

APP-201

Donor Template SNV -> REV

Protospacer Sequence

SNV

INK2S00119R_APP_V717I_A03_AA
6468 bp

5'
3'

GGAAATTTTCAGAACACCACAATAAAAACAAATTTTGGCAACAAAATGTCACACAAAATTTTTGGCTTCCCAGTACATATAAAAAGTTA
CCTTTAAAGTCTTGTGGTGTATTATTTGTTTAAAACGTTGTTTTACAGTGTGTTTTAAAAAACCGAAGGGTCATGTATATTTTTCAAT

85

APP

APP-201

TGTTGGCCAGGCACGGTGGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCCAAGGCAGGTGGATCACAAGGTCAGGAGGTCGAG
ACAACCGGTCCGTGCCACCGAGTACGGACATTAGGGTCGTGAAACCTCCGGTTCGTTCCGTCACCTAGTGTTCAGTCTCCAGCTC

170

APP

APP-201

AGCATCCTGGCCAACATGGTGAAATGCCATCTCTACTAAAAATACAAAATTAGCTGCATATGGTGGCATGTGCCTGTAGTCCCA
TCGTAGGACCGGTTGTACCACTTTACGGTAGAGATGATTTTTATGTTTTAATCGACGTATACCACCGTACACGGACATCAGGGT

255

APP

APP-201

GCTACTCGGGAGGCTGAGGCAAGAGAATCTCTTGATCCCAGGAGGCAGAGGTTGCAGTGAGCCGAGATCGCACTCCAGCCTGGGC
CGATGAGCCCTCCGACTCCGTTCTCTTAGAGAACTAGGGTCTCCGTTCTCCAACGTCCTCGGCTCTAGCGTGAGGTCGGACCCG

340

APP

APP-201

AACAAGAGCAAGACTCTGTCTCAAAAAAAAAAAAAAAAAAGTTTGTATTTACTATACTCTAGTCTGTTACATGTGCAATAGCATTAT
TTGTTCTCGTTCTGAGACAGAGTTTTTTTTTTTTTCAAACATAAATGTGATATGAGATCAGACAATGTACACGTTATCGTAATA

425

APP

APP-201

GTCTAAAAAAAAAAAAAAAAAGCCATGTATATACTTTACTTTTTATTAGTAAAAATTTTTATTACTAAAAATGCTAATGATCATCTGAGT
CAGATTTTTTTTTTTTTTCGGTACATATATGAAATGAAAAAATCATTTTTATAAAAAAATGATTTTTACGATTACTAGTAGACTCA

510

APP

APP-201

CTTCAGCAAGTCCTAATCTTTTTGCTCGTGGAGAATCTTGCCTTGATGTTCACTGCTGCAGGCTAATCCAGGTGGTGGCTGCTGA
GAAGTCGTTTCAGGATTAGAAAAACGAGCACCTCTTAGAACGGAACCTACAAGTGACGACGTCGATTAGGTCCACCACCGACGACT

595

APP

APP-201

GGTTTGTGGTGGCTGTGGCAATTTCTTAAAAAAGACAATAATGAAGTTTGGCTGCATCGATTGACTCTTCTTTTCACAAAAAGAT
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680

APP

APP-201

GTCTCTGTAGCATGTGATGCTGTTTGATAGCATTTTGCTCACAGTAGAACAGCTTTCAAAATTGGAGTCAGTCTTCGCAAACTCA
CAGAGACATCGTACACTACGACAAAATATCGTAAAACGAGTGTTCATCTTGTGCGAAAGTTTTAACCTCAGTCAGAAGCGTTTTAGAT

765

APP

APP-201

GCCACTGCTTCATCGAGTTTATGTGATATCCTAAATCCTTTGTTGTCATTTCAACAGTGTTTCATAGCATCTTCACCAAGAGTAGA
CGGTGACGAAGTAGCTCAAATACACTATAGGATTTAGGAAACAACAGTAAAGTTGTCACAAGTATCGTAGAAGTGTTTCTCATCT

