

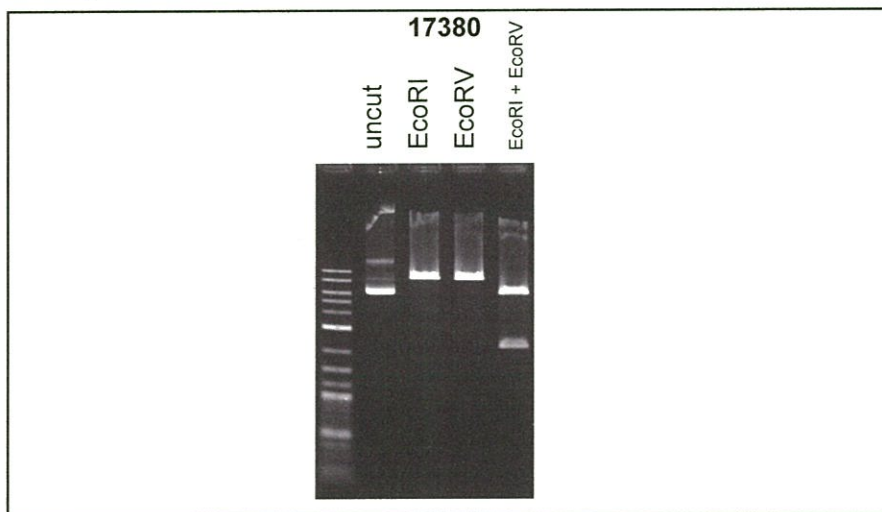
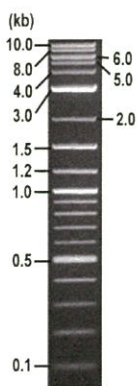


**RIKEN DNA BANK**

clone name : **CDCA7px330**

- **Clone ID** : RDB\_17380
- **Lot** : 17380\_A9Fh
- **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
<u>EcoRI</u>	<u>8.5</u> kbp
<u>EcoRV</u>	<u>8.5</u> kbp
<u>EcoRI + EcoRV</u>	<u>6.2, 2.3</u> 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	Pr0108	SV40pA3'element reverse	U6 pro, gRNA (CDCA7) gRNA scaffold, CBh pro 5'
Sequence - B	Pr0650	Cas9-N_R1	Cas9 5', SV40 NLS, 3xFLAG, CBh pro 3'
Sequence - C	Pr0606	BGH_rev2	bGH p(A), nucleoplasmin NLS, Cas9 3'
Sequence - D	-	-	-
Sequence - E	-	-	-
Sequence - F	-	-	-
Sequence - G	-	-	-
Sequence - H	-	-	-

APPROVED BY :











S/N G:161 A:92 T:96 C:130 primer name B : Cas9-N\_R1

KB.bcp 5'-TGATGCTGTGCCGGTCCGGTG-3'

KB 1.4.1.8 Cap:8

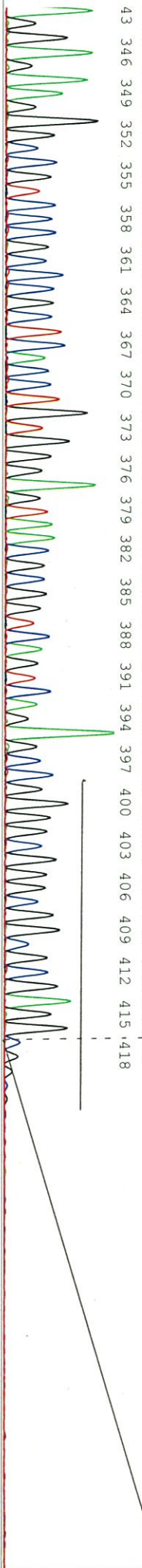
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5 118 121 124 127 130 133 136 139 142 145 148 151 154 157 160 163 166 169 172 175 178 181 184 187 190 193 196 199 202 205 208 211 214 217 220 223 226 2  
 GATACCGACCTTCGGCTTCTTCTTCTTGGGCGCATCTTATCGTCATCGTCTTGTAAACAATATCATGATCCTTGTA GTCTCCGTCGTGGTCCCTTATACTGATGGTGGCACC GG

29 232 235 238 241 244 247 250 253 256 259 262 265 268 271 274 277 280 283 286 289 292 295 298 301 304 307 310 313 316 319 322 325 328 331 334 337 340 3  
 TCCAACTGAAAAAAGTGTTCAGGCAAGGTGCTCCAGGTAATTAACAATTAATACCCCAACCAACCATCCCTTAACCCCTTACCCTTGCTCAAGCTAAATTACAGCCCGG

43 346 349 352 355 358 361 364 367 370 373 376 379 382 385 388 391 394 397 400 403 406 409 412 415 418  
 AGGAGAAAGGGCCGTCCTCCCGCCCGCTCACCTGTGGGAGTAAACGCCGGTCA GTCAAGAGCCGGGGGGGGGGGGGGGGGGAGGNN



1188 594 0  
 1188 594 0

1188 594 0  
 1188 594 0







