

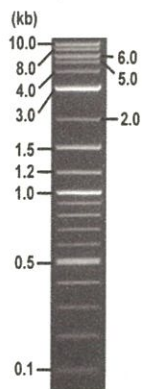


RIKEN DNA BANK

clone name : pAxCAhIFN-beta (forward)

- Clone ID : RDB \_ 02285
- Lot : 14299 \_ A6Ht
- DNA Concentration : 25 nanogram/microliter
- Volume : 40 microliter
- Form : DNA solution in TE buffer
- Host : DH5 alpha / Gigapack III
- 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 : 200 nanogram DNA per lane ; 1% agarose gel , 1 x TAE Buffer

Restriction enzyme	Expected size of fragment
<u>ClaI</u>	<u>44.8, 0.7</u> kbp
<u>HindIII</u>	<u>17.5, 8.0, 5.3, 4.6, 3.8, 2.9, 2.1, 1.1, 0.1</u> kbp

● Confirmation of the insertion sequence

Sequence name	Primer name	Sequence name	Primer name
Sequence - A	pAxCA_F1	Sequence - E	-
Sequence - B	Amp_R	Sequence - F	-
Sequence - C	-	Sequence - G	-
Sequence - D	-	Sequence - H	-

APPROVED BY :



S/N G:124 A:81 T:96 C:107

primer name A : pAxCA\_F1

02285\_14299\_A6Ht\_pAxCA\_F1

Sep 13,2016 10:02AM, JST

KB.bcp

5- GGCTTCTGGCGGTGACCGGC -3'