850

APP

APP-201

TTCCATTGCAAGAAACCCTTTCTTTGCTCATCCATAAGAGGTAACCTCCTCATCCATTCAAGTTTGATCATGAGATGGTAGCAGT
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935

APP

APP-201

TCAGTCATGTCTTTAGGCTCCACTTCTAATTCTAGTTCTCTTGTATTTCACCACATCTGCTGTGACATCCTTTACTGAAGTCT
AGTCAGTACAGAAATCCGAGGTGAAGATTAAGATCAAGAGAACAATAAAGGTGGTGTAGACGACACTGTAGGAAATGACTTCAGA

1020

APP

APP-201

TGAACTCCTCAAAGTCACCCATGAGAGTTGGAATCATTGTCTTCCAAACTCCCCTTATTGCTGATGTTTTGATCTCCCATGGATC
ACTTGAGGAGTTTCAGTGGGTACTCTCAACCTTAGTAACAGAAGGTTTGAGGGCAATAACGACTACAAAACCTAGAGGGGTACCTAG

1105

APP

APP-201

ACAGATGTTCTTAATGGCATCTGGAATGGTGAATCCTTTTTAGAAAGGTTTGCAGTTTACTTTGCCAGATCCATTAGAGGAATCG
TGTCTACAAGAATTACCGTAGACCTTACCACTTAGGAAAAATCTTCCAAACGTCAAATGAAACGGGTCTAGGTAATCTCCTTAGC

1190

APP

APP-201

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1275

APP

APP-201

TTGATCCACAGGCTGCAGAATGGATGTTGTGTTAGCAGGCATGAAAACAACAGTAATCTCCCTGTACCTGTCTTCAGAGCTCTT
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1360

APP

APP-201

GGGTGACTAGGTGCTTTGTTAGTGAGCAGTAATATTTGAAATGAGTCTTTTCTGAGCAGTAGGTCTCAACAGTGGGTTTAAAT
CCCCTGATCCACGAAACAATCACTCGTCATTATAAACTTTACTCGAGAAAAGACTCGTCATCCAGAGTTGTACCCAAATTTTA

1445

APP

APP-201

ATTCAGTAAACCATGCTGTTAAGAGATGTGCTGTCATCCAGTTTTGTTCCATTTATAGAGCATGGGCACAGTAGATTTAGAATA
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1530

APP

APP-201

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1615

APP

APP-201

GTCAGCTTGTCTTTGAAGCTTTGAAGGCAGGACTTCTCTCTAGTGATGAAAGTCTTAGATGGCATTCTTCCAAATAAGTCTG
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1700

APP

APP-201

TTTCATCTCCACTGAAAAATCTGTTATTTAGTGTAGCCACCTTCATCAGGAGTCTTATCTAGATCTTCTGGATAACTTATTGCAGC
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1785

APP

APP-201

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1870

APP

APP-201

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1955

APP

APP-201

GCTTTGGCCTAAGAGAATGTTGTGGTTGGTTTGATCTTCCATCCAGACAACCTAAAAGTTTTCTCCGTATCAGCAATAAGGCTGTT
CGAAACCGGATTCTCTTACAACACCAACCAAACTAGAAGGTAGGTCTGTTGATTTTCAAAGAGGCATAGTCGTTATTCCGACAA

2040

APP

APP-201

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2125

APP

APP-201

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2210

APP

APP-201

CTTCCTTTCACCTTGAATACTTAAGAGGCCACTGTAGGGTTATTTACTGGCCCACTTTCAATATTGTTGTCTCTCAAGGGATAGGG
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2295

APP

APP-201

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2380

APP

APP-201

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2465

APP

APP-201

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2550

APP

APP-201

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2635

APP

APP-201

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2720

APP

APP-201

AACTCCTTATATAACAAGATGATAAAAAGTTGTGGTGTGCTTCTATAATTTTCATTCAAGTAGATAAAGTTGAAAAATAATGACTTG
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2805

