

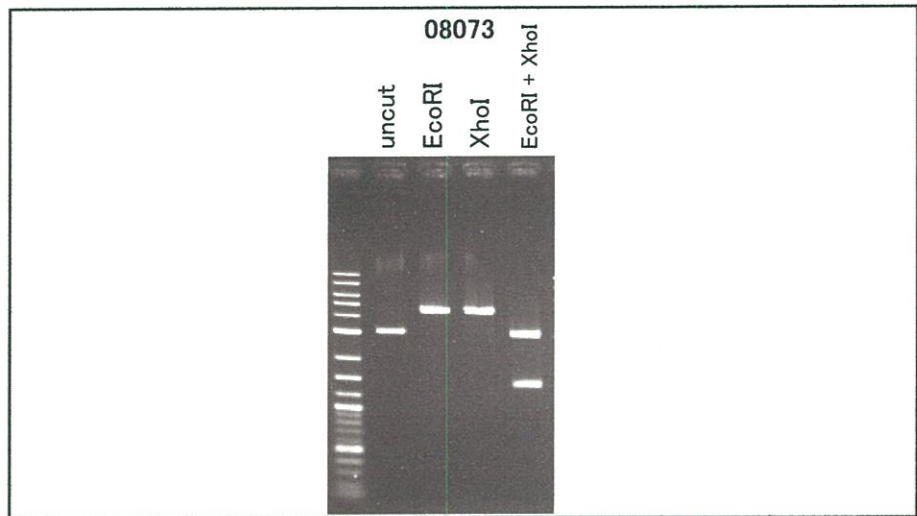
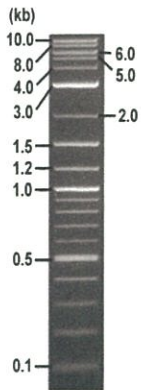


RIKEN DNA BANK

clone name : pmmFLIP (Mouse c-FLIP)

- Clone ID : RDB _ 08073
- Lot : 08073 _ A6Ik
- 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

