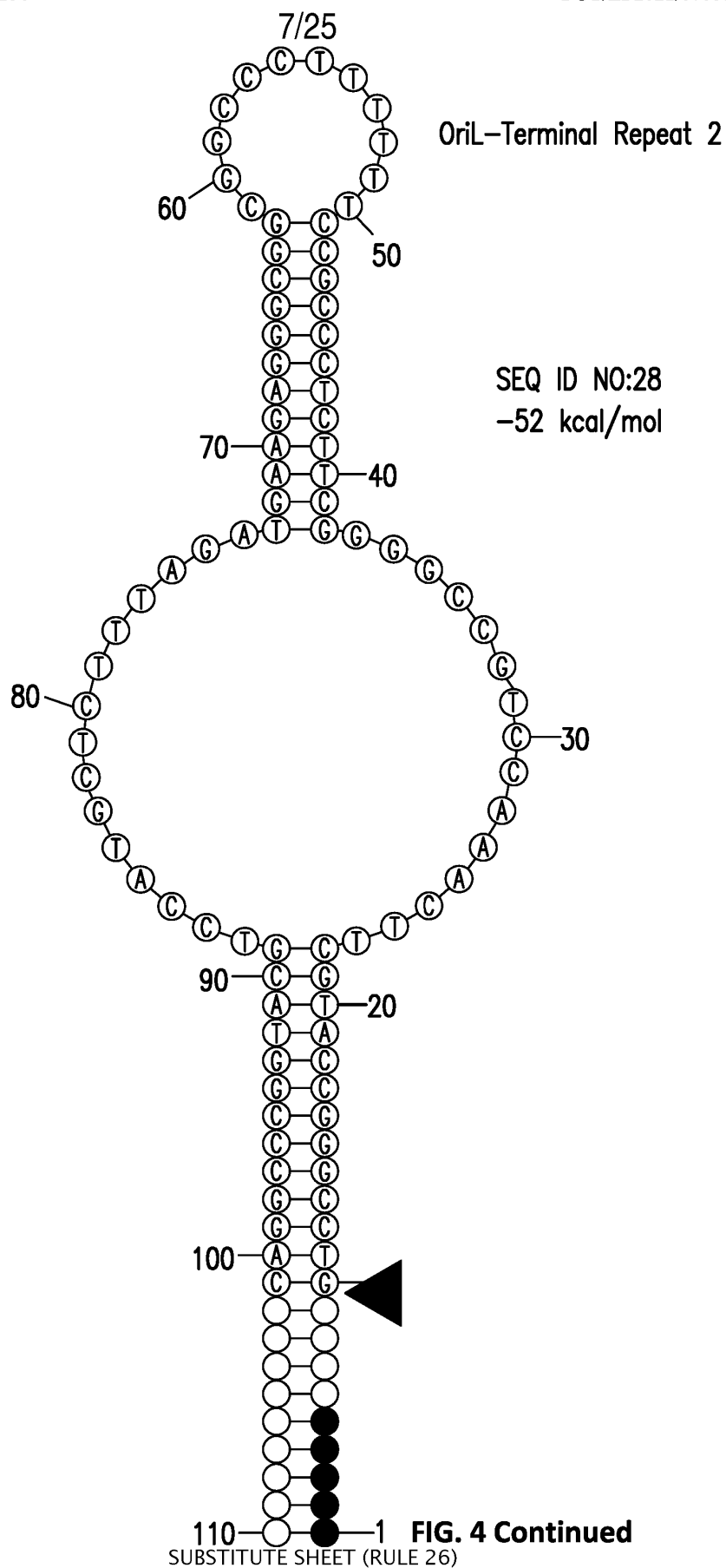


FIG. 3

SUBSTITUTE SHEET (RULE 26)





