

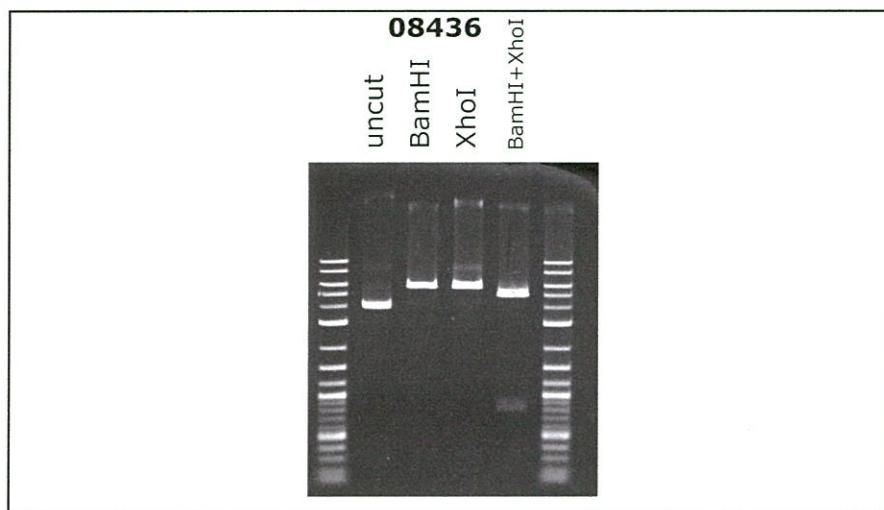
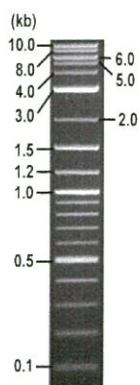


**RIKEN DNA BANK**

clone name : pGEX-mG3-1a

- Clone ID : RDB \_ 08436
- Lot : 19444 \_ B1Kq
- 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>BamHI</b>	<b>5.8</b> kbp
<b>XhoI</b>	<b>5.8</b> kbp
<b>BamHI+XhoI</b>	<b>4.9, 0.8</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	Pr0403	pGEX-F2	GST, thrombin site, insert 5'
Sequence - B	Pr0084	pJB8-Forward	insert 3'
Sequence - C	Pr0724	lacIpro-B_F	lacI pro, lacI
Sequence - D	-	-	-
Sequence - E	-	-	-
Sequence - F	-	-	-
Sequence - G	-	-	-
Sequence - H	-	-	-

APPROVED BY :













S/N G:28 A:24 T:18 C:21

primer name **B** : pJb8-Forward

D08453A7\_B1Kq\_1\_pJb8-Forward

Dec 07, 2021 10:11AM, JST

KB.bcp

5'-ACGAGGCCCTTTCGCTTC-3'

