

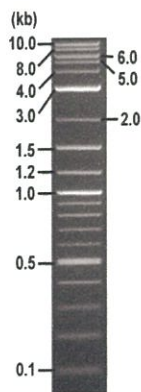


RIKEN DNA BANK

clone name : pGL4-phMMP9

- Clone ID : RDB _ 07537
- Lot : 14426 _ A6Kg
- 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

2-Log DNA Ladder
(NEB#N3200L),
125 ng/well



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

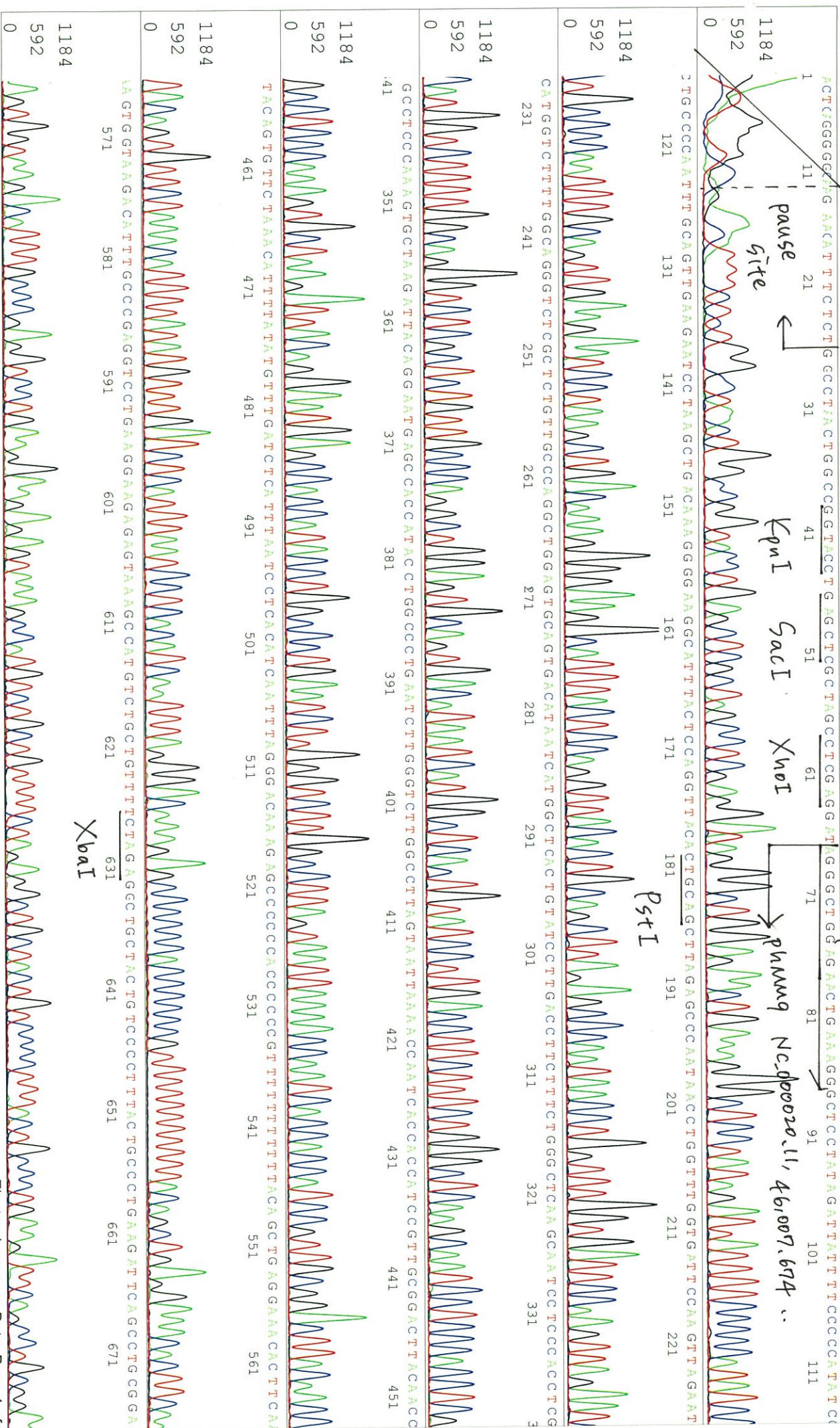
Restriction enzyme	Expected size of fragment
<u>BglII</u>	<u>5.6</u> kbp
<u>XhoI</u>	<u>5.6</u> kbp
<u>BglII + XhoI</u>	<u>4.2, 1.4</u> kbp
	kbp
	kbp

