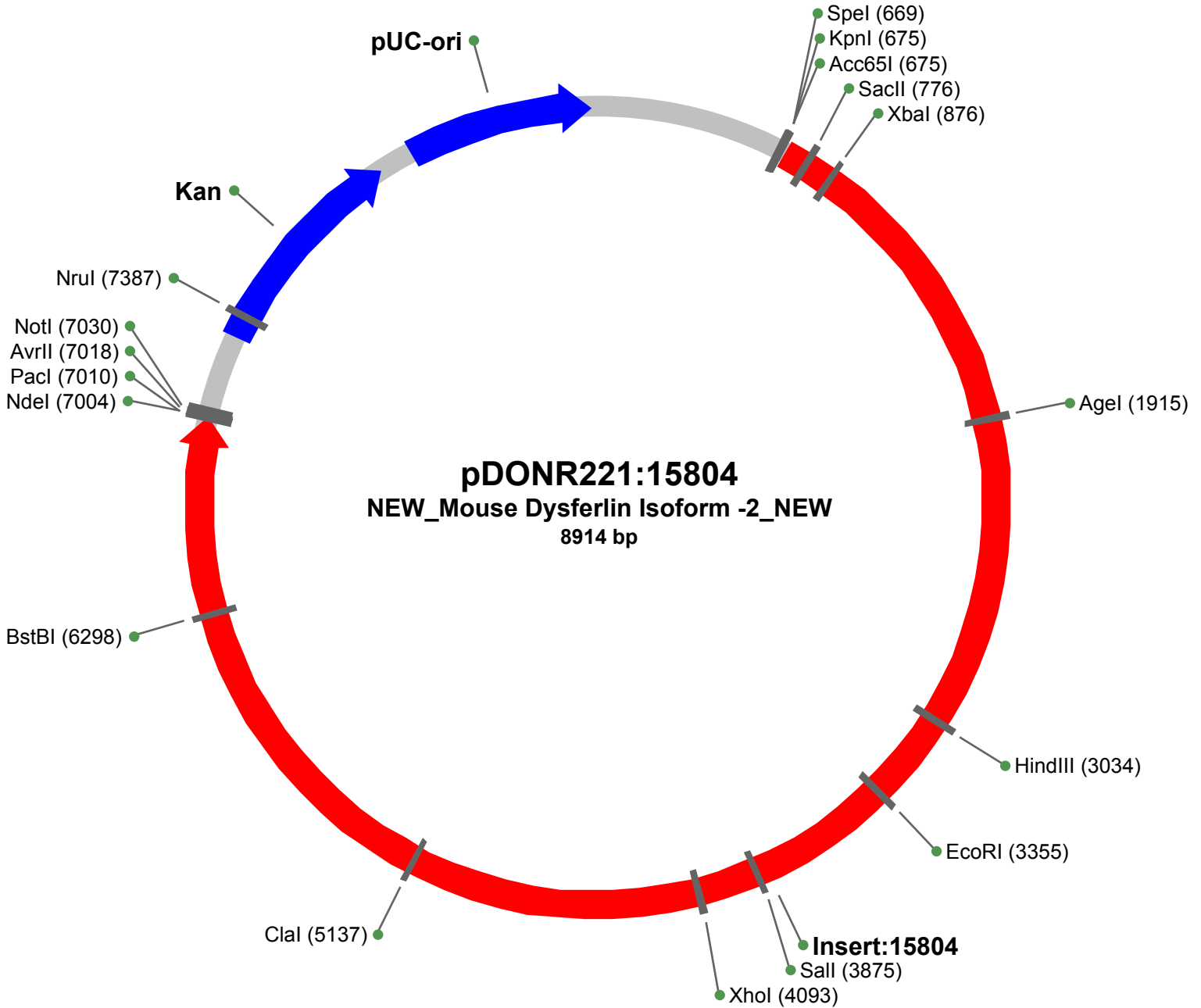


Plasmid Map

pDONR221:15804 - NEW_Mouse Dysferlin Isoform -2_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:15804 - Start:682 End:7025
- Kan - Start:7308 End:8117
- pUC-ori - Start:8237 End:8911