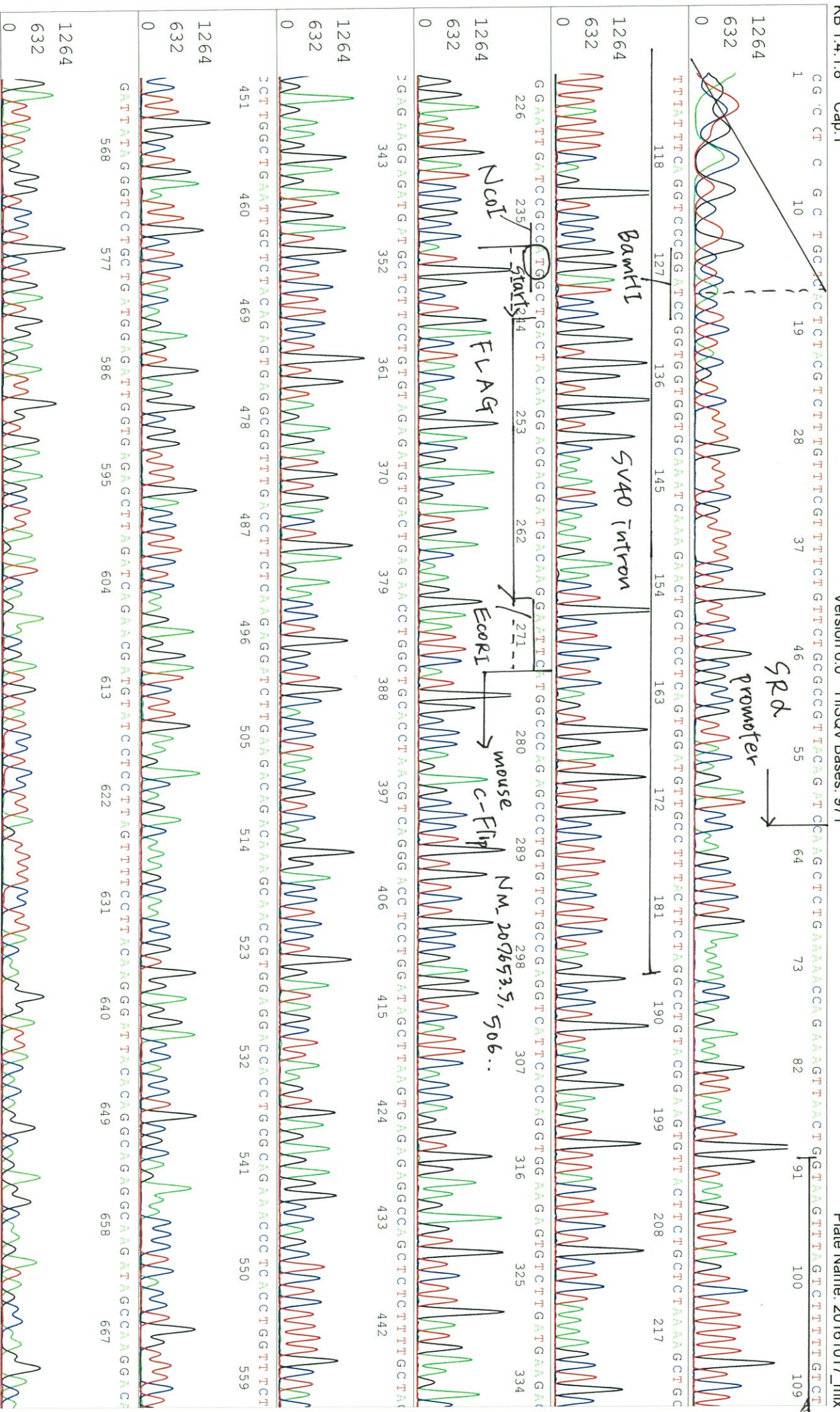
Restriction enzyme	Expected size of fragment
<u>EcoRI</u>	<u>4.5</u> kbp
<u>XhoI</u>	<u>4.5</u> kbp
<u>EcoRI + XhoI</u>	<u>3.1, 1.5</u> kbp
_____	_____ kbp
_____	_____ kbp

● Confirmation of the insertion sequence

Sequence name	Primer name	Sequence name	Primer name
Sequence - A	SRapro_F	Sequence - E	-
Sequence - B	ME-1290Rv	Sequence - F	-
Sequence - C	-	Sequence - G	-
Sequence - D	-	Sequence - H	-

APPROVED BY :