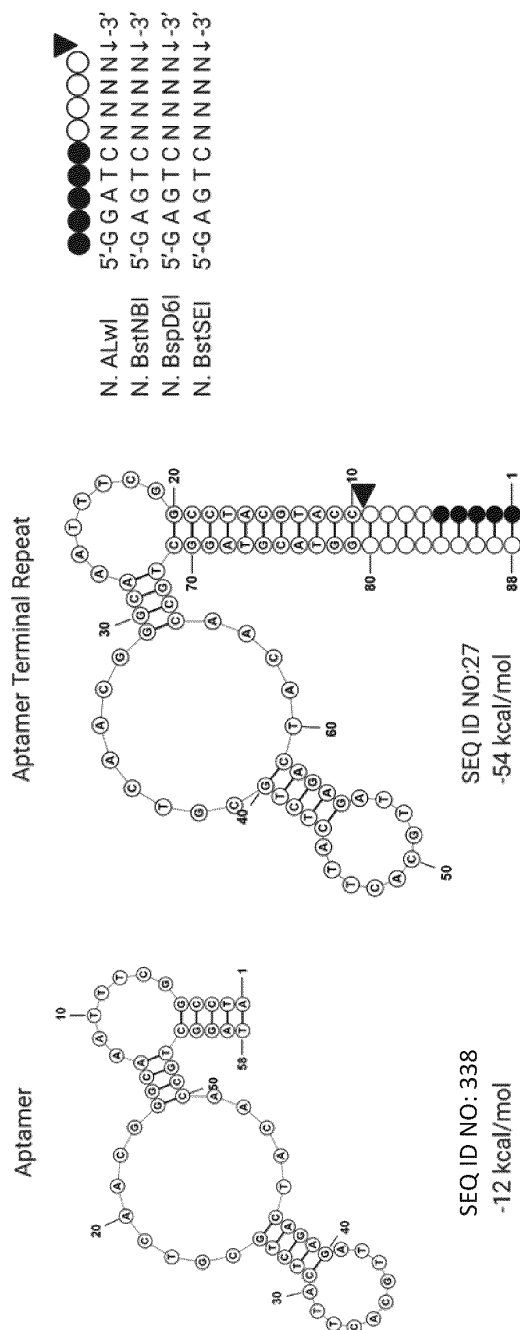


FIG. 5

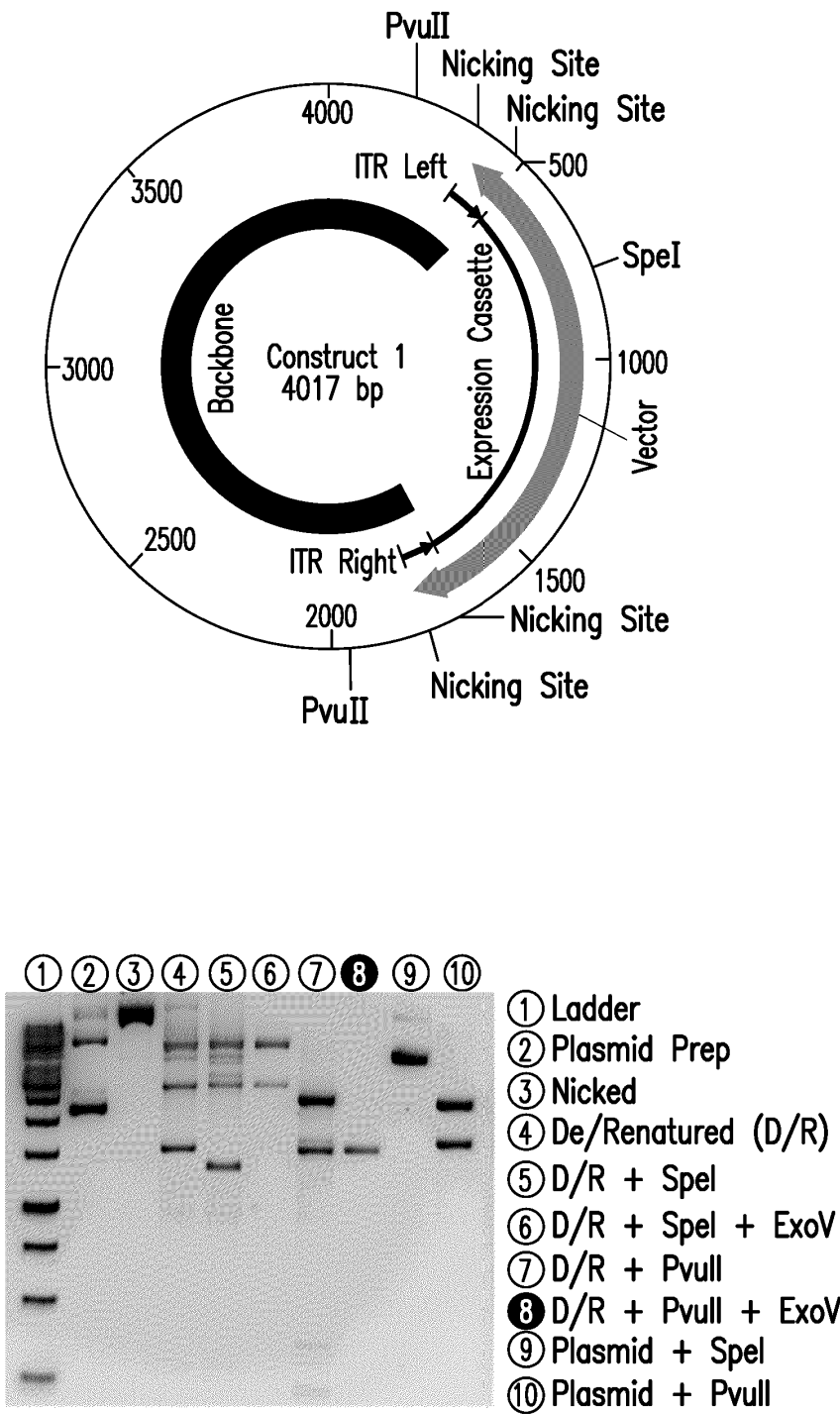


FIG. 6

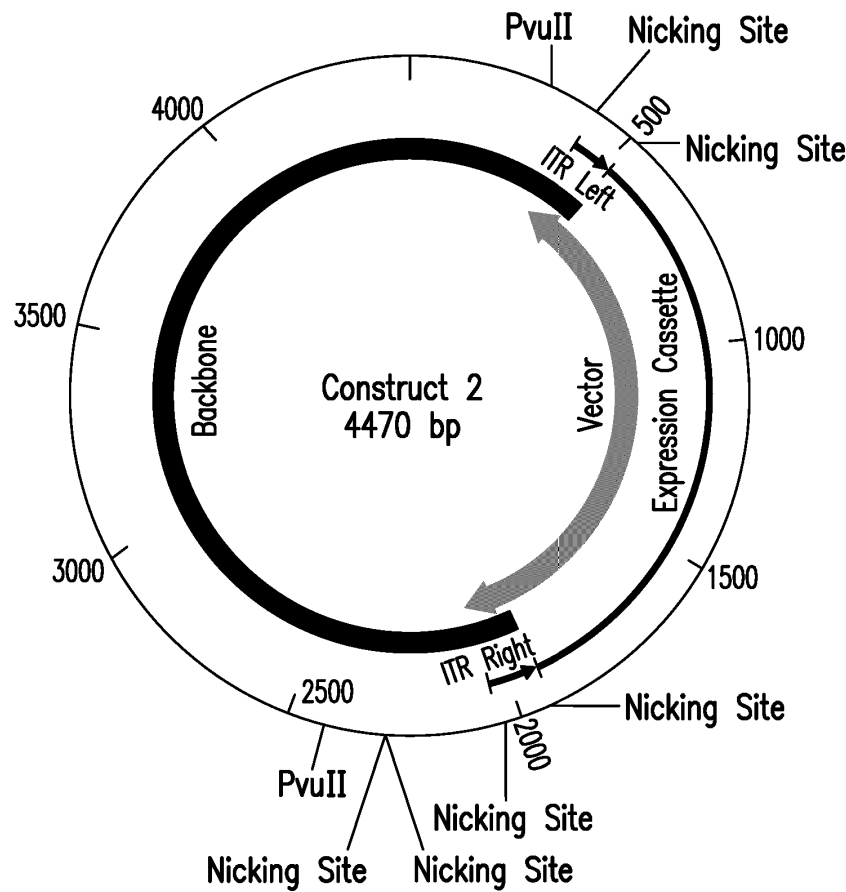
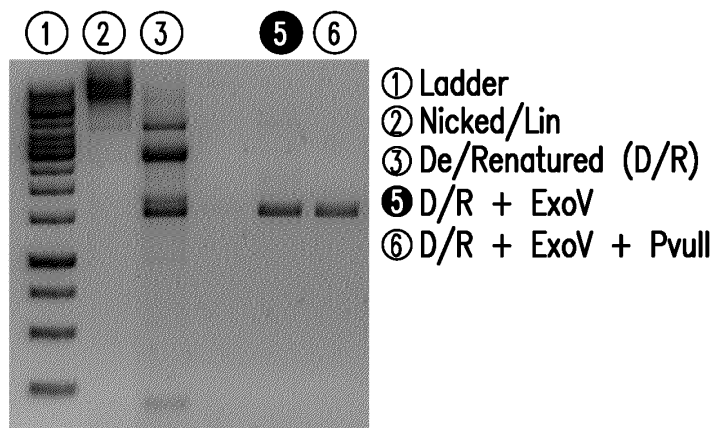


