

BGT04_T02a-pET16-DEST

5' CACACCTCTCCATTGCAGCAGCCCCGGAAGTGGTGAATTCAGGCGGCATGAACGTCTACATTCTTTTCGACCG
 3' GTGTGGAGAGGTAACGTCGTCGGGCCTTGACCACTAAGTCCGCCGTACTTGCAGATGTAAGAAAGCTGGC

5530

BGT04_T02a

BGT04_T02a-pDONR221

H T S P L Q Q P G T G D S G G M N V Y I L S T
 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71

NheI

BmtI

StuI

5' CGACTGAGCTAGCGAAACAGGGTATCGAGGTCGATATTTACACTCGTGCCACGAGGCCTTCTCAGGGTGA
 3' GCTGACTCGATCGCTTTGTCCCATAGCTCCAGCTATAAATGTGAGCACGGTGCTCCGGAAGAGTCCCACT

5600

BGT04_T02a

BGT04_T02a-pDONR221

A T E L A K Q G I E V D I Y T R A T R P S Q G E
 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95

5' GATCGTGAGAGTAGCTGAGAATTTGCGGGTCATTAATATCGCTGCGGGCCGTATGAGGGGCTTTCCAAA
 3' CTAGCACTCTCATCGACTCTTAAACGCCAGTAATTATAGCGACGCCCCGGCATACTCCCCGAAAGGTTT

5670

BGT04_T02a

BGT04_T02a-pDONR221

I V R V A E N L R V I N I A A G P Y E G L S K
 96 97 98 99 100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118

5' GAGGAGCTTCTACTCAGTTGGCGGCGTTTACCGCGGAATGTTGTCGTTTACGCGCCGGGAGAAGGTTA
 3' CTCCTCGAAGGATGAGTCAACCGCCGAAATGGCCGCCTTACAACAGCAAATGCGCGGCCCTCTTCCAAT

5740

BGT04_T02a

BGT04_T02a-pDONR221

E E L P T Q L A A F T G G M L S F T R R E K V
 119 120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141

5' CTTATGATCTGATCCATTCTCACTATTGGCTGTCTGGTCAGGTGGGGTGGTTGCTGCGGATTTGTGGCG
 3' GAATACTAGACTAGGTAAGAGTGATAACCGACAGACCAGTCCACCCACCAACGACGCGCTAAACACCGC

5810

BGT04_T02a

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T Y D L I H S H Y W L S G Q V G W L L R D L W R
 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156 157 158 159 160 161 162 163 164 165

BGT04_T02a-pET16-DEST

5' GATTCCCCTTATTCATACGGCACACACTTTGGCGGCGGTGAAGAATTCTTATCGGGATGATTCGGACACT
 0 ++++++
 3' CTAAGGGGAATAAGTATGCCGTGTGTGAAACCGCCGCACTTCTTAAGAATAGCCCTACTAAGCCTGTGA

5880

BGT04_T02a

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I P L I H T A H T L A A V K N S Y R D D S D T
 166 167 168 169 170 171 172 173 174 175 176 177 178 179 180 181 182 183 184 185 186 187 188

5' CCGGAGTCGGAGGCGCGTCCGATTTGTGAGCAGCAGCTGGTGGATAACGCTGACGTGTTGGCGGTGAACA
 0 ++++++
 3' GGCCTCAGCCTCCGCGCAGCGTAAACACTCGTCGTCGACCACCTATTGCGACTGCACAACCGCCACTTGT

5950

BGT04_T02a

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P E S E A R R I C E Q Q L V D N A D V L A V N
 189 190 191 192 193 194 195 196 197 198 199 200 201 202 203 204 205 206 207 208 209 210 211

Nsil

5' CTCAGGAGGAGATGCAGGATTTGATGCATCACTACGATGCGGATCCGGATCCGATTTCTGTGGTGTCAAC
 0 ++++++
 3' GAGTCCTCCTCTACGTCCTAAACTACGTAGTGATGCTACGCCTAGGCCTAGCCTAAAGACACCACAGTGG

6020

BGT04_T02a

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T Q E E M Q D L M H H Y D A D P D R I S V V S P
 212 213 214 215 216 217 218 219 220 221 222 223 224 225 226 227 228 229 230 231 232 233 234 235

DrallI

5' GGGTGC GGACGTGGAAC TTTATAGCCCTG GAAATGATCGCGGACGGAACGTTCCCGTCGTGAGCTGGGC
 0 ++++++
 3' CCCACGCCTGCACCTTGAAATATCGGGACCTTTACTAGCGCGCTGCCTTGCAAGGGCAGCACTCGACCCG

6090

BGT04_T02a

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G A D V E L Y S P G N D R A T E R S R R E L G
 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256 257 258

CspCI

CspCI'

AarI

5' ATTCCGCTGCACACAAAGGTAGTGGCTTTTGTGGGTCGGTTGCAGCCGTTTAAGGGCCCGCAGGTGCTGA
 0 ++++++
 3' TAAGGCGACGTGTGTTTCCATCACC GAAAACACCCAGCCAACGTCGGCAAATTC CCGGGCGTCCACGACT

6160

BGT04_T02a

BGT04_T02a-pDONR221

I P L H T K V V A F V G R L Q P F K G P Q V L
 259 260 261 262 263 264 265 266 267 268 269 270 271 272 273 274 275 276 277 278 279 280 281

BGT04_T02a-pET16-DEST

RsrII

5' TCAAGGCGGTTGCGGCGTTGTTTGATCGCGATCCGGACCGAAATCTGCGCGTCATTATTTGTGGCGGCC

3' AGTTCCGCCAACGCCGCAACAACTAGCGCTAGGCCTGGCTTTAGACGCGCAGTAATAAACACCGCCGGG

6230

BGT04_T02a

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I K A V A A L F D R D P D R N L R V I I C G G P

