

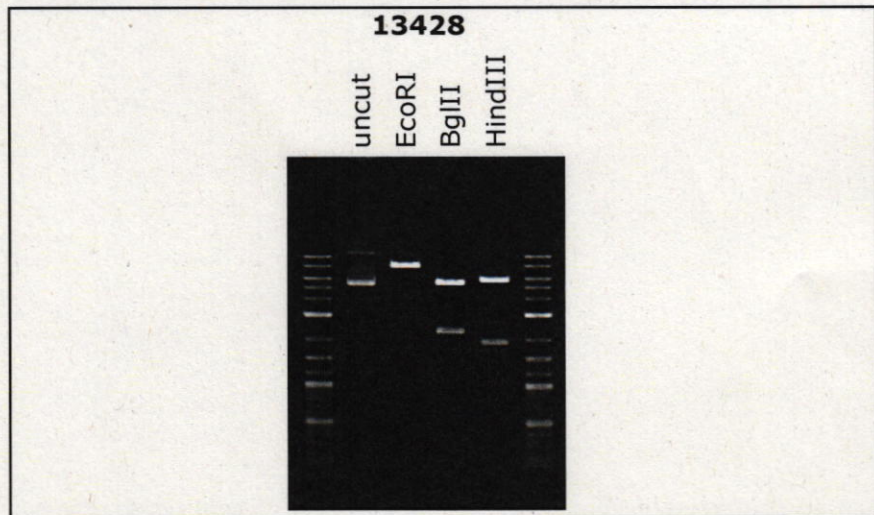
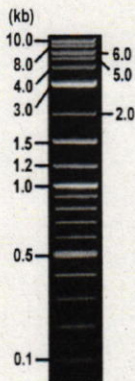


**RIKEN DNA BANK**

clone name : pcDNA3.1/hGnT-V-myc-his

- Clone ID : RDB\_13428
- Lot : 13428\_A6Dq
- 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 : 100 nanogram DNA per lane ; 1% agarose gel , 1 x TAE Buffer

Restriction enzyme	Expected size of fragment
<b>EcoRI</b>	<b>7.7</b> kbp
<b>BglII</b>	<b>5.4, 2.3</b> kbp
<b>HindIII</b>	<b>5.7, 1.9</b> 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	Pr0016	CMV_Forward	CMV pro,insert 5'
Sequence - B	Pr0606	BGH_rev2	bGH pA,6xHis,myc,insert 3'
Sequence - C	Pr0110	SV40pro_F	NeoR
Sequence - D	-	-	-
Sequence - E	-	-	-
Sequence - F	-	-	-
Sequence - G	-	-	-
Sequence - H	-	-	-

APPROVED BY :