FIG. 7

SUBSTITUTE SHEET (RULE 26)

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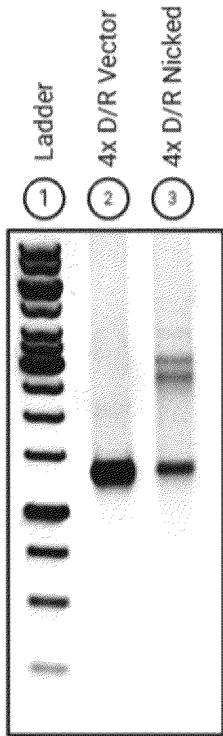


FIG. 8A

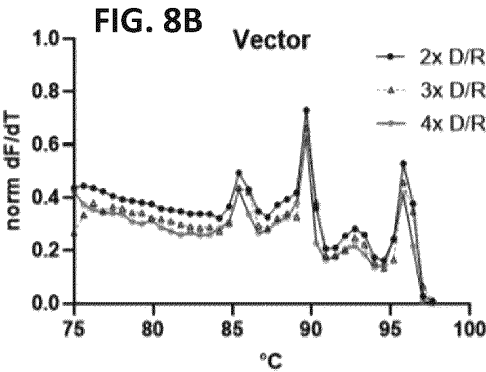
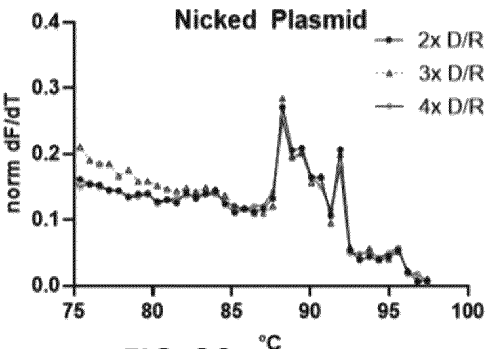


FIG. 8C

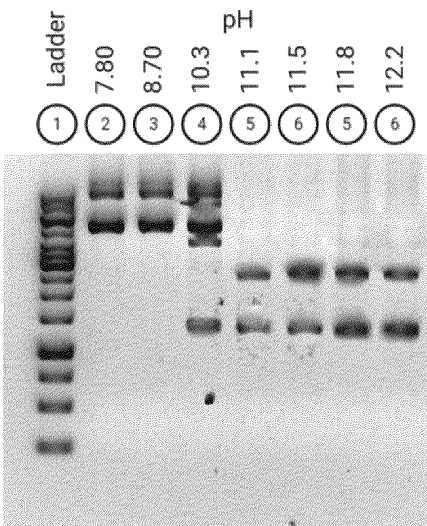


FIG. 9A

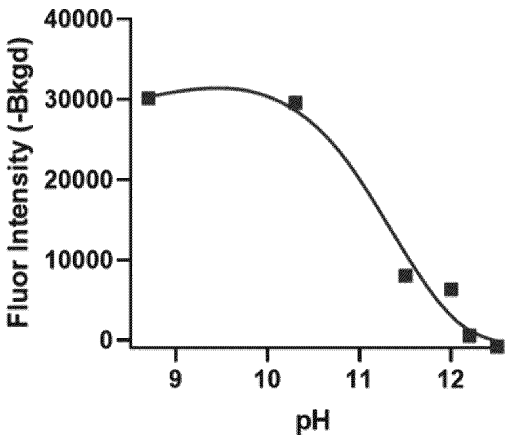


FIG. 9B

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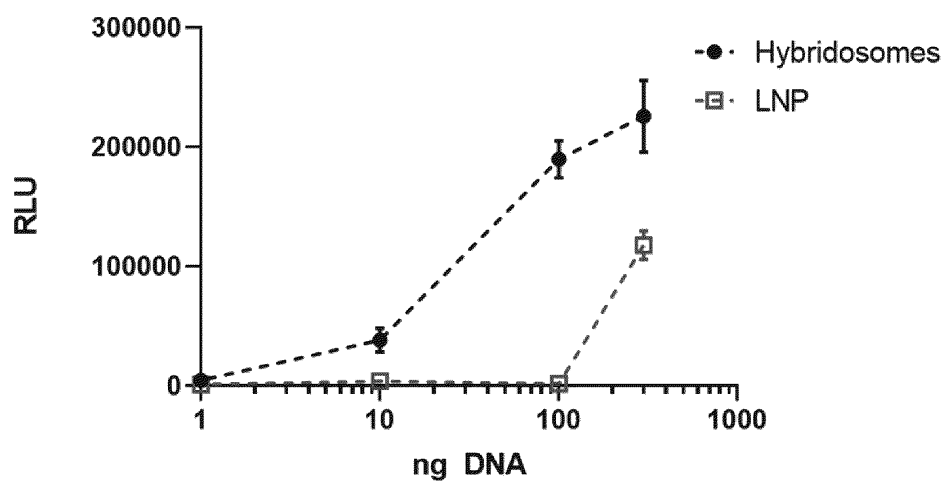


