

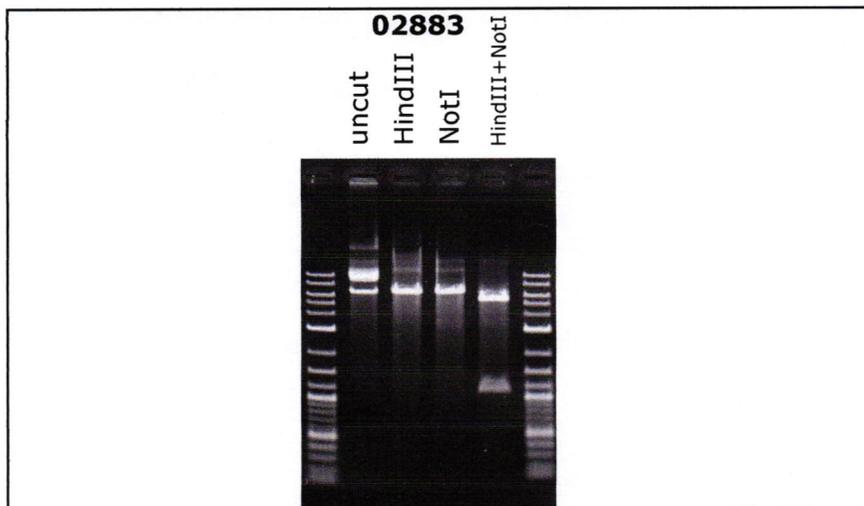
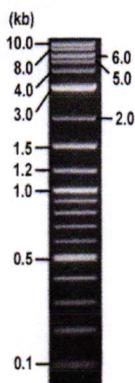


RIKEN DNA BANK

clone name : Human HLA-C*01:02 cDNA

- Clone ID : RDB_02883
- Lot : 02883_02Kq
- DNA Concentration : 25 nanogram/microliter
- Volume : 40 microliter
- Form : DNA solution in TE buffer
- Host : HB101
- 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
HindIII	6.5 kbp
NotI	6.5 kbp
HindIII+NotI	5.4, 1.1 kbp
	kbp

● Confirmation of the insertion sequence

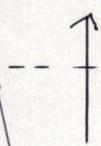
Please be sure to check our sequence analysis results before your request.

Sequence name	Primer ID	Primer name	Confirmed feature
Sequence - A	Pr0016	CMV_Forward	CMV pro, insert 5'
Sequence - B	Pr0885	bGH_rev3	bGH pA,insert 3'
Sequence - C	Pr0731	SV40pro_ori_F	SV40 pro_ori,NeoR
Sequence - D	-	-	-
Sequence - E	-	-	-
Sequence - F	-	-	-
Sequence - G	-	-	-
Sequence - H	-	-	-

APPROVED BY :



HLA-C*01:02 cDNA



3GCCCCCTCCCACTTGCCGCTGGGTGATCTGAGCCCGGGTTCGCCGGCGGTCAGGAGCCGANGTCCCTCGTTCANGGCCGATGTAATCCCTTGCCCGTCCGTANGCCACTGCTC
676 685 694 703 712 721 730 739 748 757 766 775

TACCCCGGAGGANGNCCGTCGCGNCCAGGTCCGANCCNACNTCNACTGGANGNGTGANANCCNGNCTCGCTCTGNNGTANTANCCNCCGNNTTNCNNNGCTC
784 793 802 811 820 829 838 847 856 865 874 883 892

ANNCCNTCANNTGNGCCTNGNCGTNGNCCNTNNGNNCTNCCANNNNNNNNCCANTNNNNNNNNCCNNNNNNTNNNNNCCNNNNNNNNNNNN
901 910 919 928 937 946 955 964 973 982 991 1000

NN
100 1009 1018 1027 1036 1045 1054 1063 1072 1081 1090

NN
1099 1108 1117 1126 1135 1144 1153

NN
1162 1171

1128
564
0

S/N G:499 A:434 T:335 C:438

02883_02883_B3AgF_SV40pro_ori_F

Jan 27, 2023 02:51PM, JST

KB:bcpcap

KB_3500_POP7_BDTV3.mob

Jan 27, 2023 03:21PM, JST

KB 1.4.1.8 Cap:5

Pts 1610 to 14165 PK1 Loc:1587

Spacings: 12.44 Pts/Panel1350

CGTGAACCCATGGCGGATGCTGCTTGGCCGAATATCATGGTGGAATAAGCCCGCTTTCTGGATTCCGACTGTGCCCGCTGGGTGCGGANCCGCTATCCANACATAAGCGT

NcoI

Not

→

0 1188 594

GCTACCCCGTGANATTGCTGAAGAGCTTGNCGCGAANGGCGTGAACCGCTTCCTCGGTCTTACGGNATCCGCCCTCCCGATTCCGACANCGCATCGCCNTCTCGA

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ANNANTTCNTCTGANCCGGNNTCNGGGNTCCGAANTGACCGANCAAGCGANNCCNNNNNGCCNTCNCNANATTTCNATTCNNNGNNGCNATGAANGNGNNN

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CNNANNCNTTTNNNGNNGNN

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