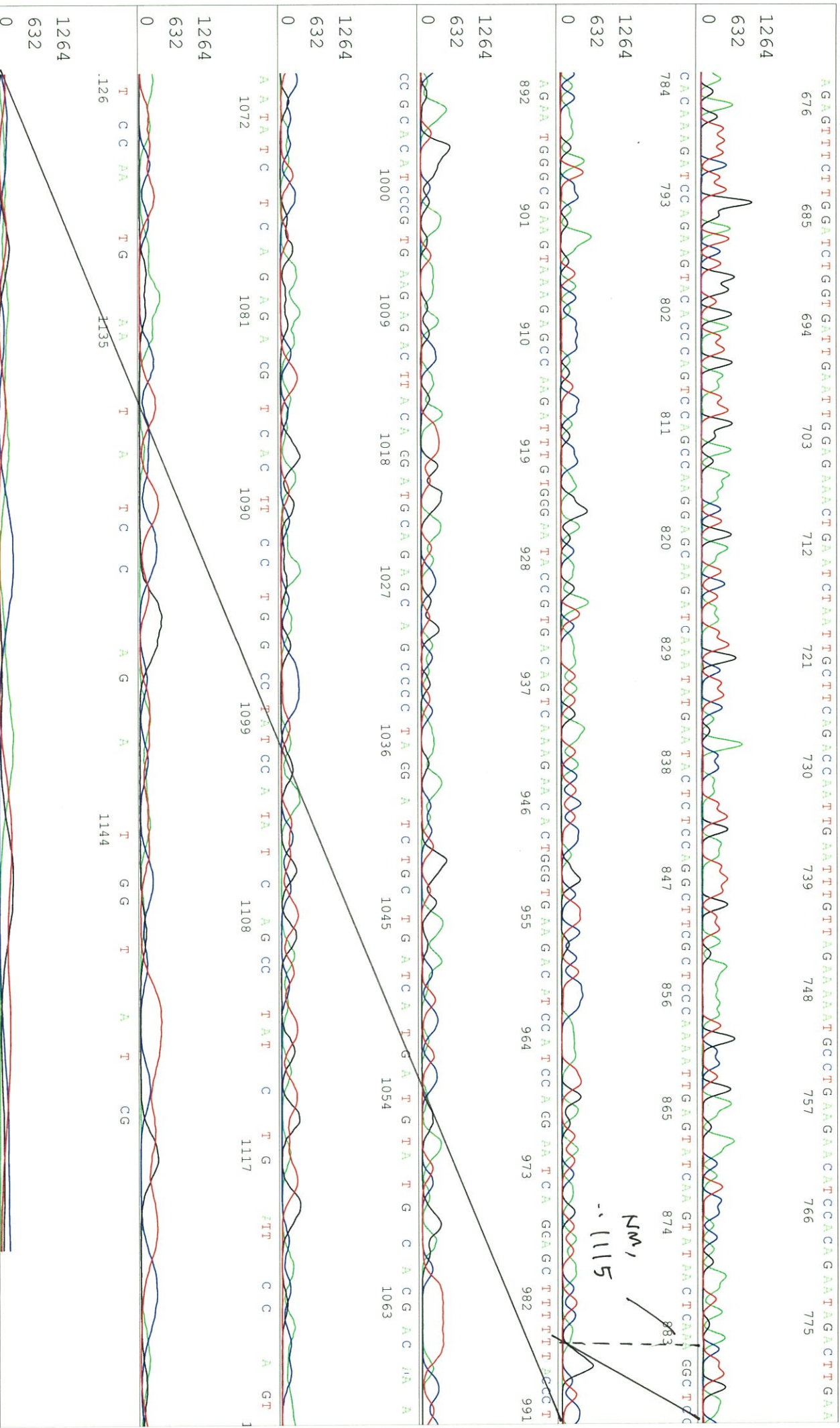




S/N G:495 A:468 T:524 C:423

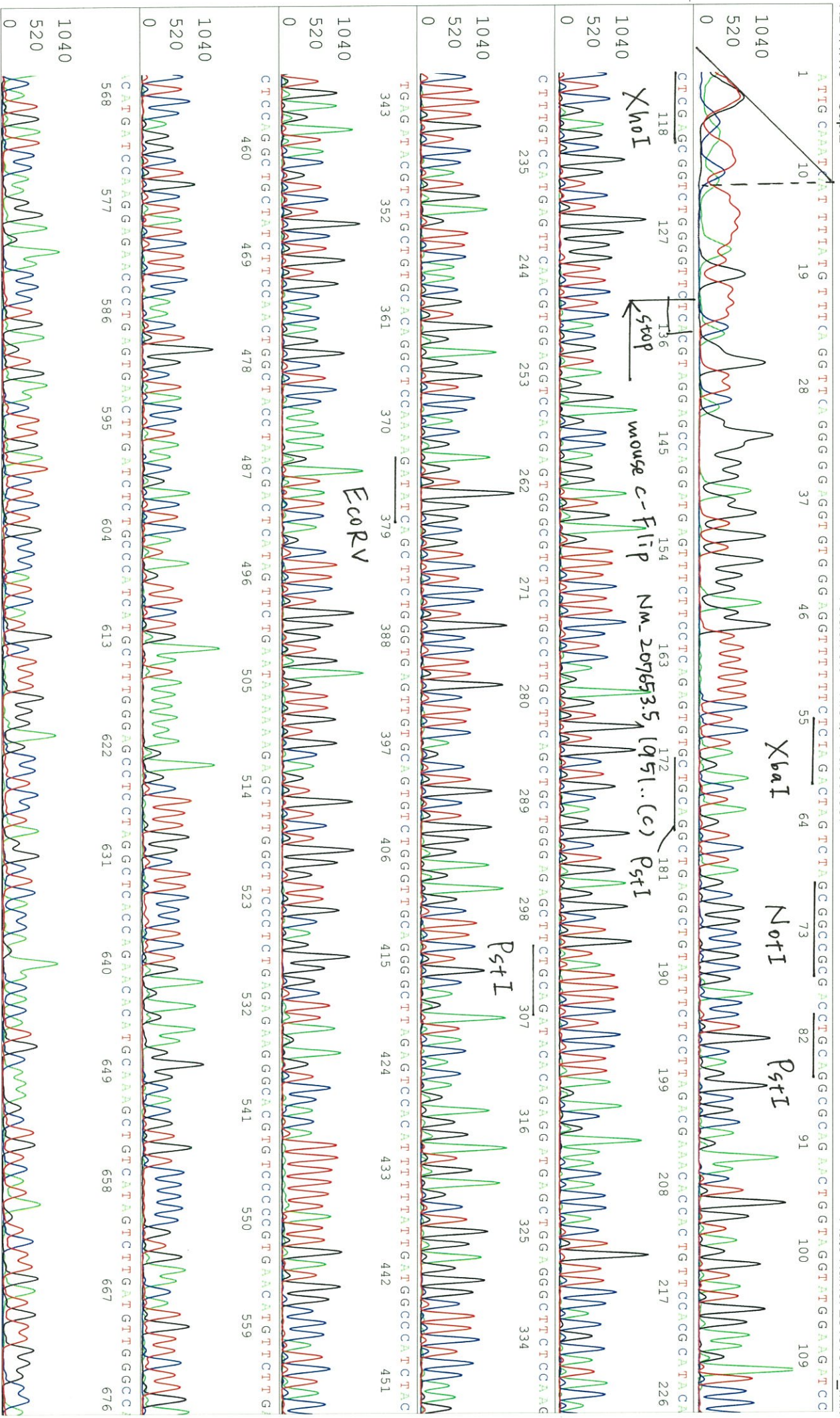
KB.bcp

KB 1.4.1.8 Cap:1



SN G:185 A:149 T:195 C:169 primer name **B** : ME-1290RV
 KB bcp 5'-ATAAGCTGCAATAAACCAAGTTAACAAAC-3' Version 6.0
 KB 1.4.1.8 Cap2

Oct 17, 2016 04:37PM, PDT
 Oct 17, 2016 05:17PM, PDT
 Spacing:12.31 Pts/Panel1350
 Plate Name: 20161017_mix



S/N G:185 A:149 T:195 C:169

KB bcp

KB 1.4.1.8 Cap2

GTACTTGCATATCGGCGAACCAATCTGGGTTATGTCATGTGACTTGGGAAACAAGAAAGCTGGATATGATAGCCCAAGGAAAGTGAAGGTCCTTGAAGATATTTGTG

1040

520

C GTTGC CAATTA CAATCA AATG ATCA AGCA TATCC TTAG GGGG CTTG CTTG CA TCC TGT AA GTCTCTT CACG GATGT GCCG AGG TAAAAA AGCT CC TG ATT CCTGG

793 802 811 820 829 838 847 856 865 874 883

1040

520

VTGGATGTC TTCA CCA G TG TTCTTTGAC TGTCA CCGGTTA TTCC AACAAA TC TTGG CTCTTTAC TTCCGCCCA TTCTGG AGCC TTGAG TTATAC TTGATA C

892 901 910 919 928 937 946 955 964 973 982

.. (159)

1040

520

TC AA TTTT GGAGCG AAGCCCTGC A GAGTA TCA TTA TGA TC TGC TC TTGG CTG ACCTGG CGTACC TCTG ATCT T

991 1000 1009 1018 1027 1036 1045 1054 1063

1040

520

VG TCAEGGTC TATTC TGTGC A GGTC CTTCA GCA GCA TTTTGTGAC AAATTTCC A TTG TTCTC

1072 1081 1090 1099 1108

1040

520

TG AA GCA A

1117