

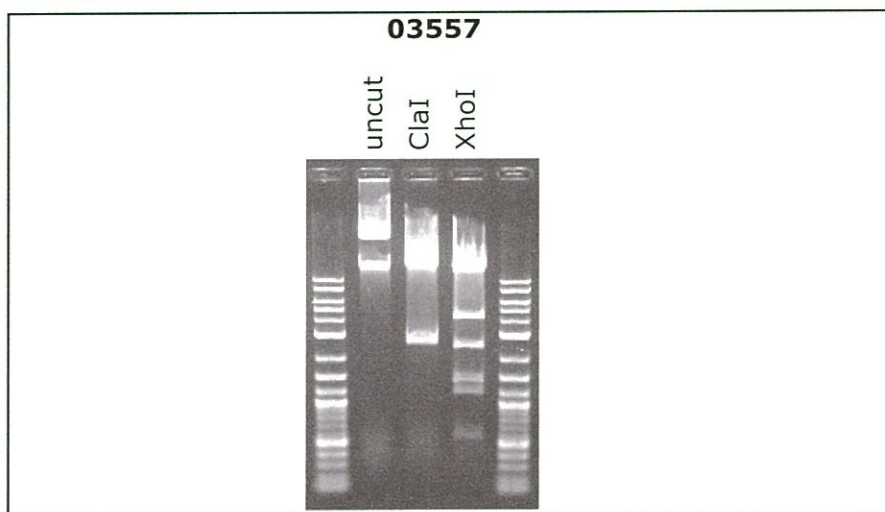
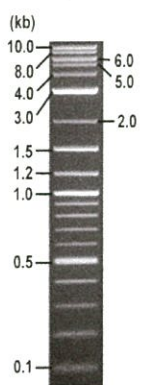


**RIKEN DNA BANK**

clone name : **pAxCALNLhCCNA1 (forward)**

- Clone ID : RDB \_ 03557
- Lot : 03557 \_ B0Fm
- 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 : 500 nanogram DNA per lane ; 1% agarose gel , 1 x TAE Buffer

Restriction enzyme	Expected size of fragment
<u><b>ClaI</b></u>	<u><b>42.5, 2.8, 2.7</b></u> kbp
<u><b>XhoI</b></u>	<u><b>23.5, 14.5, 4.3, 2.5, 1.4, 1.2, 0.6</b></u> kbp
_____	_____ 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	Pr0878	pAxCLNLW_F	SV40 pA, loxP, insert 5'
Sequence - B	Pr0614	pCAGGS_R1	beta-globin pA, insert 3'
Sequence - C	Pr0732	tetpro-B_R	ITR, Ad5 psi
Sequence - D	Pr0164	pAxCA_F1	loxP, NeoR
Sequence - E	-	-	-
Sequence - F	-	-	-
Sequence - G	-	-	-
Sequence - H	-	-	-

APPROVED BY :







