

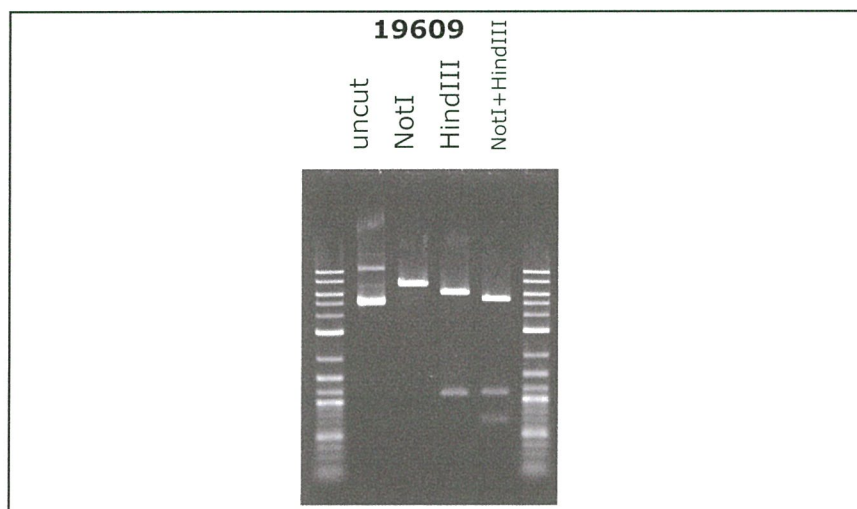
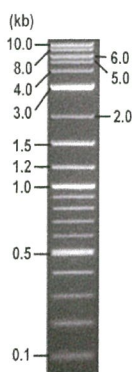


RIKEN DNA BANK

clone name : tdStayGold/pcDNA3

- Clone ID : RDB_19609
- Lot : 19609_B2Ea
- DNA Concentration : 25 nanogram/microliter
- Volume : 40 microliter
- Form : DNA solution in TE buffer
- Host : DH5 alpha
- Culture : LB medium
- Antibiotics : 100 microgram/ml Ampicillin
- Purification : QIAGEN QIAprep Spin Miniprep kit
- Digestion by restriction enzyme

1 kb Plus DNA Ladder
(NEB#N3200L),
250 ng/well



Electrophoresis : 100 nanogram DNA per lane ; 1% agarose gel , 1 x TAE Buffer

Restriction enzyme	Expected size of fragment
NotI	7.2 kbp
HindIII	6.1, 1.1 kbp
NotI+HindIII	5.4, 1.1, 0.7 kbp

● **Confirmation of the insertion sequence**

Please be sure to check our sequence analysis results before your request.

Sequence name	Primer ID	Primer name	Confirmed feature
Sequence - A	Pr0016	CMV_Forward	CMV pro,insert 5'((GGGGS)x3,tdStayGold 5')
Sequence - B	Pr0606	BGH_rev2	insert 3'(tdStayGold 3')
Sequence - C	Pr1213	hStayGold_F1	insert mid
Sequence - D	Pr1214	EV_linker_F1	insert mid
Sequence - E	Pr0731	SV40pro_ori_F	SV40 pro_ori,NeoR
Sequence - F	-	-	-
Sequence - G	-	-	-
Sequence - H	-	-	-