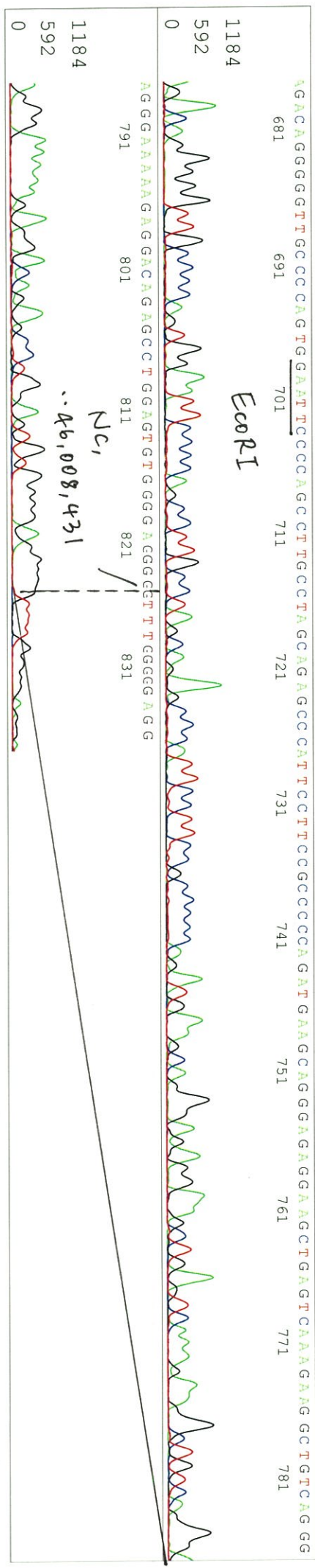
● Confirmation of the insertion sequence

Sequence name	Primer name	Sequence name	Primer name
Sequence - A	pGL4-4174F	Sequence - E	-
Sequence - B	pGL4-136R	Sequence - F	-
Sequence - C	pAxCALNL_F1	Sequence - G	-
Sequence - D	-	Sequence - H	-

APPROVED BY :







S/N G:1302 A:852 T:716 C:1073

07537_A6Kg_4_pGL4-136R

Dec 02, 2016 10:02AM, JST

KB.bcp

primer name β : pGL4-136R

KB_3500_POP7_BDTV3.mob

Dec 02, 2016 10:28AM, JST

KB 1.4.1.8 Cap:14

5-CTTCGAGTGGTAGAATGGC-3'

