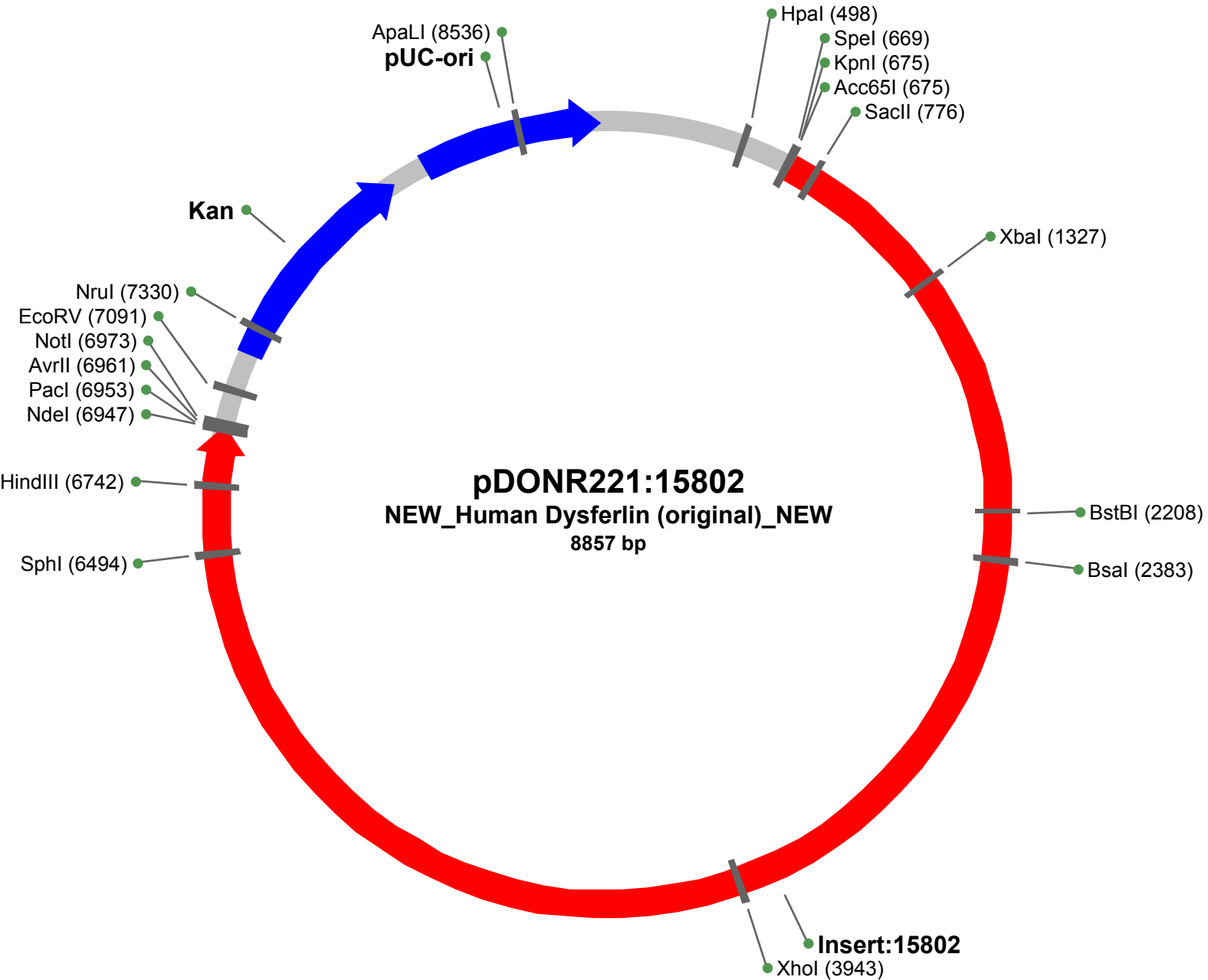


Plasmid Map

pDONR221:15802 - NEW_Human Dysferlin (original)_NEW

Only single cutters are shown in the map, for a more complete list see table below.

pDONR221 is a Gateway® vector



Original Author

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Feature Map

- Insert:15802 - Start:682 End:6968
- Kan - Start:7251 End:8060
- pUC-ori - Start:8180 End:8854

