

RIKEN Clone ID : W01A062C17

Vector : pENTR/D-TOPO

Gene	MYOD1		
Accession No.	BC064493.1	1699 bp	1..1699
	<i>CDS</i>	963bp	63..1025

● Plasmid DNA purification

Date : 121219

Culture : LB (25 ug/ml Kanamycin) 5 ml -> at 37 deg C over night

Date : 121220

Purification : QIAGEN Miniprep kit -> dH₂O 100 ul

● Digestion by restriction enzyme/Concentration calibration

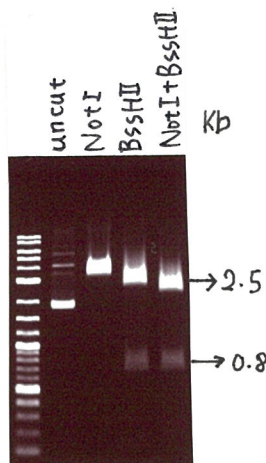
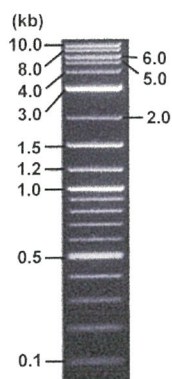
Date : 121220

DNA concentration (O.D.): 69 ng/ul

DNA	0.7	ul
Enzyme (NotI + BssHII)	0.5 + 0.5	ul
Buffer H	1	ul
dH ₂ O	7.3	ul
Total	10	ul

Erectrophoresis : 1% agarose gel, 1x TAE Buffer

Marker : 2-Log DNA Ladder (NEB#N3200L)



<Expected size of fragment from BC064493.1 CDS>

2553, 215, 82, 2, 764 bp

● Confirmation of the insertion sequence

Date : 121225

Primer A	M13
Primer B	T7 long
Primer C	-
List of Sequencing Primers	http://dna.brc.riken.jp/en/GNPclone3en.html

● Adjustment of DNA concentration

Shipping amount : 40 ul

Concentration at the time of preparation of plasmid DNA : 25 ng

Date : 130108

DNA (<u>69</u> ng/ul)	85.0	ul
10x TE	23.5	ul
dH ₂ O	126.1	ul
Total	234.6	ul



APPROVED BY :