FIG. 10

Dividing Cells (293T)

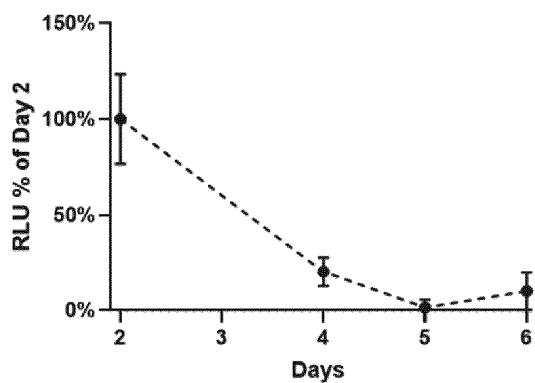


FIG. 11A

Non-Dividing Cells (Ir. MEFs)

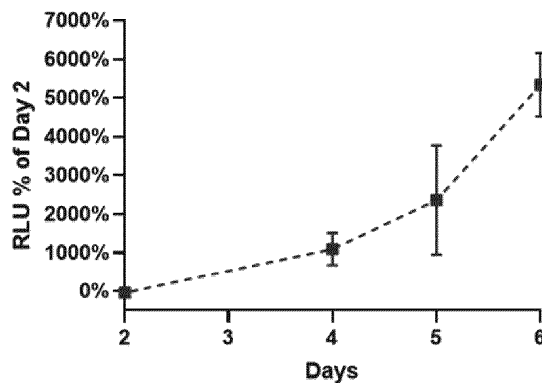


FIG. 11B

Non-Dividing Cells (IrMEFs) (0.1pmol/well)

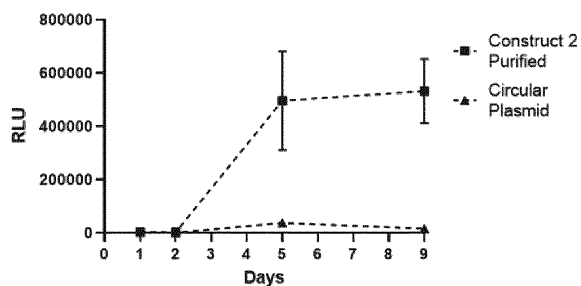


FIG. 11C

Dividing Cells (293T) (0.1pmol/well)