KB\_3500\_POP7\_BDTV3.mob

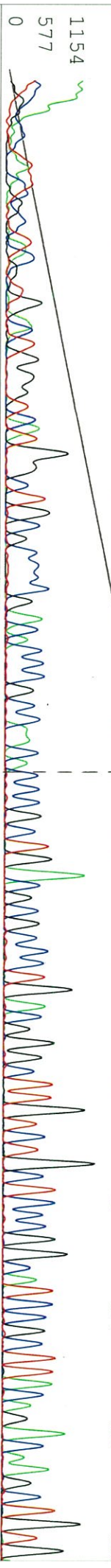
Dec 07, 2021 10:39AM, JST

KB 1.4.1.8 Cap:2

Pts 1413 to 13351 PK1 Loc:1390

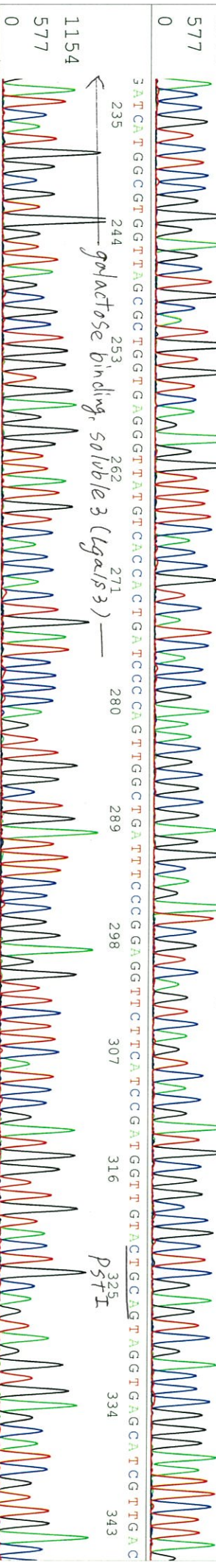
Spacing:11.97 Pts/Panel1350

Version 6.0 HiSeqV Bases: 930



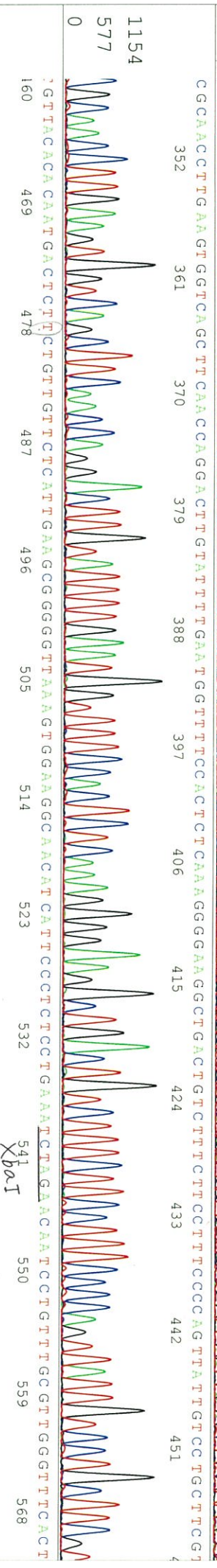
stop codons

NotI XhoI SalI SmaI EcoRI

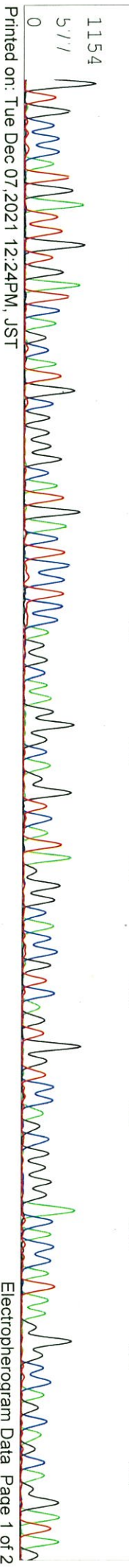
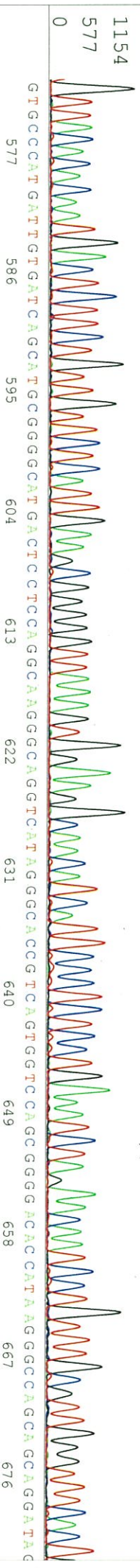


galactose binding, soluble 3 (LgalS3)

PstI



XbaI













S/N G:69 A:43 T:37 C:43

D08453A7\_B1Kq\_1\_Lacpro-B\_F  
KB\_3500\_POP7\_BDTV3.mob

Dec 08, 2021 10:03AM, JST

KB.bcp

Pts 1458 to 13505 PK1 Loc:1435

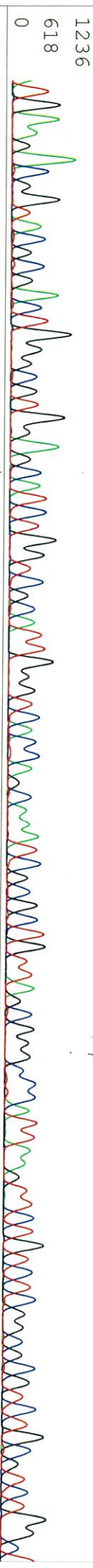
Dec 08, 2021 10:30AM, JST

KB 1.4.1.8 Cap:6

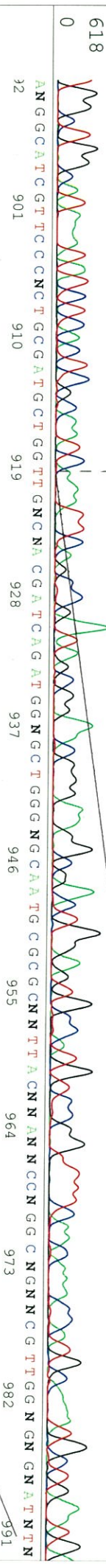
Version 6.0 HISQV Bases: 968

Spacing: 12.0 Pts/Panel/350  
Plate Name: 20211208\_kitaku

ATGAAAGACGGGTACCGCGACTGGGCGGTGGAGGCATCTGGTCGCATTTGGGTCACCAACAATCGCCCTGTTAGCGGGCCCATTAAGTTCGTCTCGCGCCGTCCTGGC	6	685	694	703	712	721	730	739	748	757	766	775	784
									Apal				



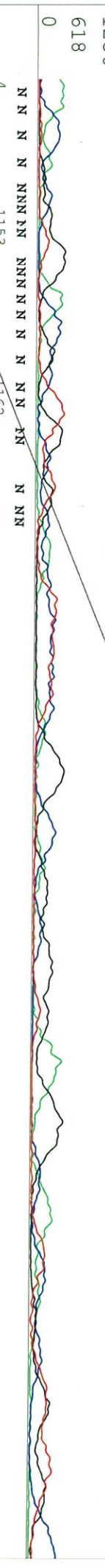
TGGCTGGCATAAATATCTCAGCTCGCAATCAATTCAGCCGATAGCCGGAAGCGGANTTGGAGTGCCATGTCGGNTTCAACNAACCATGCAAAATGCAATG	793	802	811	820	829	838	847	856	865	874	883
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ANGGCATCGTTCCNCCTGCGATGCTGGTTGNCNACGATCAGATGGNGCTGGGNGCAATGCGCGCNCNTTACNNANNCCNGCNGNCCGTTGGNGNATNTN	901	910	919	928	937	946	955	964	973	982	991
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NCGGNANTGGGANNNNNNNANNNNNNNNC TCANGNNNNNNNCNNNNNNNNNNNNNNNNNNNTTNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	1000	1009	1018	1027	1036	1045	1054	1063	1072	1081
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ANNN	1090	1099	1108	1117	1126	1135	1144
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NN	4	1153	1162
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