<http://dna.brc.riken.jp/index.html>

S/N G:30 A:14 T:17 C:28

Primer A : M13

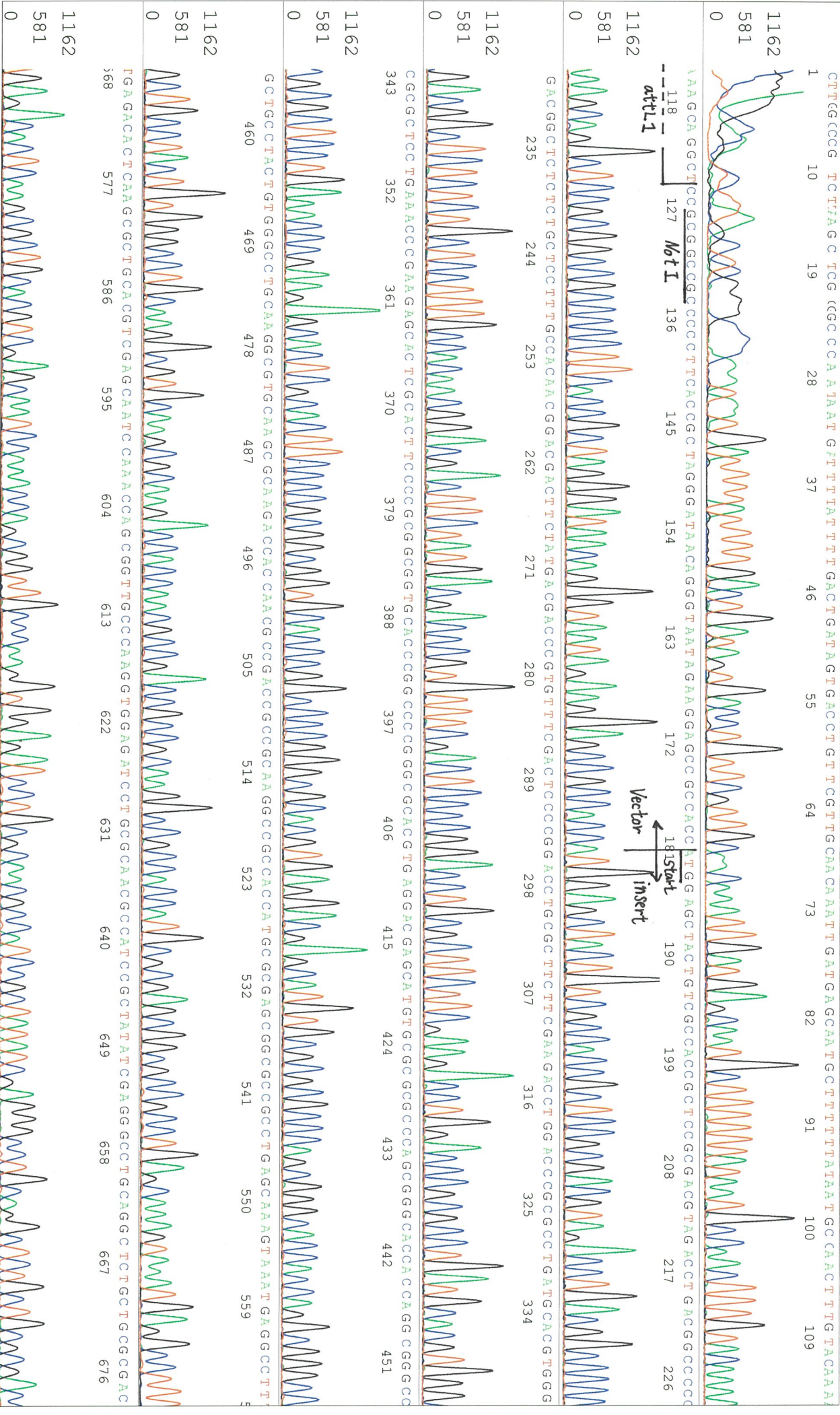
KB.bcp

KB 1.4.1.8 Cap:9

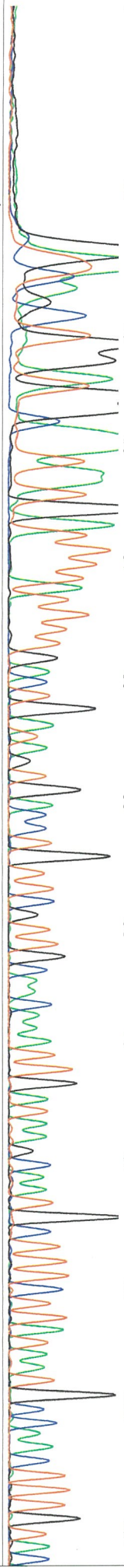
5' GTTTTCCACGTCACGACGTTGTA 3'

W01A062C17_M13(-40)_C06_09
W01A062C17_M13(-40)
KB_3500_POP7_BDTV3.mob
Pls 1284 to 13907 PK1 Loc:1261
Version 5.4 HISQV Bases: 1041

Inst Model/Name 3500/3500 Instrument
Dec 25,2012 05:15PM, JST
Dec 25,2012 05:41PM, JST
Spacing:12.36 Pls/Panel1350
Plate Name: 20121225_GNP



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



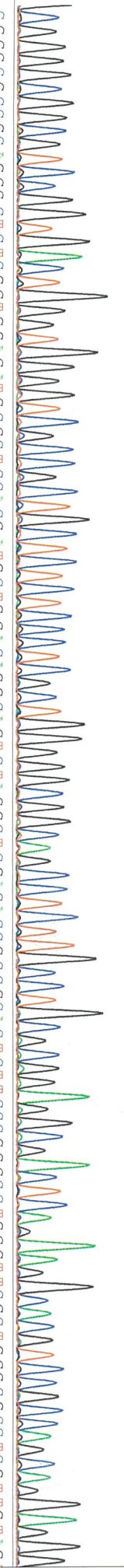
att1.2

BshII

vector

SPB81

226 3CGGGGGGGGTCCTCCGGTACTGGTGGGTCCGCCCTGCTCTCCCTCGCTGGGGGGCGCAAGCCCTCTTGCCCTGGCGCGGAGGC GACTCA GAAGGCACCTCCGCCAAGGAGG



343 CGGGGGGGGCAGGGCTCTCCGGTGAAGATGGCGCTCCAAGATGCTGGAAGCAAGTCTAGGGCTCGAACAAGCCCGCACTCTCCCCGGGCTGGGCTTCGCTGGCGCCCTCGTTGTAG

460 F A G G C C C C T T C G T A G C A G T C C C C C G C C G G C C G C C C C T C G G G G G C C C C T G T A G T C C A T C A T G C C C T C G G A G C A G T T G G A G C C C G G G C T G A C C C T C G G A G T C G C C C C T G T

568 G T G C T C G C C C G C C G C C G C C G G G G C A G C C G G G C C C G C G C A T A G A A G G C C G C T G C G G C C C A G G G G G C C G C C G G C G T C C T G T C C C G C A G C A G C C T G C A G G C C C T C G A T A T A