APP

APP-201

CTTTTATAAACAGTATGAAGCAATGTAGTGCAGTAAATGAAATTTTATTCTTTCTTTACAATGTTCTCAAAATTATTTTTATGTT
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2890

APP

APP-201

TAATCCAAATAAAGAGCAAGAATAAAGCAACATTTTCAGATTTTGGTTTCTGGAGACAATAGTTAGAAAGCATGAGTTATGAGTGA
ATTAGGTTTATTTCTCGTTCTTATTTTCGTTGTAAGTCTAAAACCAAAGACCTCTGTTATCAATCTTTCGTACTIONACTCACT

2975

APP

APP-201

CTTAAATTTCTTGTTCCTGTACTTCACTTTGAAATAACATTATGCTTTAAAAGCATTACACTGCTAAAGGTTAATTAGAATTC
GAATTTTAAGAACAACGGACATGAAGTGAACCTTTATTGTAATACGAAATTTTTTCGTAATGTGACGATTTCCAATTAATCTTAAG

3060

APP

APP-201

TGCAGAATTACTATAGCTAAAAGTAGGTAACAAGATATCTTTTTTTCTATTGTTTAACTCCTTTGTTTCAGAATGCCTATTCCTG
ACGTCTTAATGATATCGATTTTCATCCATTGTTCTATAGAAAAAAGATAACAAATTGAGGAAACAAAGTCTTACGGATAAGGAC

3145

APP

APP-201

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3230

APP

APP-201

AGGGCTCAGAGTGCTATGACTAGGAAAATGCTGACCTCCTTTTCATTAGTATGATCGTGCCTTTCCAGCTTTTGATAGATCCAAGC
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3315

APP

APP-201

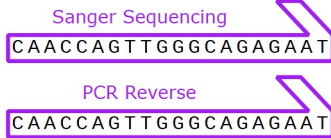
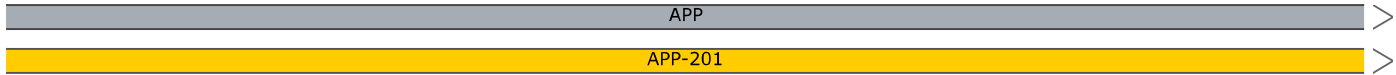
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3400