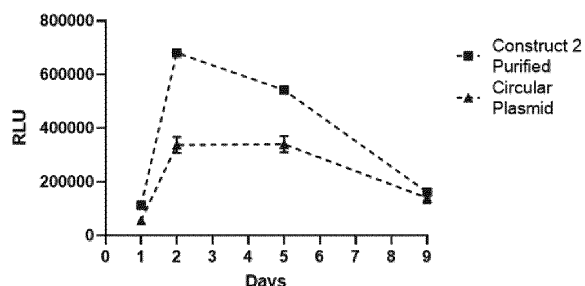


FIG. 11D

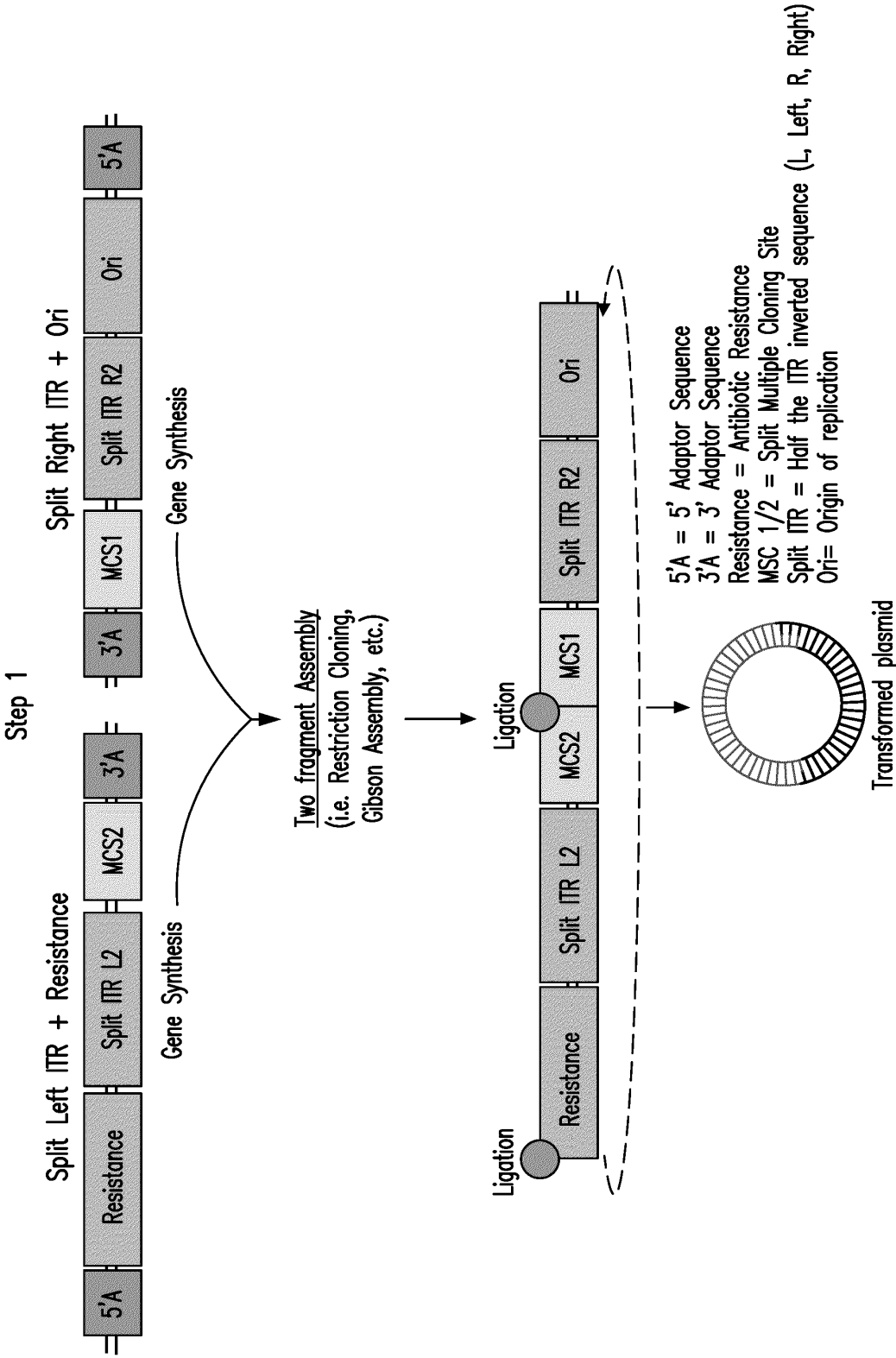


FIG. 12

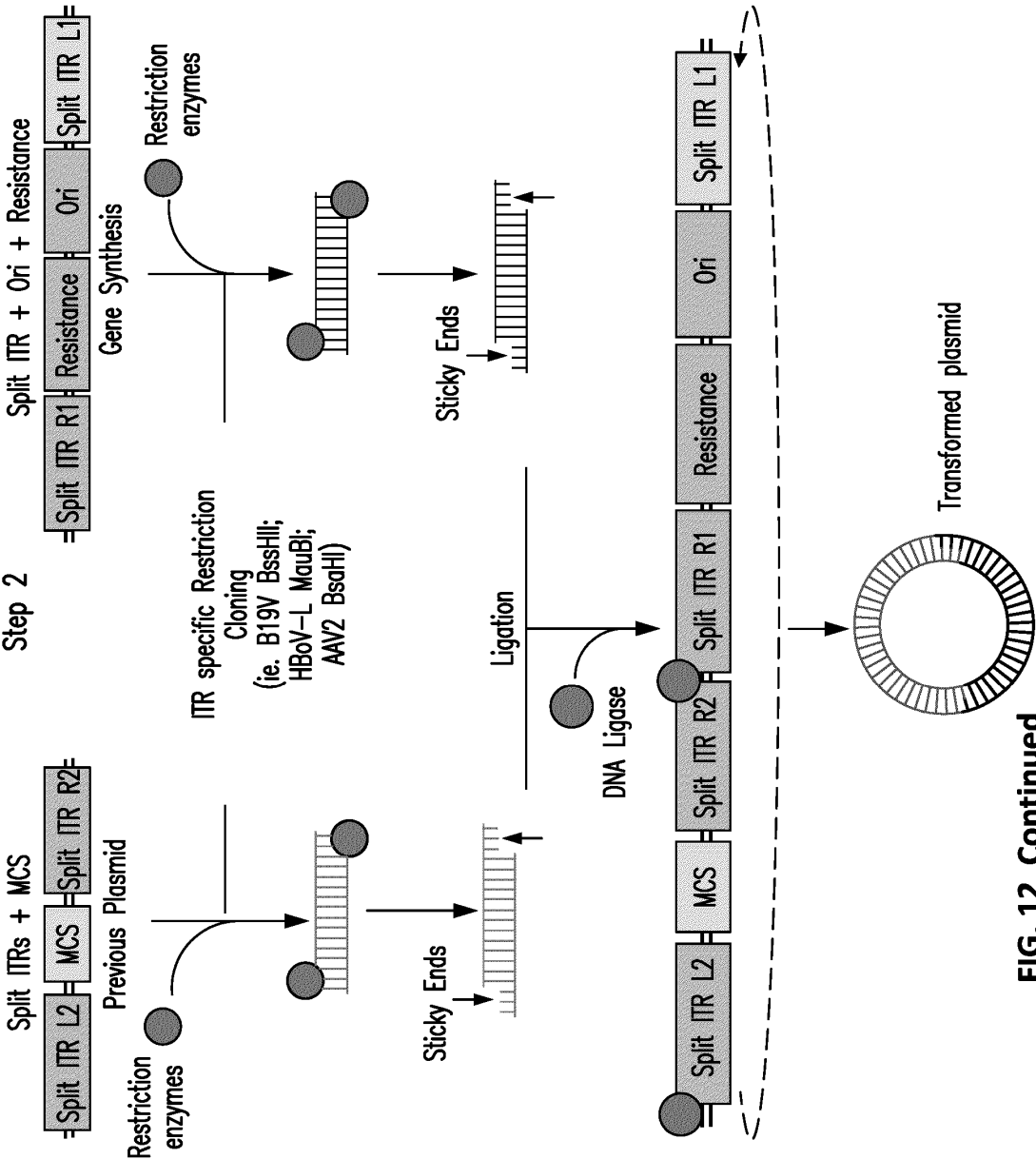


FIG. 12 Continued

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AV1 wt	TTGCCCACTCCCTCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BbvCI Top	AACGGGTGAGGGAGAGACGGCG-GAGCGAGCGAGCGACCGGCTGGTTCACAGCGCTCTGCAGACGAGCGCGGGGTGGCTCGCTCGCTC-GCGGCTCTCTCCCTCACCCGTT	125
Nb.BbvCI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BsmI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BsmI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BsrDI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BsrDI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.EssSI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.EssSI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BsiI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nb.BsiI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.AwiI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.AwiI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BbvCI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BbvCI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BemI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BemI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BspQI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BspQI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125
Nt.BsiNI Top	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	127
Nt.BsiNI Bottom	TTGCCCACTCCCGCTCTCGCGG-CTCGCTCGCTCGGTGGGGCTTCGGACCAAAAGTTCGGCAGACGGCAGAGCTCTGCTCTGCGGGCCCCACCGACGGAGTGGGCAA	125

