

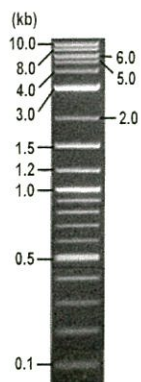


RIKEN DNA BANK

clone name : pGMH10

- Clone ID : RDB _ 01955
- Lot : 13033 _ A4L8
- 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

2-Log DNA Ladder
(NEB#N3200L),
125 ng/well



Electrophoresis : 41 nanogram DNA per lane ; 1% agarose gel , 1 x TAE Buffer

Restriction enzyme	Expected size of fragment
<u>BamHI</u>	<u>4.8</u> kbp
<u>PstI</u>	<u>4.0, 0.8</u> kbp
_____	_____ kbp
_____	_____ kbp
_____	_____ kbp

● Confirmation of the insertion sequence

Sequence name	Primer name
Sequence - A	ADHrev
Sequence - B	HIS3pro_F
Sequence - C	Amp_R
Sequence - D	-

APPROVED BY :



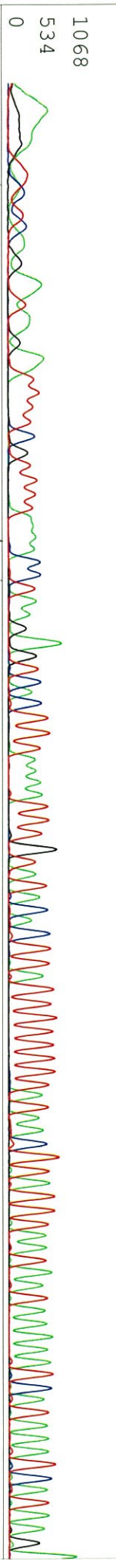
SN G:130 A:187 T:158 C:119 primer name A : ADH rev

KB.bcp

5'-CTATACCTGAGAAAGCAACCTGAC-3'

KB 1.4.1.8 Cap:1

1 G G A G G T T C C A A G A T A G A A T T T C G T T T T A A A C C T A A G A G T C A C T T T A A A A T T T G T A T A C A C T T A T T T T T A A C T T A T T A A T A A A A T C A T A A G



1068 534 0
 A A P T C G C C C G A A T T A G C T T G G C T G C A G G T C G A D T C A C A G A T C C T C T T C G G A A A T C A G T T T C T G T C A G A A T T C G A G C T C G G T A C C C G G G A T C C T A G A G T C A A T T C G A T
 ADH1
 121 131 141 151 161 171 181 191 201 211 221
 ← GAL1, 10 promoter
 PstI SalI Myc EcoRI SacI KpnI BamHI XbaI
 1068 terminator

3 C G G G T T T T T T T C T C T G T G A C G T T A A A G A T A G A G G T A T T A A C A A T T T T T T G T T G A T A C T T T A T T A C A T T T G A A T A A G A A G T A A T A C C G A A A A T G T T G A A A G T A T T A G
 231 241 251 261 271 281 291 301 311 321 331 341

1068 534 0
 T T A A A G T G G T T A T G C A G T T T T T G C A T T T A T A T C T G T T A A T A G A T C A A A A A T C A T C G G T T C G C T G A T T A A T T A C C C C A G A A P T A A G G C T A A A A A A C T A A T C G C A T T A T C A T C C
 41 351 361 371 381 391 401 411 421 431 441 451

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 ← UAS

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 C A G T G C G C C G C G A G G C A C A T C T G C G T T C A G A A C G C G A C C G G T G A A G A C G A G A C G C A C G G A G A G A G T C T T C C T T C G G A G G G C T G T C A C C G C T C G G C G C T T C T A A
 571 581 591 601 611 621 631 641 651 661 671