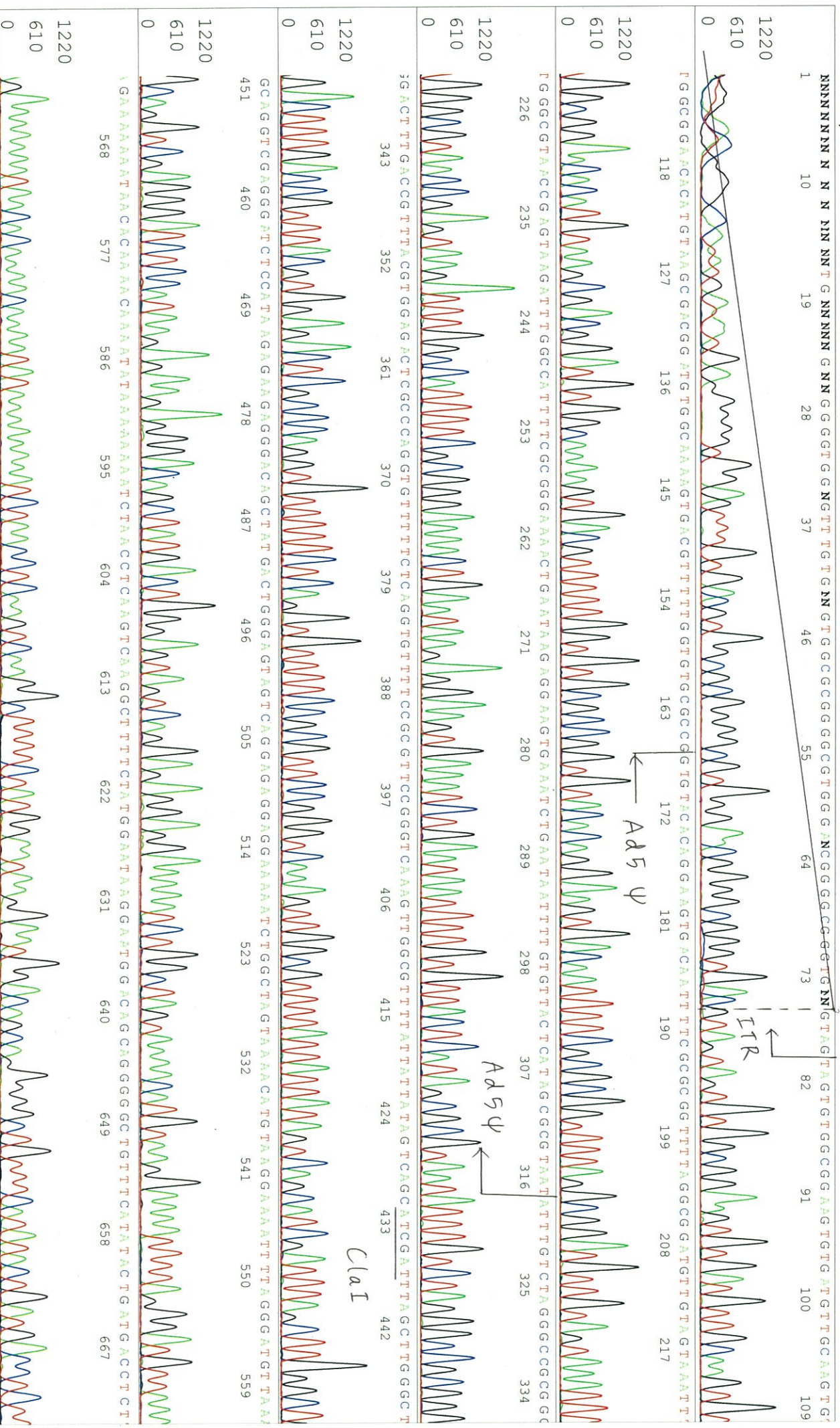


primer name C : tetpro-B\_R

5- TATCAGGAGGCCCTTCGTCT -3'

S/N G:93 A:75 T:67 C:62  
KB.bcp  
KB 1.4.1.8 Cap:4

Spacing: 11.03 Pts/Panel1350  
Plate Name: 20200702mix



S/N G:93 A:75 T:67 C:62

03557\_03557\_B0Fm\_3\_tetpro-B\_R

Jul 02, 2020 02:47PM, JST

KB.bcp

KB\_3500\_POP7\_BDTv3.mob

Jul 02, 2020 03:21PM, JST

KB 1.4.1.8 Cap:4

Pls 1397 to 12041 Pk1 Loc:1374

Spacing: 11.03 Pts/Panel1350

TTATA GCCAACCTTTTGTTCATGGCAGCCAGCATATGGGCATATGTTGCCAAACTCTAAACCAAATACCTCATCTCTGATGTTTAAATGATTTGCCCTCCCATATGTC

676 685 694 703 712 721 730 739 748 757 766 775



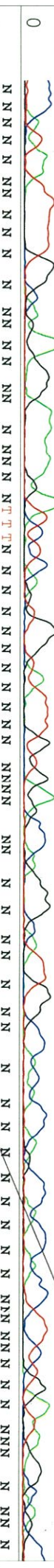
T TCCNAGTGRGAGACACAAAATTCCAACACACTTATTTGCCAATGAAAATAAATTTCTTTATTNAGC NN GAN TCAGANGCTCAANGGCTTTCATGANGTCC

784 793 802 811 820 829 838 847 856 865 874



JCCNTAATTTTGGCAGANGGAAAANATNNCANNNGNNTTTGTGANN CAGGG NNTTN GNCACA CCANNCC NNC NNTNTGANN NN NN NC

883 892 901 910 919 928 937 946 955 964 9



NNNNNNNNNNNNNNNNNNNTTNN

73 982 991 1000 1009 1018 1027 1036 1045



NN

1054 1063 1072 1081 1090 1099



NN

0 1220 610 1099