Restriction Map

Name	Sequence	Cut Positions
Acc65I	GGTACC	676
AlwNI	CAGNNNCTG	1679,2198,2872,2920,4355,5980,8442
Apal	GGGCCC	567,2262,4048,5936
ApaLI	GTGCAC	8537
Aval	CYCGRG	560,1240,1295,1595,3944,4185
AvrII	CCTAGG	6962
BamHI	GGATCC	1461,4593,5913
BbsI	GAAGAC	1182,1785,2015,3515,4016,4268,5070,5376,437(C),706(C),1849(C)
BglI	GCCNNNNNGGC	2792,3672,4935
BglII	AGATCT	1914,1998,4329,5514,6429
BsaI	GGTCTC	2378(C)
BsmBI	CGTCTC	4096,4243,7692,914(C),5678(C)
BspEI	TCCGGA	1605,3502,4824
BsrDI	GCAATG	632,6357(C),7019(C),7174(C)
BstBI	TTCGAA	2210
BstXI	CCANNNNNNTGG	2395,2738,2908,3675,4375,4565,4774,6925
BtsI	GCAGTG	214,2804,7624,2042(C),7537(C)
Clal	ATCGAT	4920,5127
EagI	CGGCCG	693,3465,6975
EcoRV	GATATC	7094
HindIII	AAGCTT	6743
HpaI	GTTAAC	501
KasI	GGCGCC	1389,3236,3984,6759
KpnI	GGTACC	680
MluI	ACGCGT	230,8155
NdeI	CATATG	6949
NheI	GCTAGC	239,505
NotI	GCGGCCGC	6975
NruI	TCGCGA	7333
PacI	TTAATTAA	6958

PstI	CTGCAG	1062,1441,2194,2517,2864,3004,3099,4024,5475,6027,6054,6480
PvuI	CGATCG	692,7676
PvuII	CAGCTG	174,1396,2734,2836,2929,4025,4747,5227,5410,6341,6541,6942,7089
SacI	GAGCTC	2328,2850,2997,4530,4890
SacII	CCGCGG	780
SalI	GTCGAC	5693,6062
SanDI	GGGWCCC	3784,5423
SpeI	ACTAGT	670
SphI	GCATGC	6499
XbaI	TCTAGA	1328
XhoI	CTCGAG	3944
XmaI	CCCGGG	1240,1295,4185