KB\_3500\_POP7\_BDTV3.mob

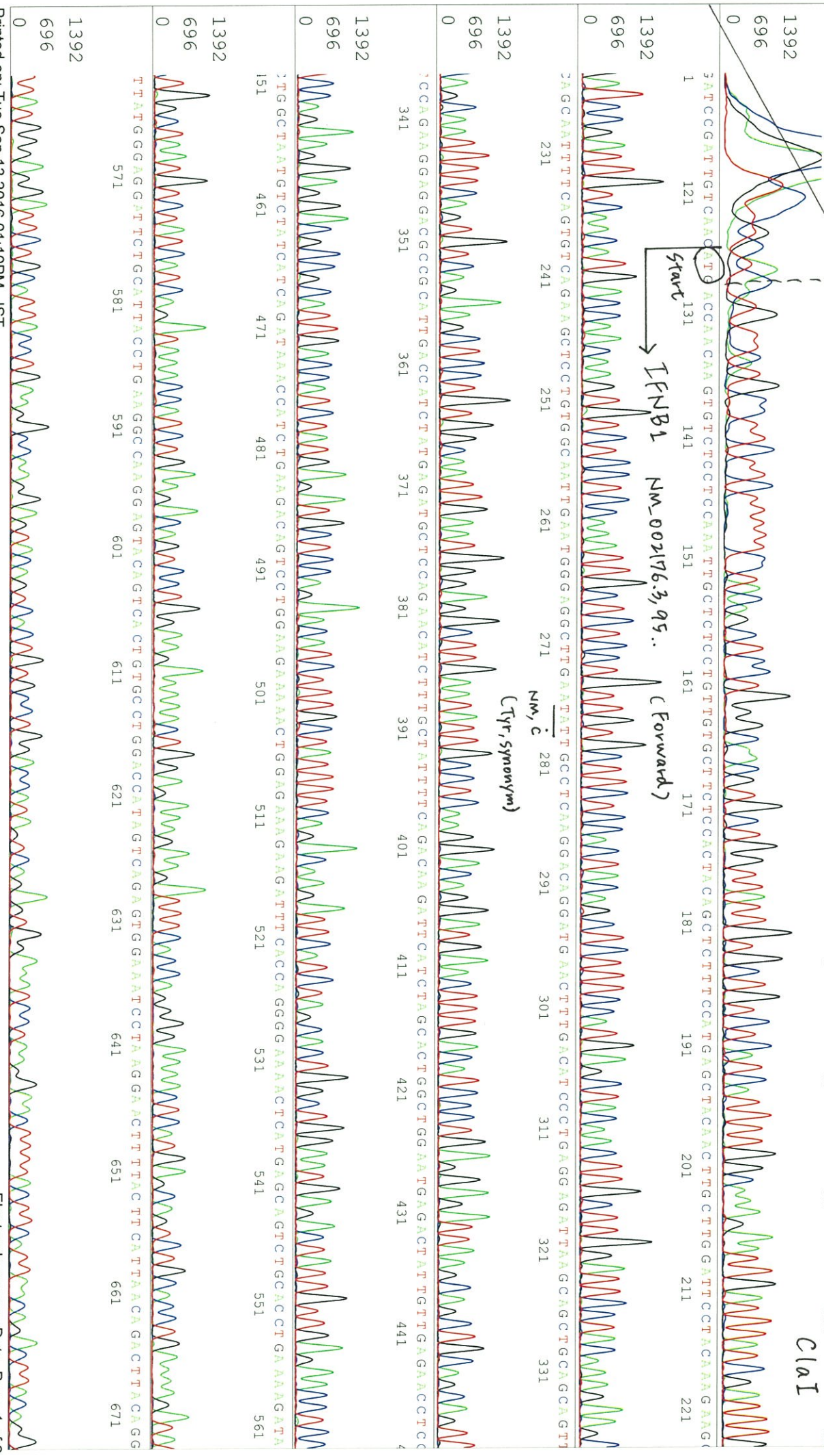
Sep 13,2016 10:28AM, JST

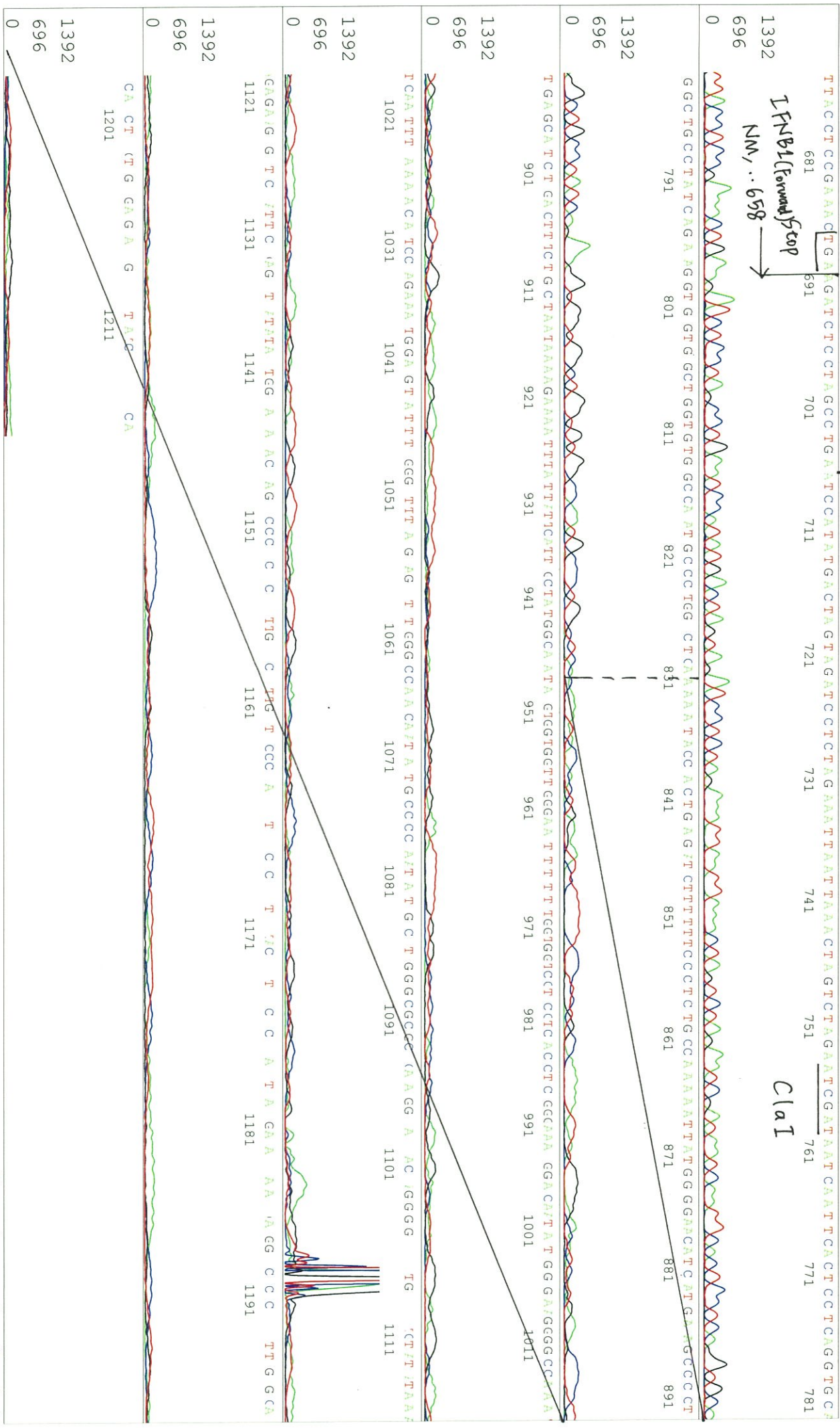
KB 1.4.1.8 Cap:16

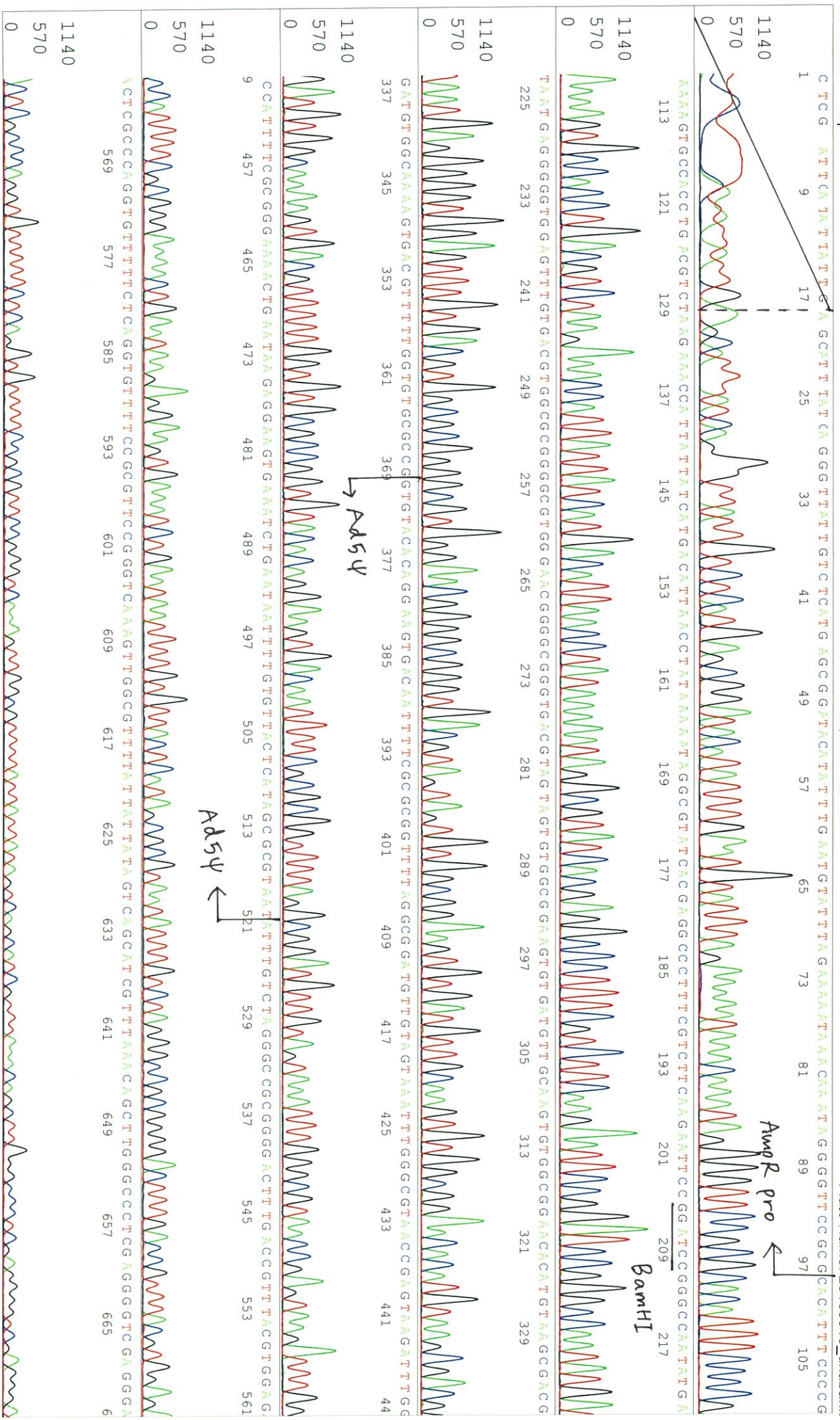
Version 6.0 HiSeqV Bases: 938

Pts 1438 to 13113 PK1 Loc:1415

Plate Name: 20160913\_dhbank







S/N G:304 A:183 T:160 C:114

02285\_14299\_A6Ht\_Amp\_R

Sep 13, 2016 10:02AM, JST

KB\_3500\_POP7\_BDTV3.mob

Sep 13, 2016 10:28AM, JST

Pts 1549 to 13176 PK1 Loc: 1526

Spacing: 11.75 Pts/Panel1350

KB 1.4.1.8 Cap:19

Version 6.0 HiSeqV Bases: 905

Plate Name: 20160913\_dnabank

TCTCCATTAAGAGAAAGAGGACAGCTTATGACTGGGAGTACAGGAGGAGGAAAACTCTGGCTAGTAAACATGTAAGGAAAAATTTTAGGGGATGTTAAAGAAAAAA

1140  
570

TAAACAACAACAATATAAAAAAACTTAACCTCAAGTCAAAGCTTTTCTATGGATTAAGCAACAGCAAGGGGCTGTTTCAATATACCTGATGACCTGTTA/T

1140  
570

AGCCAACTTTGTTCAATGGCAAGCCAGCATAATGGGCA/T/TGTTGCCAAACTCTAAACAAATACCTCACTGATGTTTAAATGAAATTTGGCCCTCAATTT

1140  
570

FGTGTCTCAGAGTGAAGACACAATAAACTCAACAACCTAATCTCAACACTTAATACTGAAATTTGGCCCTCAATTTAAAGCTGAGTCTGAG

1140  
570

CT/C GGCCTCTGTGATGTGCCCTAACTTAACTTCTATGAAAGATAAATCTTAAATAGCTGAGTCAAG

1140  
570

CT/C GGCCTCTGTGATGTGCCCTAACTTAACTTCTATGAAAGATAAATCTTAAATAGCTGAGTCAAG

1140  
570