FIG. 13

FIG. 14

FIG. 15

FIG. 16

FIG. 17

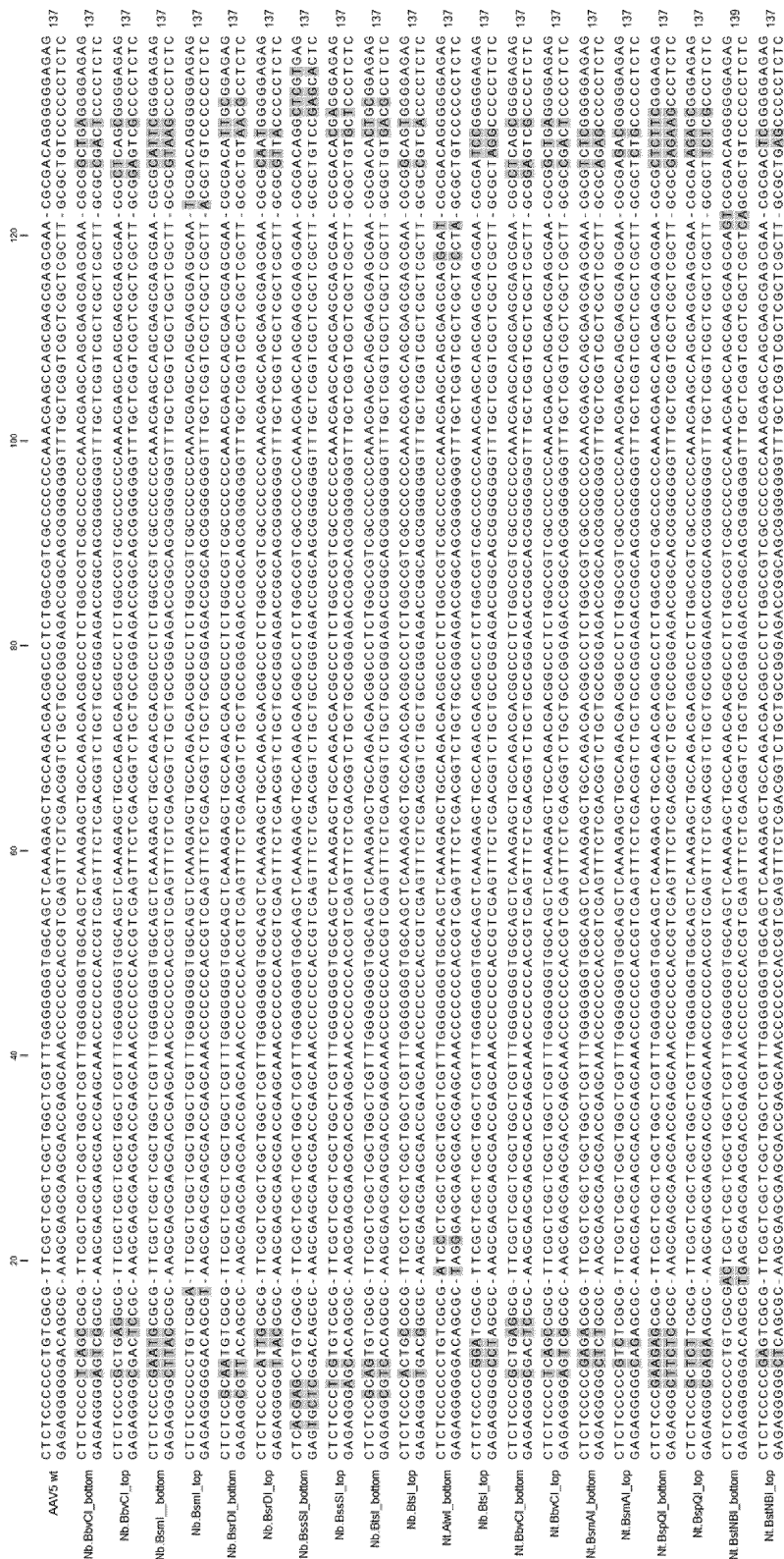


FIG. 18



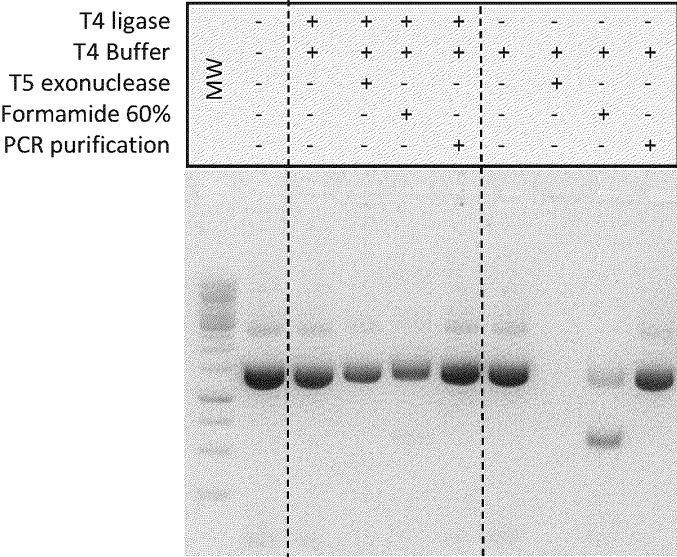


FIG. 20A

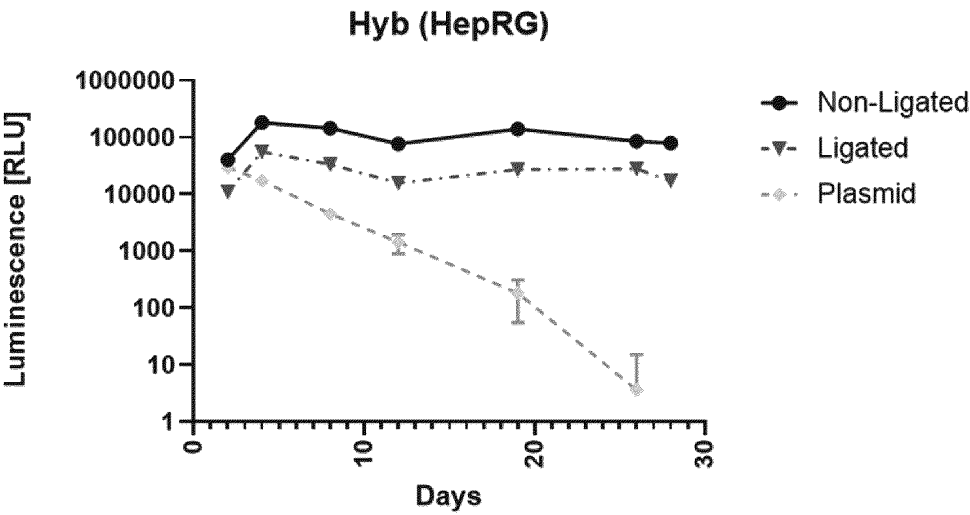


FIG. 20B

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