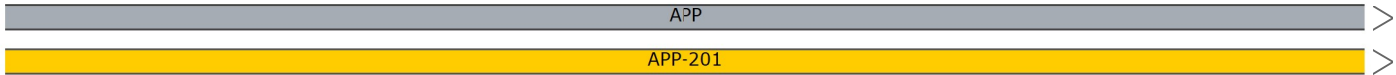
APP

APP-201

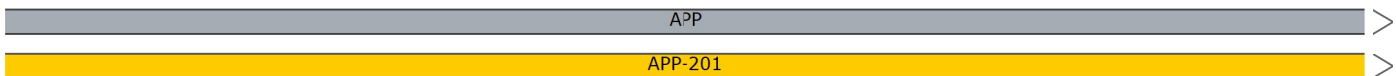
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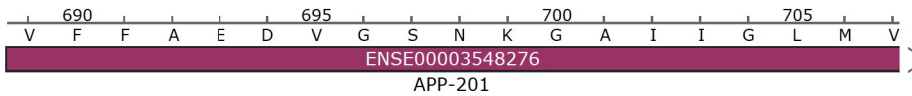
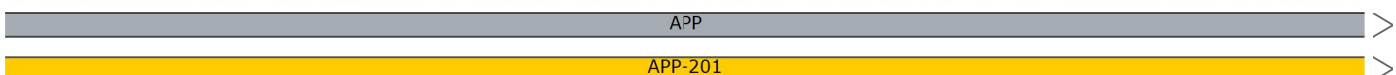
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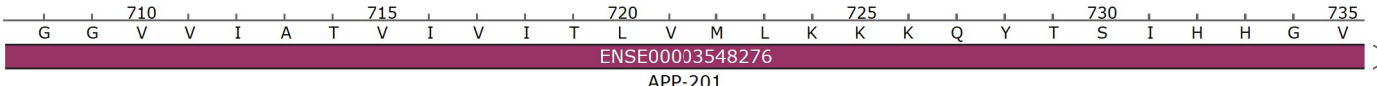
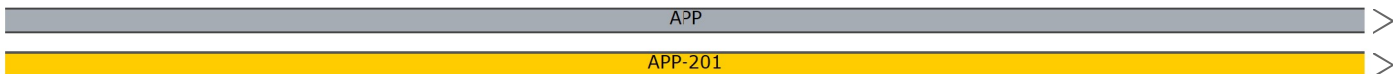
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AAAATTCCTTTTCTTAATTTGTTTTCAAGGTGTTCTTTGCAGAAGATGTGGGTTCAAACAAAGGTGCAATCATTGGACTCATGGT
TTTTAAGAAAAAGAATTAACAAAAGTTCCACAAGAAACGTCTTCTACACCCAAGTTTGTTCACGTTAGTAACCTGAGTACCA



GGCGGTGTTGTCATAGCGACAGTGATCGTCATCACCTTGGTGATGCTGAAGAAGAAACAGTACACATCCATTCATCATGGTGTG
CCCGCCACAACAGTATCGCTGTCACTAGCAGTAGTGGAACCACTACGACTTCTTCTTTGTCATGTGTAGGTAAGTAGTACCACAC



Donor Template SNV -> REV



CCCGCCACAACAGTATCGCTGTCACTAGCAGTAGTGGAACCACTACGACTTCTTCTTTGTCATGTGTAGGTAAGTAGTACCACAC
Donor Template SNV -> REV

GTGGAGGTAGGTAAACTTGACTGCATGTTTCCAAGTGGGAATTAAGACTATGAGAGAATTAGGCTTAGCTTTTTGCTAAGAACTA
CACCTCCATCCATTTGAACTGACGTACAAAGGTTACCCCTTAATTCTGATACTCTCTTAATCCGAATCGAAAAACGATTCTTGAT

3910

APP

APP-201

V E V G K L D C M F P S G N *
ENSE... (in frame with APP-201)
APP-201

Donor Template SNV -> REV

CACCTCcatccattt

Donor Template SNV -> REV

GCTAAGTATCTCTTTTAAAAAACGAATCAGTGTGCTTCCATGATGCTTGGGTTACAGTTGTTCTTTCTTGTTTTGGTTTTTCATTC
CGATTCATAGAGAAAATTTTTGCTTAGTCACACGAAGGTAAGTACTACGAACCCAATGTCAACAAGAAAGAACAAAACCAAAGTAAG

3995

APP

APP-201

ACGAAGGTAAGTACTACGAACCCA
PCR Forward

ATTGCAACTTACCGTGAATATTCTGCTCAAGGTATTGAGAGTGTGTTGTTATCTCAACTTACAATTTGTGTTGAAGTTATCAA
TAACGTTGAATGGCACTTATAAGACGAGTTCCATAACTCTCACACACAACAATAGAGTTGAATGTTAAACACAACCTTCAATAGTT

4080

APP

APP-201

AATAATACAAATGATAATGCATGACTTTAAAAAAGCATGATTACTCTTAGACTTTTTTTTTTCAAACACAAGCAACAAAAACAG
TTATTATGTTTACTATTACGTAAGTAAATTTTTTTCGTAATAAGAGAACTGAAAAAAGTTTGTGTTTCGTTGTTTTTGGTC