Pts 1552 to 13548 PK1 Loc:1529

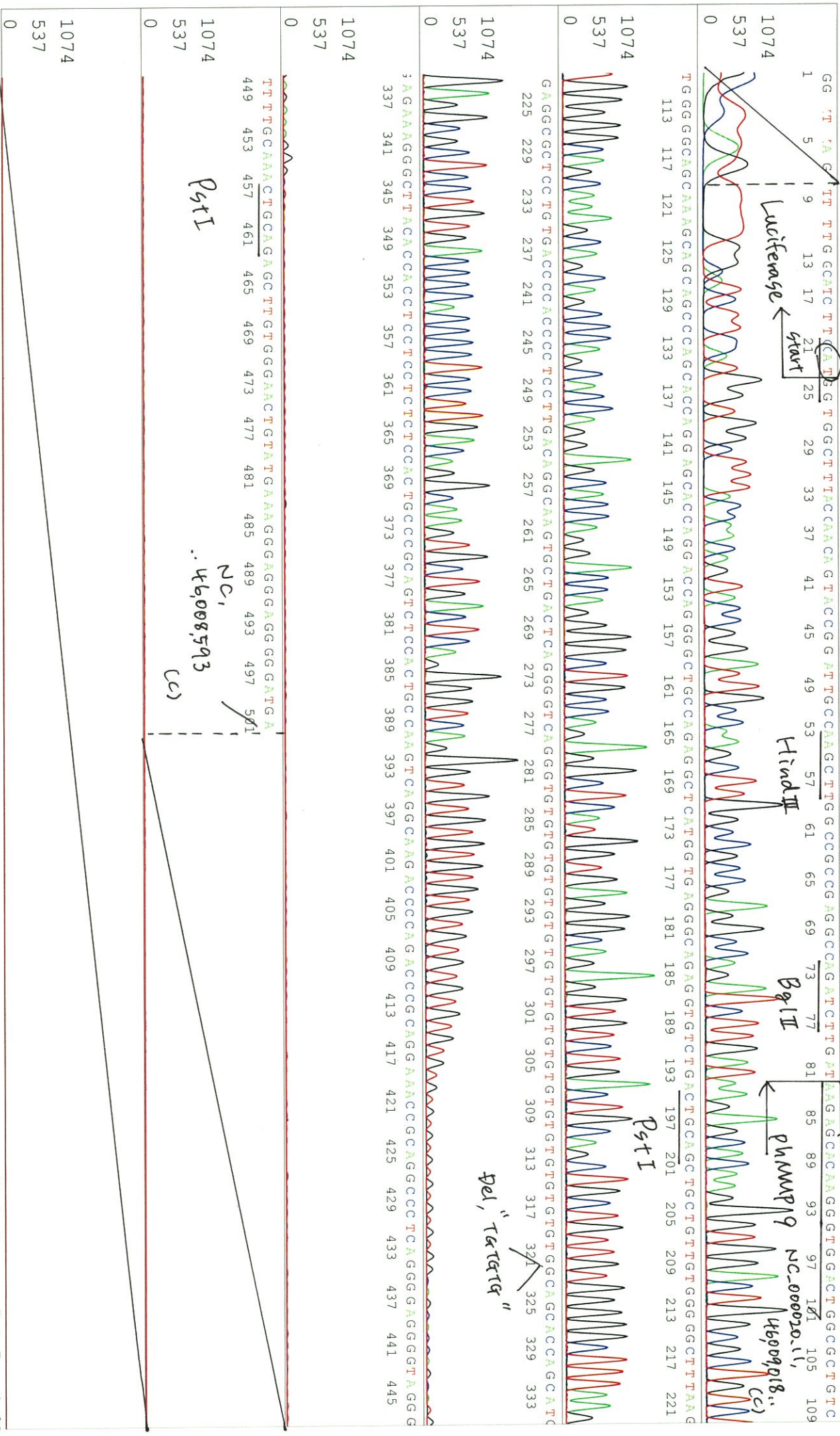
Spacing:11.96 Pts/Panel1350

Version 6.0 HiSeqV Bases: 481

Version 6.0 HiSeqV Bases: 481

R primer

Plate Name: 20161202_mix



S/N G:81 A:93 T:81 C:133

07537_A6Kg_4_paxCALNL_F1

Dec 02, 2016 10:02AM, JST

KB.bcp

primer name C : paxCALNL_F1

KB 3500_POP7_BDTV3.mob

Dec 02, 2016 10:28AM, JST

KB 1.4.1.8 Cap:17

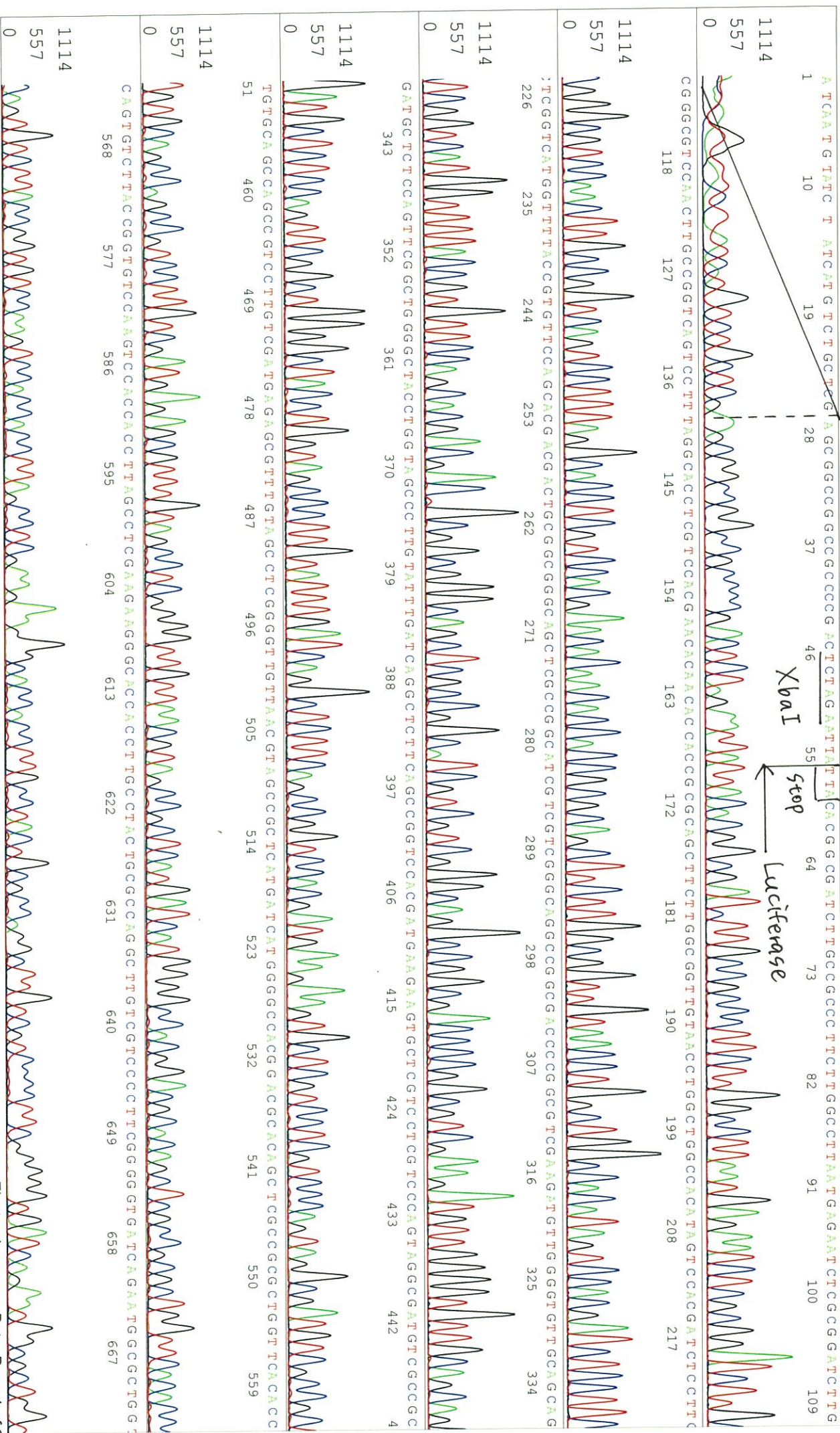
5'-CACTGCATTCTAGTTGTGTTGTCC-3'

