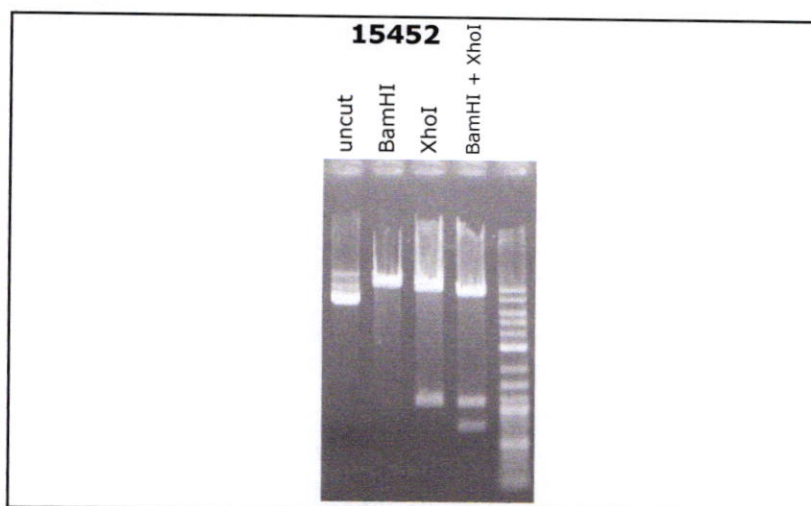
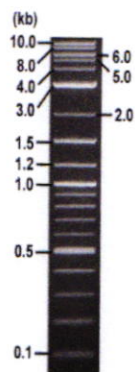


**RIKEN DNA BANK**clone name : **tFucci(CA)2/pCSII-CMV**

- Clone ID : RDB_15452
- Lot : 15452_A7Jv
- DNA Concentration : 25 nanogram/microliter
- Volume : 40 microliter
- Form : DNA solution in TE buffer
- Host : NEB stable
- 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
BamHI	10.8 kbp
XhoI	9.6, 1.1 kbp
BamHI + XhoI	8.9, 1.1, 0.7 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	Pr0674	mRFP-M_R	insert 5'(tFucci(CA)2 5',CMV2)
Sequence - B	Pr0250	PRE_rev	PRE,insert 3'(tFucci(CA)2 3')
Sequence - C	Pr0453	mCherry-C_F	insert mid (tFucci(CA)2 mid)
Sequence - D	Pr0673	HIV-B_R	HIV- psi,5'LTR,CMV1
Sequence - E	Pr0132	HIVgp_R1	HIV- psi,5'LTR
Sequence - F	Pr 725	PPT/CTS-E_R	cTTP_CTS
Sequence - G	Pr0021	CSII-CMV5'	insert mid(tFucci(CA)2 mid)
Sequence - H	Pr 1232	EGFP-B_R2	insert mid (tFucci(CA)2 mid)
Sequence - I	Pr0885	bGH_rev3	3'LTR,PRE
Sequence - J	Pr0110	SV40pro_F	EM7 pro,Zeo,SV40 pA

