















o  
 5' accgcacctgtggcgccggtgatgccggccacgatgCGTCCGGCGTAGAGGATCGAGATCGATCTCGATC  
 o  
 3' tggcgtggacaccgCGGCCACTACGGCCGGTGCTACGCAGGCCGCATCTCCTAGCTCTAGCTAGAGCTAG

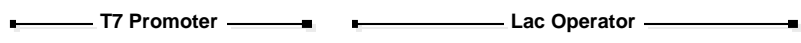
5110

SgrAI

XbaI

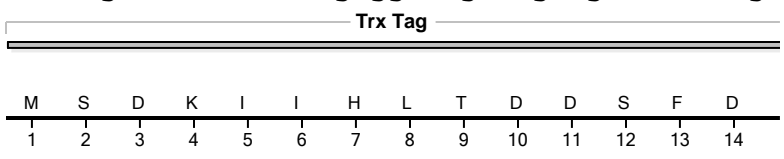
o  
 5' ccgcgaaattaatacGACTCACTATAGGGGAATTGTGAGCGGATAACAATTCCTCTAGAAATAATTTT  
 o  
 3' ggCGCTTAAATTATGCTGAGTGATATCCCTTAACACTCGCCTATTGTTAAGGGGAGATCTTTATTAATAA

5180



o  
 5' gTTTAACTTTAAGAAGGAGATACATATGAGCGATAAAATTATTCACCTGACTGACGACAGTTTGTACA  
 o  
 3' CAAATTGAAATTCCTCCTATATGTATACTCGCTATTTAATAAGTGGACTGACTGCTGTCAAAACTGT

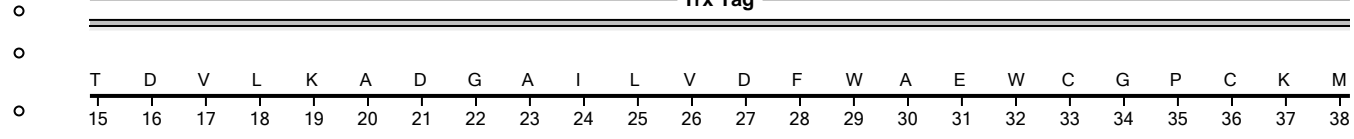
5250



o  
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 o  
 3' gcctacatgagtttcgCCTGCCCGCTAGGAGCAGCTAAAGACCCGCTCTCACCACGCCAGGCAGTTTTA

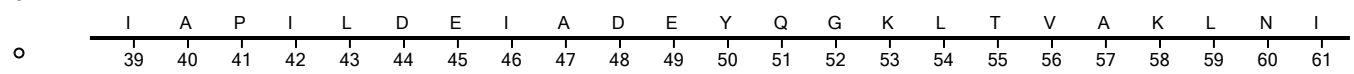
5320

RsrII



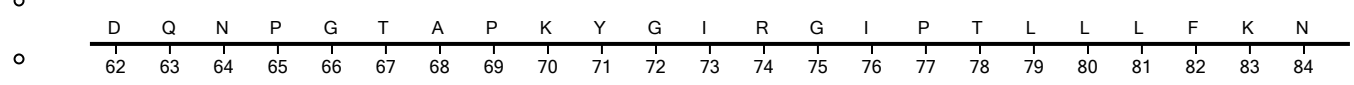
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 5' gatcgcCCCGATTCTGGATGAAATCGCTGACGAATATCAGGGCAAACCTGACCGTTGCAAAACTGAACATC  
 o  
 3' ctagcggggctaagacctactttagcGACTGCTTATAGTCCCGTTTGACTGGCAACGTTTGTACTTGTAG

5390



o  
 5' gatcaaaaccctggcactgcGCCGAAATATGGCATCCGTGGTATCCCGACTCTGCTGCTGTTCAAAAACG  
 o  
 3' ctagttttgggaccgtgacCGGCTTTATACCGTAGGCACCATAGGGCTGAGACGACGACAAGTTTTTGC

5460





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5' gtgaagtggcggcaaccaaagtgggtgcaactgtctaaaggtcagttgaaagagttcctcgacgctaacct  
 3' cacttcaccgcccgttggtttcaccacgtgacagatttccagtcacctttctcaaggagctgcgattgga

Trx Tag

5530

G E V A A T K V G A L S K G Q L K E F L D A N L  
 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101 102 103 104 105 106 107 108

5' ggccggttctgggttctggccatatgcaccatcatcatcattcttctgggtctgggtgccacgcggttct  
 3' ccggccaagaccaagaccggtatacgtggtagtagtagtaagaagaccagaccacgggtgcgccaaga

-T...g

His6-tag      Thrombin

5600

A G S G S G H M H H H H H S S G L V P R G S  
 109 110 111 112 113 114 115 116 117 118 119 120 121 122 123 124 125 126 127 128 129 130 131

BglII      Acc65I  
 KpnI

5' ggatgaaagaaccgctgctgctaaattcgaacgccagcacatggacagcccagatctgggtaccgacg  
 3' ccatactttctttggcgacgacgatttaagcttgcggtcgtgtacctgtcgggtctagacctatggctgc

En...e

5670

G M K E T A A A K F E R Q H M D S P D L G T D  
 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154

NcoI

5' acgacgacaaggccatggcgatATCAACAAGTTTGTACAAAAAAGCAGGCTctgaaaacttgtactttca  
 3' tgctgctgttccggtaccgctaTAGTTGTTCAAACATGTTTTTTCGTCCGAgacttttgaacatgaaagt

5740

Enterokinase      attB1      TEV site

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D D D K A M A I S T S L Y K K A G S E N L Y F Q  
 155 156 157 158 159 160 161 162 163 164 165 166 167 168 169 170 171 172 173 174 175 176 177 178