4165

APP

APP-201

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4250

APP

APP-201

TTAAATAGTCTGGCCTCTCAATACAACCAGGTAATAGTAAATAATGTGACTTTTCAGTGCAGACTCTTACCTTTAAAGTTTCTCA
AATTTATCAGACCGGAGAGTTATGTTGGTCCATTATCATTTATTACACTGAAAAGTCACGTCTGAGAATGGAAATTTCAAAGAGT

4335

APP

APP-201

GGAAATCACACTATACCAAATGGCCACCAAAAACATACATGATTGTTATTTGAGAGTTCTGCAACCTATTACCCTGCACTTCCAC
CCTTTAGTGTGATATGGTTTACCGGGTGGTTTTGTATGTAACAATAAACTCTCAAGACGTTGGATAATGGGACGTGAAGGTG

4420

APP

APP-201

CTGAAATATCTTAGTTTCAAGGTGCTATACTAAAAACAAGAAAGAGAGAAAGAAAATTTACGAATAAGTTATTTTTATGCATTTGA
GACTTTATAGAATCAAAGTTCACGATATGATTTTTGTTCTTTCTCTTTCTTTTAAATGCTTATTCAATAAAAAATACGTAAACT

4505

APP

APP-201

AATGCCATTTGGATGGGGGAAAAAGTGTTAAAAATCTCACAGTATATGCAACTTTTTAGGAAACTGTCTTGCAGAGAGTATAGTGT
TTACGGTAAACCTACCCCTTTTTTCACAATTTTAGAGTGTACATATACGTTGAAAAATCCTTTGACAGAACGTCTCTCATATCACA

4590

APP

APP-201

AAACATAACTTTTTATATGCACAGGGAAACTGAAAAATTTGTGTGACATTTAATTGCGATATTTTAAGAGTTATGGAATTTTTAGA
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4675

APP

APP-201

GCTGAAAGATTTATTGATGATGACCAATCAAATTGATCTTCTCATCTTGTAGATGAGGGAACTGAGGCCGAGACAGAAGGGTTCA
CGACTTTCTAAATAACTACTACTGGTTAGTTAACTAGAAGAGTAGAACATCTACTCCCTTGACTCCGGCTCTGTCTTCCCAAGT

4760

APP

APP-201

AATTATGCAAGGTCAAGCTCAGAGTCCAGATTTCCCTCATATTTCCCATTTGCTTTTTTTTTTTTTTTTTTTTGGAGACAGTCTCGCGC
TTAATACGTTCCAGTTCGAGTCTCAGGTCTAAAGGGGAGTATAAGGGGTAACGAAAAAAAAAAAAAAAAAACTCTGTCAGAGCGCG

4845

APP

APP-201

TGTTGCCCAGGCTGGATCTTGGCTCACTGCAACCTCTGCCTCCTGGGTTCAAGCAATTCTCTTGTCTCAGCTTCTGAGTAGCTG
ACAACGGGTCCGACCTAGAACCGAGTGACGTTGGAGACGGAGGACCCAAGTTCGTTAAGAGAACAGAGTCGAAGGACTCATCGAC

4930

APP

APP-201

GGACTATGGGCACACGCCACCATGCCCGGCTAATTTTTGTTTTTAGTAGAGACAGGGTTTCACCATATTGGTCAGGCTGGTCTCG
CCTGATACCCGTGTGCGGTGGTACGGGCGGATTAAAAAACAAAAATCATCTCTGTCCCAAAGTGGTATAACCAGTCCGACCAGAGC

5015

APP

APP-201

AACTCCTGACCTCAGGTGATCCACCCGCCTTGGCTTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCCGCGTCTGGCCCCCATT
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5100

APP

APP-201

GCTTTTTCCATCAGAGAAAAAGTCCAGGTGTTTCTAGGGAGTTGAAGATTCCAATATCCCAAGGAAATGGGCTCTAAACCATTCT
CGAAAAAGGTAGTCTCTTTTTCAGGTCCACAAAGATCCCTCAACTTCTAAGGTGATAGGGGTTCTTTTACCCGAGATTTGGTAAGA