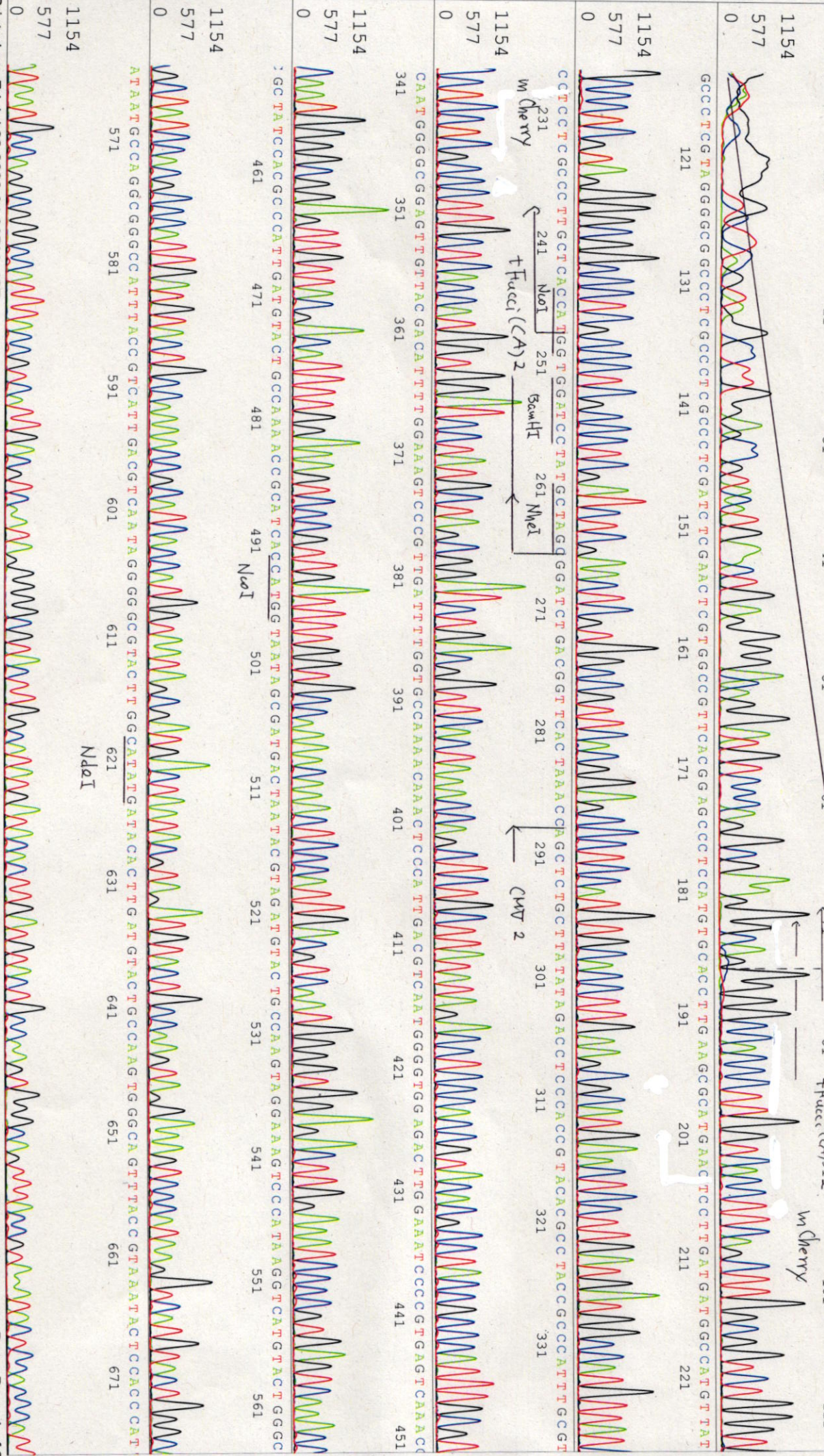
APPROVED BY :



S/N G:46 A:36 T:31 C:36
KB.bcp
KB 1.4.1.8 Cap:8

primer name A : mRFP-M_R
5' GAAGGACAGCTTCAAGTAGTCG 3'

NNNNNNNNNN NN NNNNNN NNN GNC T TG G NNCCG TNNNTG AACTG A GGGGNNAGG FTGTCCCAAGCG AAGGGEAN GGGGCCACCCTTGGTCACTTCAGCTTGCGCGTCTGGGT
1 11 21 31 41 51 61 71 81 91 101 111



0 577 1154
ATAATGCCAAGGGCCATTACCGTCACTT GACGTCATA TAGGGGCGGTACTTGGCATTATGATACACTTGTATGACTTGGCCAAGTGGCA GTTTACCCTAAATAC TCCACCAT
571 581 591 601 611 621 631 641 651 661 671
NdeI



3GGGCGCCACCTGCTTAGAGATTTCCTCCACACTGACTANNANGGTCTGANGGNNCTCTTAGTTACCAGANTCNCAACANACGACGGGCAACACACTACTTGAAGCACTCAAG

5' LTR

3CAAGCTTATTGAGGCTTAAGCAGNGGTTNNCCNANTTAGNNNGANANCTNCNNNGCNCANANCNNGNCTAACNNGANANANCNNNNAANAAANG

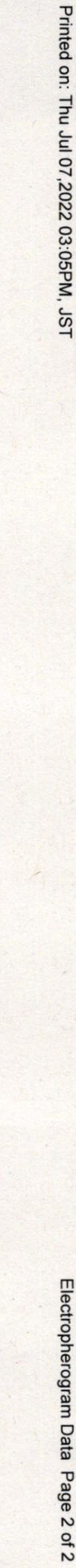
CGCCNNNNNATAANNNNNNCCNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN

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Primer name I : bGH_rev3

5' TTAGGAAAGGACAGCTGGGAGTG 3'



CCACCCCAACTTTGTTTATTCGACGCTTATAATGGTTACCAAAATAAAGCAATAGCATCCAAATTTCCACAAATAAAGCATTTTTTCCACTGCATTCTAGTTGGTTT

ST 40 p (A)

GTCCAAACCTCAATCAATGATGTTATCTTATTCATGCTCTGTATACNGTCNACCTCTANCTAGANCCTGGCGGTAATCATGGTCAATNCTGTTNCCCTGTGAAAT

ST 40 p (A)

TGTTATTNNGCNCNCAATTNCCNCAACAATACNANCNNGAAACATAAAGNGNNNNCNGGGGNNCCNNANNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN

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