282 283 284 285 286 287 288 289 290 291 292 293 294 295 296 297 298 299 300 301 302 303 304 305

5' TTCTGGTCCGAATGCGACACCGGATACCTATAGGCATATGGCAGAGGAACTGGGCGTCGAAAAGCGAATT

3' AAGACCAGGCTTACGCTGTGGCCTATGGATATCCGTATACCGTCTCCTTGACCCGCAGCTTTTCGCTTAA

6300

BGT04_T02a

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S G P N A T P D T Y R H M A E E L G V E K R I

306 307 308 309 310 311 312 313 314 315 316 317 318 319 320 321 322 323 324 325 326 327 328

5' CGCTTTTGGACCCGCGCCCGCCGAGCGAGCTAGTGGCCGTGTATCGGGCGGCGGACATCGTGGCCGTGC

3' GCGAAAACCTGGGCGCGGGCGGCTCGCTCGATCACCGGCACATAGCCCGCCGCCTGTAGCACCGGCACG

6370

BGT04_T02a

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R F L D P R P P S E L V A V Y R A A D I V A V

329 330 331 332 333 334 335 336 337 338 339 340 341 342 343 344 345 346 347 348 349 350 351

AgeI

5' CAAGTTTTAATGAGTCCTTCGGACTCGTCGCCATGGAGGCGCAAGCCAGCGGCACACCGGTCATTGCGGC

3' GTTCAAATTAATCAGGAAGCCTGAGCAGCGGTACCTCCGCGTTTCGGTCGCCGTGTGGCCAGTAACGCCG

6440

BGT04_T02a

BGT04_T02a-pDONR221

P S F N E S F G L V A M E A Q A S G T P V I A A

352 353 354 355 356 357 358 359 360 361 362 363 364 365 366 367 368 369 370 371 372 373 374 375

XmaI
SmaI

5' CCGGGTTGGCGGCCTGCCCATCGCAGTCGCGGAAGGGGAGACGGGATTGCTTGTGCGACGGCCACTCCCCG

3' GGCCCAACCGCCGGACGGGTAGCGTCAGCGCCTTCCCCTCTGCCCTAACGAACAGCTGCCGGTGAGGGGC

6510

BGT04_T02a

BGT04_T02a-pDONR221

R V G G L P I A V A E G E T G L L V D G H S P

376 377 378 379 380 381 382 383 384 385 386 387 388 389 390 391 392 393 394 395 396 397 398

BGT04_T02a-pET16-DEST

5' CATGCCTGGGCCGACGCCTTAGCCACACTCTTGGACGATGACGAAACGCGCATCAGAATGGGTGAAGACG
 0 ++++++
 3' GTACGGACCCGGCTGCGGAATCGGTGTGAGAACCTGCTACTGCTTTGCGCGTAGTCTTACCCACTTCTGC

6580

BGT04_T02a

BGT04_T02a-pDONR221

H A W A D A L A T L L D D D E T R I R M G E D
 399 400 401 402 403 404 405 406 407 408 409 410 411 412 413 414 415 416 417 418 419 420 421

5' CCGTCGAACACGCCAGAACATTCTCCTGGGCGGCCACCGCCGCACAGCTATCGTTCGCTGTACAACGACGC
 0 ++++++
 3' GGCAGCTTGTGCGGTCTTGTAAGAGGACCCGCCGGTGGCGGCGTGTTCGATAGCAGCGACATGTTGCTGCG

6650

BGT04_T02a

BGT04_T02a-pDONR221

A V E H A R T F S W A A T A A Q L S S L Y N D A
 422 423 424 425 426 427 428 429 430 431 432 433 434 435 436 437 438 439 440 441 442 443 444 445

5' TATTGCCAACGAAAATGTCGACGGTGAACGCATCACGGCTAGGACCCAGCTTTCTTGTACAAAGTGGTT
 0 ++++++
 3' ATAACGGTTGCTTTTACAGCTGCCACTTTGCGTAGTGCCGATCCTGGGTGCAAAGAACATGTTTCACCAA

6720

BGT04_T02a

attB2

BGT04_T02a-pDONR221

I A N E N V D G E T H H G
 446 447 448 449 450 451 452 453 454 455 456 457 458 459

5' GATtcgaggatccggctgctaacaaagcccgaaaggaagctgagttggctgctgccaccgctgagcaata
 0 ++++++
 3' CTAagctcctaggccgacgattgtttcgggctttccttcgactcaaccgacgacggtggcgactcgttat

6790

BspI

T7 Terminator

5' actagcataacccttggggcctctaaacgggtcttgaggggttttttggctgaaaggaggaactatatcc
 0 ++++++
 3' tgatcgtattggggaaccctcgagatttgcccagaactccccaaaaacgactttcctccttgatatagg

6860

T7 Terminator

5' ggatatcccgcaagaggcccggcagtagccgataaccaagcctatgcctacagcatccagggtgacggt
 0 ++++++
 3' cctatagggcgttctccgggccgctcatggccgatttggttcggatacggatgtcgttaggtcccactgcca

6930

5' gccgaggatgacgatgagcgcattgttagatttcatacacgggtgcctgactgcgtagcaatttaactgt
 0 ++++++
 3' cggctcctactgctactcgcgtaacaatctaagtagtgccacggactgacgcaatcgtaaattgaca

7000

HindIII

Clal

5' gataaactaccgcattaaagcttatcgatgataagctgtcaaacatgagaa
 0 ++++++
 3' ctatttgatggcgtaatttcgaatagctactattcgacagtttgactctt

7051