No Cuts: AgeI, AscI, EcoRI, MfeI, NcoI, SfiI, SnaBI

Sequence

1 CTTTCTGCG TTATCCCCTG ATTCTGTGGA TAACCGTATT ACCGCCTTTG AGTGAGCTGA TACCGCTCGC
 71 CGCAGCGAA CGAGTCGCG CAGCGAGTCA GTGAGCGAGG AAGCGGAAGA GCGCCCAATA CGCAAACCGC
 141 CTCTCCCCGA GCGTTGGCCG ATTCATTAAT GCAGCTGGCA CGACAGGTTT CCCGACTGTA AAGCGGGCAG
 211 TGAGCGCAAC GCAATTAATA CGCGTACCGC TAGCCAGGAA GAGTTTGTAG AAACGCAAAA AGGCCATCCG
 281 TCAGGATGGC CTTCTGCTTA GTTTGATGCC TGGCAGTTTA TGGCGGGCGT CCTGCCCGCC ACCCTCCGGG
 351 CCGTTGCTTC ACAACGTTCA AATCCGCTCC CGGCGGATTT GTCCTACTCA GGAGAGCGTT CACCGACAAA
 421 CAACAGATAA AACGAAAGGC CCAGTCTTCC GACTGAGCCT TTCGTTTTAT TTGATGCCTG GCAGTTCCTT
 491 ACTCTCGCGT TAACGCTAGC ATGGATGTTT TCCCAGTCAC GACGTTGTAA AACGACGGCC AGTCTTAAGC
 561 TCGGGCCCCA AATAATGATT TTATTTTGAC TGATAGTGAC CTGTTGTTG CAACACATTG ATGAGCAATG
 631 CTTTTTTTATA ATGCCAACTT TGTACAAAAA AGCAGGCTCA CTAGTGGTAC CGTTTTAAACG ATCGGCCGCC
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 771 ACTGCTCCGC GGTGTTTGCA GGGGTGAAGA AGAGAACCAA AGTCATCAAG AACAGCGTGA ACCCTGTATG
 841 GAATGAGGGA TTTGAATGGG ACCTCAAGGG CATCCCCCTG GACCAGGGCT CTGAGCTTCA TGTGGTGGTC
 911 AAAGACCATG AGACGATGGG GAGGAACAGG TTCCTGGGGG AAGCCAAGGT CCCACTCCGA GAGGTCTCTG
 981 CCACCCTAG TCTGTCCGCC AGCTTCAATG CCCCCTGCT CCGCACCAAAG AACGAGCCCA CAGGGGCTCG
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7281 CCGCGATTAA ATTCCAACAT GGATGCTGAT TTATATGGGT ATAAATGGGC TCGCGATAAT GTCGGGCAAT
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7561 CATTCCAGGT ATTAGAAGAA TATCCTGATT CAGGTGAAAA TATTGTTGAT GCGCTGGCAG TGTTCCTGCG
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7701 CAATCACGAA TGAATAACGG TTTGGTTGAT GCGAGTGATT TTGATGACGA GCGTAATGGC TGGCCTGTTG
7771 AACAAGTCTG GAAAGAAATG CATAAACTTT TGCCATTCTC ACCGGATTCA GTCGTCACTC ATGGTGATTT
7841 CTCACTTGAT AACCTTATTT TTGACGAGGG GAAATTAATA GGTGTATTG ATGTTGGACG AGTCGGAATC
7911 GCAGACCGATA ACCAGGATCT TGCCATCCTA TGGAACTGCC TCGGTGAGTT TTCTCCTTCA TTACAGAAAC
7981 GGCTTTTTTCA AAAATATGGT ATTGATAATC CTGATATGAA TAAATTGCAG TTTCATTTGA TGCTCGATGA
8051 GTTTTTTCTAA TCAGAATTGG TTAATTGGTT GTAACACTGG CAGAGCATTG CGCTGACTTG ACGGGACGGC
8121 GCAAGCTCAT GACCAAAATC CCTTAACGTG AGTTACGCGT CGTTCCACTG AGCGTCAGAC CCCGTAGAAA
8191 AGATCAAAGG ATCTTCTTGA GATCCTTTTTT TTCTGCGCGT AATCTGCTGC TTGCAAACAA AAAAACCACC
8261 GCTACCAGCG GTGGTTTGTG TGCCGGATCA AGAGCTACCA ACTCTTTTTT CAAAGGTAAC TGGCTTCAGC
8331 AGAGCGCAGA TACCAAATAC TGTTCTTCTA GTGTAGCCGT AGTTAGGCCA CCACCTCAAG AACTCTGTAG
8401 CACCGCCTAC ATACCTCGCT CTGCTAATCC TGTTACCAGT GGCTGCTGCC AGTGGCGATA AGTCTGTCT
8471 TACCGGGTTG GACTCAAGAC GATAGTTACC GGATAAGGCG CAGCGGTCGG GCTGAACGGG GGGTTCGTGC
8541 ACACAGCCCA GCTTGGAGCG AACGACCTAC ACCGAACTGA GATACCTACA GCGTGAGCTA TGAGAAAGCG
8611 CCACGCTTCC CGAAGGGAGA AAGGCGGACA GGTATCCGGT AAGCGGCAGG GTCGGAACAG GAGAGCGCAC
8681 GAGGGAGCTT CCAGGGGGA ACGCCTGGTA TCTTTATAGT CCTGTCCGGT TTCGCCACCT CTGACTTGAG
8751 CGTCGATTTT TGTGATGCTC GTCAGGGGG CCGAGCCTAT GAAAAACGC CAGCAACGCG GCCTTTTTAC
8821 GGTTCCTGGC CTTTTGCTGG CCTTTTGTCT ACATGTT

Only the synthesized DNA fragment (in red) has been sequence verified. We do not guarantee the vector sequence.