5' aggcTTTCCTCTAGGTTATAATTATATTACTGTTTGCAAATAATATCTTAAAGGGTGTGGTAAAC  
 3' tccgAAAGGAGATCCAATATTAATAATAATAATGTGACAAACGTTTATTATAGAATTTCCACACCATTG

T... →

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5810

G F P L G Y N Y I I T L F A N N I L K G V V N  
 179 180 181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200 201





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5' AGGATATTGAAGCAGCGGAATTGTCATTTTCTCTGGCGGGAAGGTGGTATCTGGATCCTGTGCTAAAAGG  
 3' TCCTATAACTTCGTCGCCTTAACAGTAAAAGAGACCGCCCTTCCACCATAGACCTAGGACACGATTTTCC

6580

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E D I E A A E L S F S L A G R W Y L D P V L K G  
 435 436 437 438 439 440 441 442 443 444 445 446 447 448 449 450 451 452 453 454 455 456 457 458

5' CCGGTATCCTGAAAACGCATTGAAACTTTATAAAAAGAAGGGTATTGAGCTTCTTTCCCTGAAGATGAC  
 3' GGCCATAGGACTTTTGCGTAACTTTGAAATATTTTCTTCCATAACTCGAAAGAAAGGGACTTCTACTG

6650

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R Y P E N A L K L Y K K K G I E L S F P E D D  
 459 460 461 462 463 464 465 466 467 468 469 470 471 472 473 474 475 476 477 478 479 480 481

BsmI

5' CTGAAACTTATCAGTCAGCCAATAGACTTCATAGCATTCACAATTATTCTTCGGAATTTATAAAATATG  
 3' GACTTTGAATAGTCAGTCGGTTATCTGAAGTATCGTAAGTTGTTAATAAGAAGCCTTAAATATTTTATAC

6720

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L K L I S Q P I D F I A F N N Y S S E F I K Y  
 482 483 484 485 486 487 488 489 490 491 492 493 494 495 496 497 498 499 500 501 502 503 504

AarI  
BspMI

5' ATCCGTCCAGTGAGTCAGGTTTTTTCACCTGCAAACTCCATATTAGAAAAGTTCGAAAAACAGATATGGG  
 3' TAGGCAGGTCAGTCAGTCCAAAAGTGGACGTTTGGAGTATAATCTTTTCAAGCTTTTTTGTCTATACCC

6790

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D P S S E S G F S P A N S I L E K F E K T D M G  
 505 506 507 508 509 510 511 512 513 514 515 516 517 518 519 520 521 522 523 524 525 526 527 528

5' CTGGATCATATATCCTGAAGCCTTGTATGATCTGCTTATGCTCCTTGACAGGGATTATGGAAAGCCAAAC  
 3' GACCTAGTATATAGGACTTCCGAACATACTAGACGAATACGAGGAACGTGCCCTAATACCTTTTCGGTTTG

6860

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W I I Y P E G L Y D L L M L L D R D Y G K P N  
 529 530 531 532 533 534 535 536 537 538 539 540 541 542 543 544 545 546 547 548 549 550 551

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5' ATTGTTATCAGCGAAAACGGAGCCGCCTTCAAAGATGAAATAGGTAGCAACGGAAAGATAGAAGACACAA  
 3' TAACAATAGTTCGCTTTTGCCTCGGCGGAAGTTTCTACTTTATCCATCGTTGCCTTTCTATCTTCTGTGTT

6930

BGH01\_B01b

BGH01\_B01b-pDONR221

I V I S E N G A A F K D E I G S N G K I E D T  
 552 553 554 555 556 557 558 559 560 561 562 563 564 565 566 567 568 569 570 571 572 573 574

5' AGAGAATCCAATATCTTAAAGATTATCTGACCCAGGCTCACAGGGCAATTCAGGACGGTGTAACCTTAAA  
 3' TCTCTTAGGTTATAGAATTTCTAATAGACTGGGTCCGAGTGTCCCGTTAAGTCTGCCACATTTGAATTT

7000

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BGH01\_B01b-pDONR221

K R I Q Y L K D Y L T Q A H R A I Q D G V N L K  
 575 576 577 578 579 580 581 582 583 584 585 586 587 588 589 590 591 592 593 594 595 596 597 598

5' AGCATACTACTTGTGGTTCGCTTTTGGACAACCTTTGAATGGGCTTACGGGTACAACAAGAGATTCGGAATC  
 3' TCGTATGATGAACACCAGCGAAAACCTGTTGAAACTTACCCGAATGCCCATGTTGTTCTCTAAGCCTTAG

7070

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A Y Y L W S L L D N F E W A Y G Y N K R F G I  
 599 600 601 602 603 604 605 606 607 608 609 610 611 612 613 614 615 616 617 618 619 620 621

5' GTTCACGTAAATTTGATACGTTGGAAAGAAAAATAAAGGATAGCGGCTACTGGTACAAAGAAGTAATCA  
 3' CAAGTGCATTTAAACTATGCAACCTTTCTTTTTATTTCCTATCGCCGATGACCATGTTTCTTCATTAGT

7140

BGH01\_B01b

BGH01\_B01b-pDONR221

V H V N F D T L E R K I K D S G Y W Y K E V I  
 622 623 624 625 626 627 628 629 630 631 632 633 634 635 636 637 638 639 640 641 642 643 644

EcoI  
SacI

5' AAAACAACGGTTTTTTAGGACCCAGCTTTCTTGTAACAAGTGGTTGATatcggatccgaattcgagctccg  
 3' TTTTGTGCCCCAAAATCCTGGGTCGAAAGAACATGTTTACCAACTatagcctaggcttaagctcgaggc

7210

BGH01\_B01b

attB2

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K N N G F .  
 645 646 647 648 649 650