5185

APP

APP-201

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5270

APP

APP-201

TGCAGACTACAACAGAAGCTGCTGATGGAATTTGGGGTGGAGCCTCATTTCCCATTCATACTGACGTGGATGAGCAGTGTGAGG
ACGTCTGATGTTGTCTTCGACGACTACCTTTAACCCACCTCGGAGTAAAGGGGTAAGTATGACTGCACCTACTCGTCACACTCC

5355

APP

APP-201

AAGGAAACCAAGAGAACAGCAGGCAGCAGGGTGGAAAGGAACGGACTGACCCATAGATCAGAGTGTGAGCACATAAAAGCAATGA
TTCCTTTGGTTCTCTTGTCTCGTCCGTCGTCCACCTTTCTTGCCTGACTGGGTATCTAGTCTCACACTCGTGTATTTTCGTTACT

5440

APP

APP-201

CCCCACTGTCATGTTTGGCCATTGCACCTTTTGTGATAGTATCTAAGCTTATGTCCCTCAAAACAGTGTAAAGATAGGTTTTTCATT
GGGGTGACAGTACAAACCGGTAACGTGGAAAACACTATCATAGATTCTGAATACAGGGAGTTTTGTACATTCTATCCAAAAGTAA

5525

APP

APP-201

AGTATCAGCTTATTTTCATTTCTGCTCAAATTAATTTTTAAAAAATTTAATTCATATTTGAGCAAATAAAGATGTCTTCATACT
TCATAGTCGAATAAAGTAAAGACGAGTTTAATTAATAATTTTTTAAATTAAGGTATAAACTCGTTTTATTTCTACAGAAGTATGA

5610

APP

APP-201

TGAGTAATTC AAGACCCCTTTTTGCTGCTGTTGTTGTTTTGGTTTTTTTCATGGTTTTGTTGTTTTCTTTTGTGTTGTTGTTT
ACTCATTAAAGTTCTGGGGAAAAAACGACGACAACAACAAAACCAAAAAAAGTACCAAAAACAACAAAAGAAAAACAACAACAACAAA

5695

APP

APP-201

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5780

APP

APP-201

CAGGTTCAAGCGATTCTCCTACCTCACTCTCAGCCTCCTGAGTAACTGGGATTACAGGGGCATACCACCACGCCTGGCTAATTTT
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5865

APP

APP-201

TGTATTTTTAGTAGAGACAGGGTTTTGCCATGTTGGCCAGGCTGGTCTTGAATTCCTGGCCTCAAGTGATCCGCCCTCCCAACC
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5950

APP

APP-201

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6035

APP

APP-201

TTGCCGGTCTGTCCGTTGAAGTTGCCTTCTTTGATGCCAGAGGCCAAAGACTTGGCTTCTCGGGCTTCCATAACAAAGTAACAC
AACGGCCAGACAGGCAACTTCAACGGGAAGAACTACGGTCTCCGGTTTCTGAACCGAAGGAGCCCGAAGGTATTGTTTCATTGTG

6120

APP

APP-201

AGATGGGTGGATTAAGCAACAGAAATTTACCCTCTCACATTTCTGGAGTGTAAAGATCAAGGTATCAGCCAGGTTGGTTCTCCTCCC
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6205

APP

APP-201

AGGCATCTCTCCTTGGTTTGCAGATGCCTGTCTGCTCCCTGTATCTTCACATGACCTTCACTCTGTTTTTCTGCTGTGTCCTAAC
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6290

APP

APP-201

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GAGAAGGATGTTCCCTGTGGTCAGTATCACCTAATCCTAGTTGGGACTACTGGAACAAATTGAAATTATTGGAGAAGCTGGGATAG

6375

APP

APP-201

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6460



































APP

APP-201

ACCCAGT 3'
+-----+ 6468
TGGGGTCA 5'