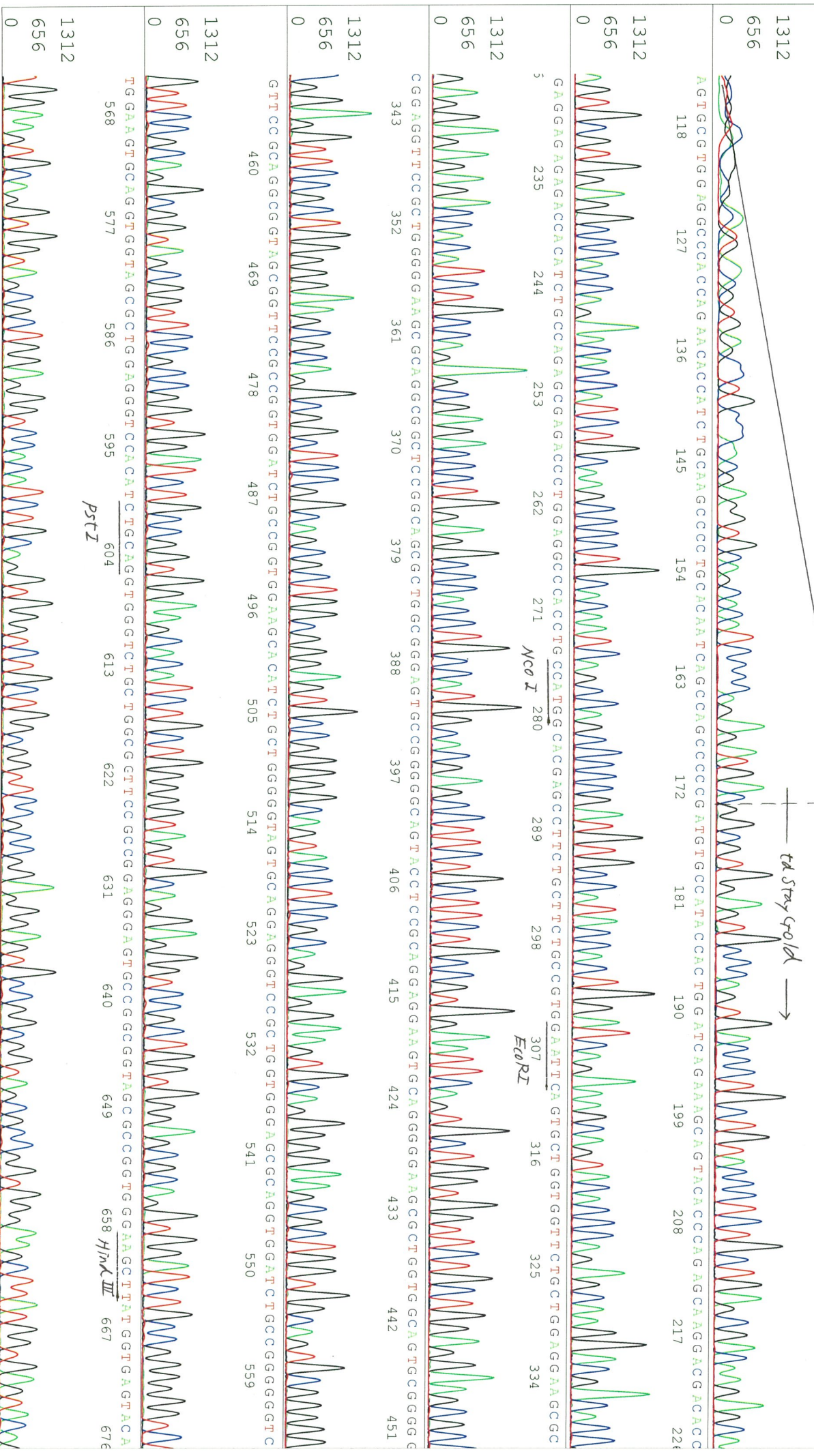
APPROVED BY :



S/N G:259 A:164 T:102 C:150 primer name C : hStayGold_F1 D08836A5_B2Ea_3_hStayGold_F1

KB.bcp KB 1351 to 13696 PK1 Loc:1328 Pts 1351 to 13696 PK1 Loc:1328

KB 1.4.1.8 Cap:10 5'-CCAGGACTTCAAGGAGAACAGC-3' Version 6.0 HiSQV Bases: 979



S/N G:200 A:184 T:146 C:144

primer name *D* : EV_linker_F1

D08836A5_B2Ea_3_EV_linker_F1

May 23, 2022 03:16PM, JST

KB.bcp

5'-TGCCGGTGGAAAGCACATCTGC-3'