Pis 1568 to 13645 PK1 Loc:1545

Spacing:12.22 Pts/Panel1350

Version 6.0 HISQV Bases: 1000

Plate Name: 20161202_mlx



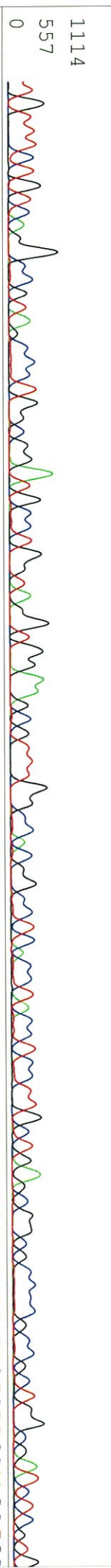
SN G:81 A:93 T:81 C:133

KB.bcp

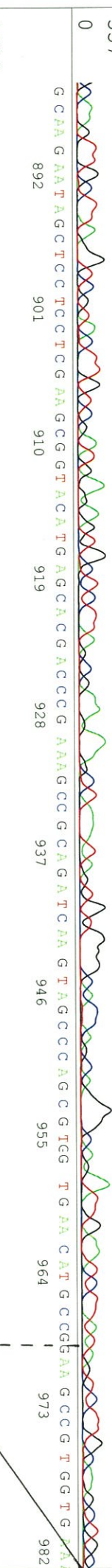
KB 1.4.1.8 Cap:17

Version 6.0 HISQV Bases: 1000

TTGTTTCTGTCAGGCCGTAGCCCTGGCCGATGCCCTGGTAGGTGGAAAGCGTTTGGCCACCGGCTTACCTCCCTTGGCTGAGCGGGCCCGCCCGCTGGCGATCTCGTTG



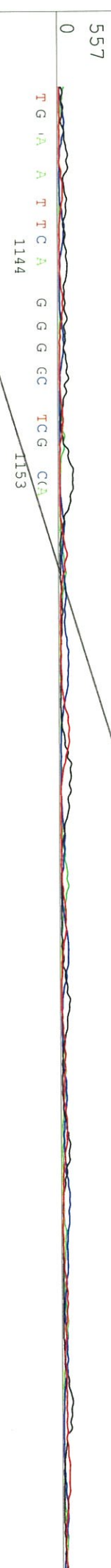
CAAGTTGCTTAGGTCCTACCTTGTCCGATGAGAGTGTCTTAGCGAAGAAAGCTAAATAGTGTGGGCACCAAGCA GGGCAGATTGAAATCTTATAGTCTTGCCAA GCTGC



GCAA GAATAGCTCC TCC TCCG AAGCGGTACATGAGCA CGAACCCG AAAAGCCG CAGATCAA GTAGCCCAAGCG TGG TGAATCATG CCGGAA GCCG TGGTGA



ATGGCA CCA CCGCTG AEGGATAGCGGTGTCGGGGATGATCTGGGTTGCCG AAGATTGGG GTC GCGGGCAATGACTGAA TCGGGACAC A



TGAAATTCAGGGGC TCGCGA