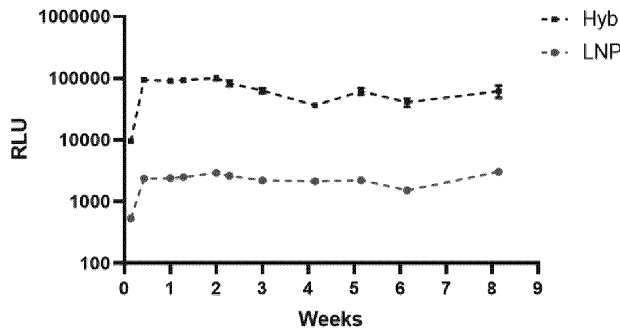


FIG. 21

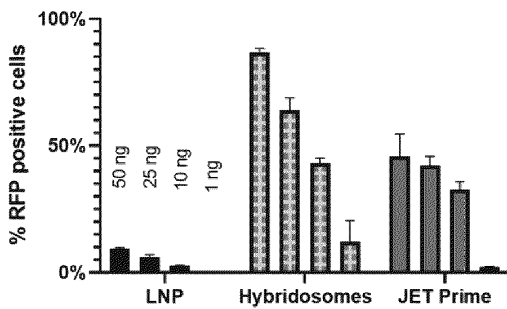


FIG. 22

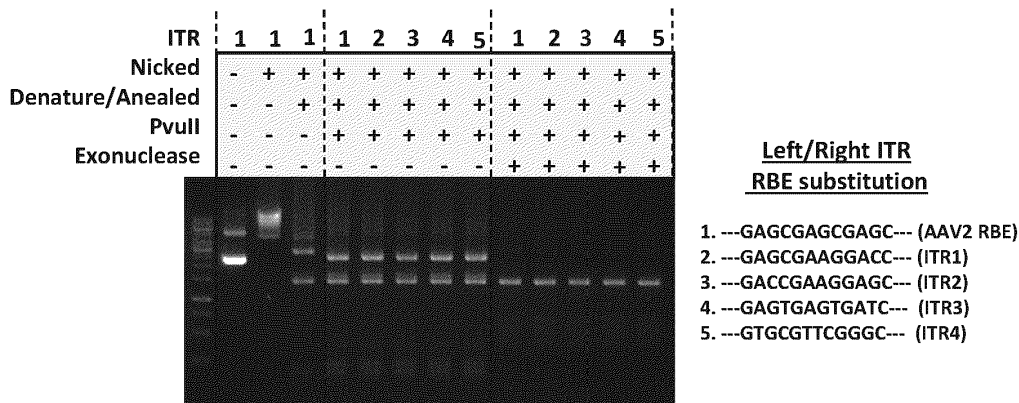


FIG. 23A

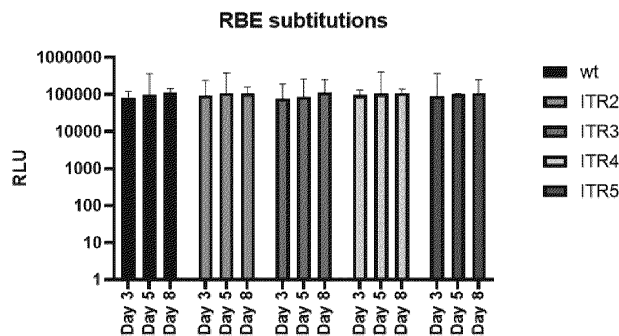


FIG. 23B

SUBSTITUTE SHEET (RULE 26)