APP

APP-201

Feature		Location	Size			Type
✓ APP		1 .. 6468	6468 bp			gene
/note	= gene ENSG00000142192 Protein coding					
APP-DT		1 .. 6468	6468 bp			gene
/note	= gene ENSG00000273492 lncRNA					
✓ APP-201		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000346798					
APP-202		1 .. 6468	6468 bp			prim_transcript
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/note	= primary transcript ENST00000358918					
APP-206		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000359726					
APP-208		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000439274					
APP-209		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000440126					
APP-213		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000464867 Retained intron					
APP-DT-201		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000608591 lncRNA					
APP-DT-202		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000609365 lncRNA					
APP-DT-203		1 .. 6468	6468 bp			prim_transcript
/note	= primary transcript ENST00000664668 lncRNA					
✓ APP-201		3685 .. 3831	147 bp			CDS
/note	= coding sequence ENSP00000284981					
/translation	= VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE 49 amino acids = 5.2 kDa					
APP-202		3685 .. 3831	147 bp			CDS
/note	= coding sequence ENSP00000345463					
/translation	= VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE 49 amino acids = 5.2 kDa					
APP-203		3685 .. 3831	147 bp			CDS
/note	= coding sequence ENSP00000346129					
/translation	= VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE 49 amino acids = 5.2 kDa					
APP-204		3685 .. 3831	147 bp			CDS
/note	= coding sequence ENSP00000350578					
/translation	= VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE 49 amino acids = 5.2 kDa					
APP-205		3685 .. 3831	147 bp			CDS
/note	= coding sequence ENSP00000351796					
/translation	= VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE 49 amino acids = 5.2 kDa					

Feature		Location	Size			Type
APP-206		3685 .. 3831	147 bp			CDS
/note	= coding sequence	ENSP00000352760				
/translation	=	VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE				
		49 amino acids = 5.2 kDa				
APP-208		3685 .. 3831	147 bp			CDS
/note	= coding sequence	ENSP00000398879				
/translation	=	VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE				
		49 amino acids = 5.2 kDa				
APP-209		3685 .. 3831	147 bp			CDS
/note	= coding sequence	ENSP00000387483				
/translation	=	VFFAEDVGSNKGAIIGLMVGGVVIATVIVITLVMLKKKQYTSIHGGVVE				
		49 amino acids = 5.2 kDa				
✓	Donor Template SNV -> REV	3741 .. 3840	100 bp			misc_feature
✓	Protospacer Sequence	3759 .. 3778	20 bp			misc_feature
✓	SNV	3769 .. 3769	1 bp			misc_feature
	/note	= REV = G SNV = A				
✓	PAM	3779 .. 3781	3 bp			misc_feature

Primer	Length	Binding Sites	Tm	Date Added
✓ PCR Reverse /sequence = CAACCAAGTTGGGCAGAGAAT 50% GC / 6175.1 Da	20-mer	3530 .. 3549	57°C	Jan 10, 2023
✓ Sanger Sequencing /sequence = CAACCAAGTTGGGCAGAGAAT 50% GC / 6175.1 Da	20-mer	3530 .. 3549	57°C	Jan 10, 2023
✓ Donor Template SNV -> REV /sequence = ttacctacCTCCACCACACCATGATGAATGGATGTGTACTGTTTCTTCTTCAGCATCACCAAGGTGATGACGATCACTGTCGCTATGACAA 40% GC / 30,542.8 Da	100-mer	3741 .. 3840	77°C	Jan 10, 2023
✓ gRNA Protospacer /sequence = GACAGTGATCATCATCACCT 45% GC / 6061.0 Da	20-mer	3759 .. 3778	47°C	Jan 10, 2023
✓ PCR Forward /sequence = ACCCAAGCATCATGGAAGCA 50% GC / 6104.0 Da	20-mer	3943 .. 3962	57°C	Jan 10, 2023