primer name A : CMV\_Forward

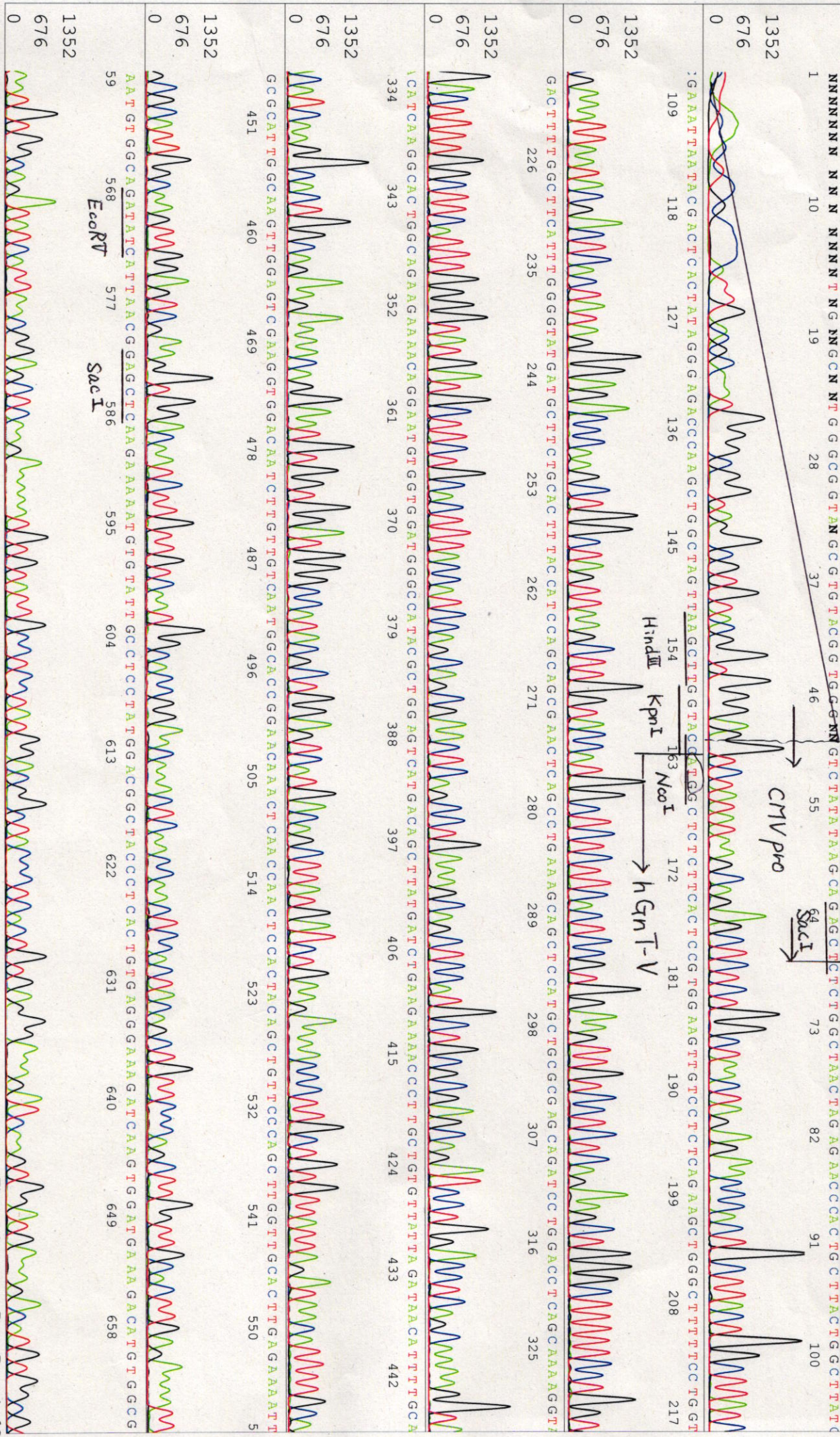
KB\_3500\_POP7\_BDTV3.mob

Pls 1562 to 12379 Pk1 Loc:1539

Version 6.0 HiSeqV Bases: 840

5'-GCACCAGAAATCAACGGGACTT-3'

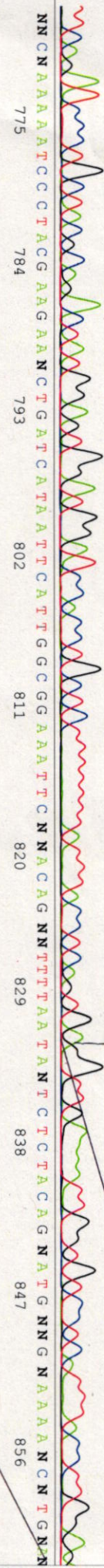
Spacing: 11.04 Pts/Panel1350  
Plate Name: 20160511\_Kitaku



TTCAGATCCCTGCTACTACCGACGACTATTGGAGTGGATGGATCCCACTCTCTCTTTTATTATTACCTCAAGTGAN GTTGAATAATTGGTGTCTCTCACTTACCTTGGAN  
667 676 685 694 703 712 721 730 739 748 757 766

*BamHI*

hGhT-V



1352  
676  
0

NNCAAATCCCTACGAAAGAAANCTGATCTACTATTGCGGAAATTCNNACAGNNTTTAAATANTCTCTACAGNATGNNGNAAAANCNTGN  
775 784 793 802 811 820 829 838 847 856

1352  
676  
0

NNANNCGGNNGNNNNNNNNNANNNNNCNGNNNNNANNN  
865 874 883 892 901 910 919 928 937 946

1352  
676  
0

INNN  
955 964 973 982 991 1000 1009

1352  
676  
0

INNN  
1018 1027 1036 1045 1054

1352  
676  
0