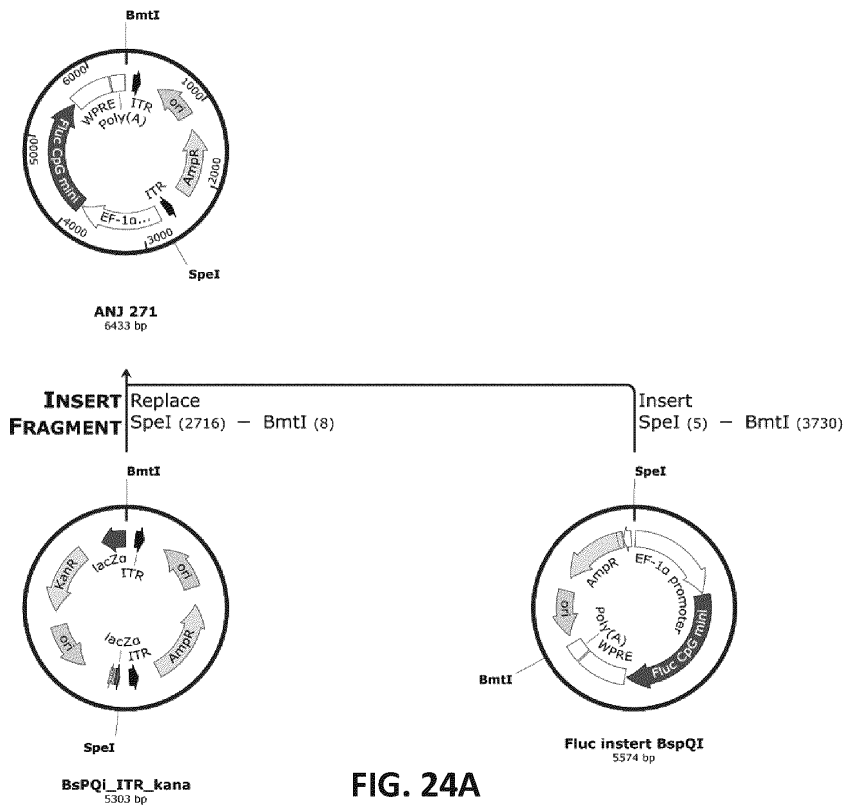


FIG. 24A

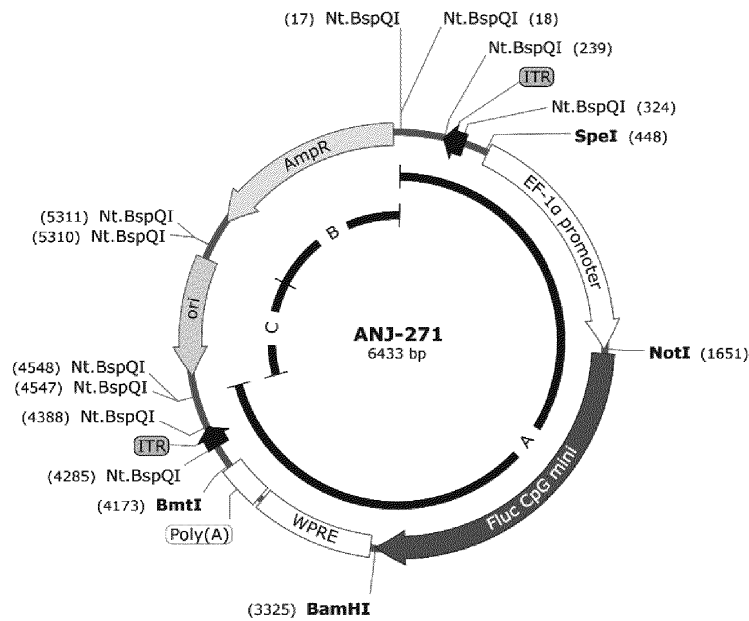


FIG. 24B

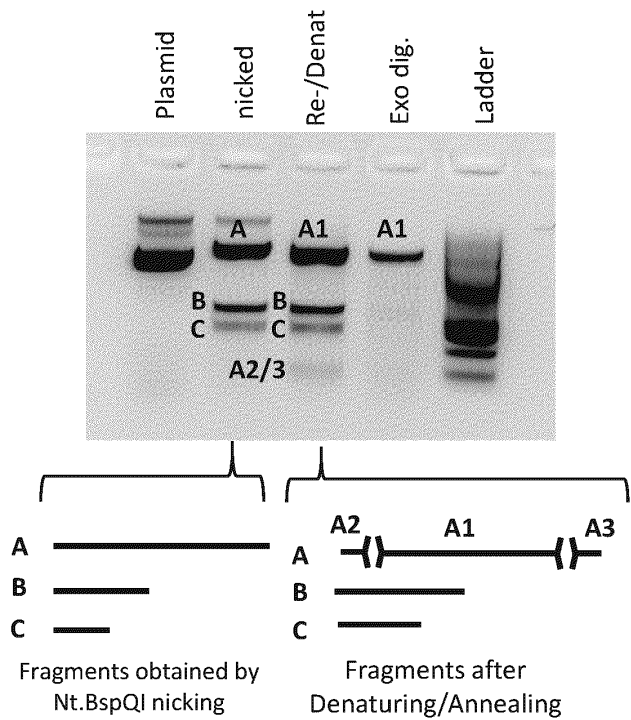


FIG. 25A

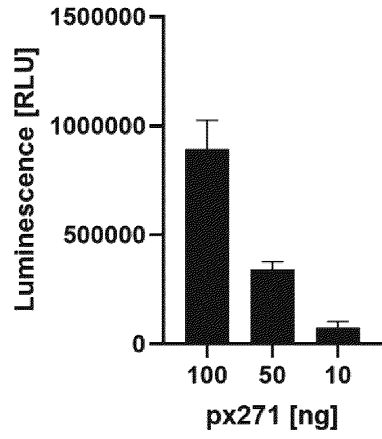


FIG. 25B

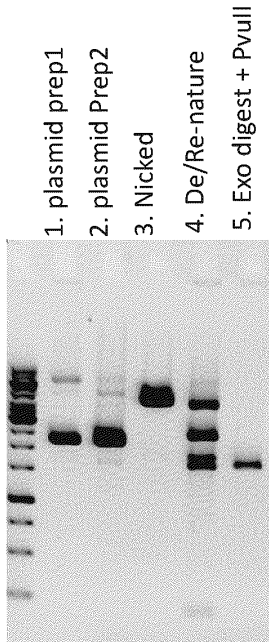


FIG. 26A

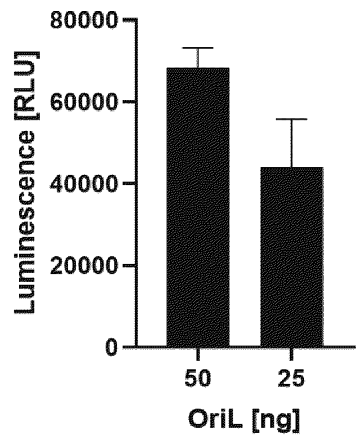


FIG. 26B