KB_3500_POP7_BDTV3.mob

May 23, 2022 03:43PM, JST

KB 1.4.1.8

Cap:13

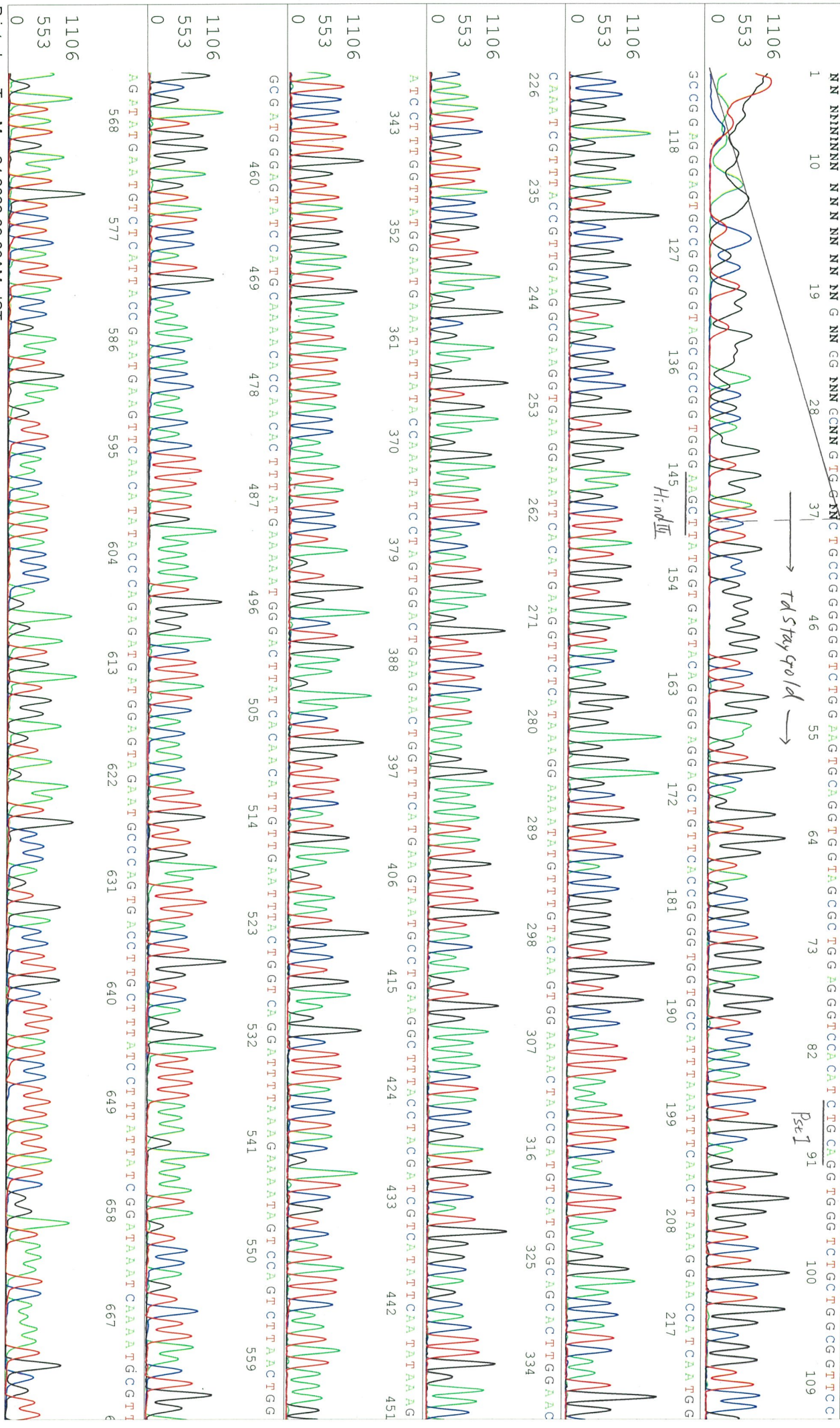
Pts 1373 to 13744 Pk1 Loc:1350

Spacing:1.1.93 Pts/Panel1350

Version 6.0

HiSQV Bases: 955

Plate Name: 20220523 Kitaku



S/N G:200 A:184 T:146 C:144

D08836A5_B2Ea_3_EV_linker_F1

KB_3500_POP7_BDTV3.mob

Pts 1373 to 1374 PK1 Loc:1350

Version 6.0 HiSQV Bases: 955

May 23, 2022 03:16PM, JST
May 23, 2022 03:43PM, JST
Spacing: 11.93 Pts/Panel1350
Plate Name: 20220523 Kitaku

KB 1.4.1.8 Cap:13

GAGGCTCACCAAAATACAAATCTGCCAAGCCCTCTTCATAATCAACCAGCCACTGATGTCCCATATCACTGGATTTCGTTAAACAATACAACAAGAAGCANAGANAGNACCGANGA
176 685 694 703 712 721 730 739 748 757 766 775 784