Restriction Map

| Name | Sequence | Cut Positions |
|---------|--------------|--|
| Acc65I | GGTACC | 676 |
| AgeI | ACCGGT | 1916 |
| AlwNI | CAGNNNCTG | 1088,1288,1538,2929,2977,3228,3240,8499 |
| Apal | GGGCC | 567,1340,2319,3221,3298,4105,5993 |
| ApaI | GTGCAC | 6326,8594 |
| AvaI | CYCGRG | 560,2346,4094,4242,4542,5114,5449 |
| AvrII | CCTAGG | 7019 |
| BamHI | GGATCC | 1560,2050,6808 |
| BbsI | GAAGAC | 1275,1884,2114,2484,2763,3572,4073,4140,4325,6323,6653,437(C),706(C),6472(C) |
| BglI | GCCNNNNNGGC | 2885,3729,4992,6572 |
| BglII | AGATCT | 2013,4386,5571,6486 |
| BsaI | GGTCTC | 5405,2435(C),4239(C) |
| BsmBI | CGTCTC | 4300,7749,914(C),4655(C),5735(C) |
| BspEI | TCCGGA | 1704,1752 |
| BsrDI | GCAATG | 632,7076(C),7231(C) |
| BstBI | TTCGAA | 6300 |
| BstXI | CCANNNNNNTGG | 1921,2239,2452,3064,3837,4831,5832,6982 |
| BtsI | GCAGTG | 214,7681,1133(C),3057(C),7594(C) |
| Clal | ATCGAT | 5139 |
| EagI | CGGCCG | 693,7032 |
| EcoRI | GAATTC | 3356 |
| EcoRV | GATATC | 2497,6091,7151 |
| HindIII | AAGCTT | 3035 |
| HpaI | GTTAAC | 501,4171 |
| KasI | GGCGCC | 4041,6816 |
| KpnI | GGTACC | 680 |
| MluI | ACGCGT | 230,8212 |
| NcoI | CCATGG | 1209,4257 |
| NdeI | CATATG | 7006 |
| NheI | GCTAGC | 239,505 |

| | | |
|-------|----------|--|
| NotI | GCGGCCGC | 7032 |
| NruI | TCGCGA | 7390 |
| PacI | TTAATTA | 7015 |
| PstI | CTGCAG | 1062,2093,2574,2921,3156,3255,4081,6084,6537 |
| PvuI | CGATCG | 692,7733 |
| PvuII | CAGCTG | 174,1495,1535,3524,4082,5284,5467,6538,6598,7146 |
| SacI | GAGCTC | 890,2385,2907,3054,4443 |
| SacII | CCGCGG | 780 |
| SalI | GTCGAC | 3876 |
| SanDI | GGGWCCC | 3841,5480 |
| SpeI | ACTAGT | 670 |
| XbaI | TCTAGA | 877 |
| XhoI | CTCGAG | 4094 |
| XmaI | CCCGGG | 4542,5114 |

No Cuts: AscI, MfeI, SfiI, SnaBI, SphI

Sequence

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1 CTTTCTGCG TTATCCCCTG ATTCTGTGGA TAACCGTATT ACCGCCTTTG AGTGAGCTGA TACCGCTCGC
71 CGCAGCCGAA CGAGTCGAGC CAGCGAGTCA GTGAGCGGAG AAGCGGAAGA GCGCCCAATA CGCAAACCCG
141 CTCTCCCCGC GCGTTGGCCG ATTCATTAAT GCAGCTGGCA CGACAGGTTT CCCGACTGGA AAGCGGGCAG
211 TGAGCGCAAC GCAATTAATA CGCGTACCGC TGACCAGGAA GAGTTTGTAG AAACGCAAAA AGGCCATCCG
281 TCAGGATGGC CTTCTGCTTA GTTTGATGCC TGGCAGTTTA TGGCGGGCGT CCGTCCCGCC ACCCTCCGGG
351 CCGTTGCTTC ACAACGTTCA AATCCGCTCC CGGCGGATTT GTCCTACTCA GGAGAGCGTT CACCGACAAA
421 CAACAGATAA AACGAAAGGC CCAGTCTTCC GACTGAGCCT TTCGTTTTAT TTGATGCCTG GCAGTTCCTT
491 ACTCTCGCGT TAACGCTAGC ATGGATGTTT TCCCAGTAC GACGTTGTAA AACGACGGCC AGTCTTAAGC
561 TCGGGCCCCA AATAATGATT TTATTTTGAC TGATAGTGAC CTGTTGTTG CAACACATTG ATGAGCAATG
631 CTTTTTTTATA ATGCCAACTT TGTACAAAAA AGCAGGCTCA CTAGTGGTAC CGTTTTAAACG ATCGGCCGCC
701 ACCATGCTGC GAGTCTTCAT CCTTTTTGCG GAGAATGTCC ACACCCCGGA CTCCGACATC AGCGATGCCT
771 ACTGCTCCGC GGTGTTTGCA GGGGTAAAGA AGAGAACCAA AGTCATCAAG AACAGTGTGA ACCCCGTGTG
841 GAATGAGGGC TTTGAGTGGG ACCTCAAAGG TATTCTCTA GATCAGAGCT CAGAACTTCT CGTGGTGGTC
911 AAGGACCATG AGACGATGGG AAGAAACAGG TTCCTGGGGG AAGCCAAGAT CCCACTCCAG GAGGTCCTTG
981 CCACCCCCAG CCTCTGTCC AGCTTCAACG CACCTGTGCT AGACGCCAAG CAGCAACCCA CGGGGGCCTC
1051 TCTGGTCTTG CAGGTGTCTT ACACGCCACC CCCAGGAGT GTGCCCTGT TCCCACCACC TGCTTCTCTA
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1471 AGTCCAGGTG ATAGAGGGG GTCAGCTGCC TGGGGTGAAT ATTAAGCCTG TGGTCAAGGT CACAGCTGCT
1541 GGGCAGACCA AGCGAACTCG GATCCAGAAG GGAAATAGCC CACTCTTCAA TGAGACTCTT TTCCTCAACG
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1891 CATTGAAGGC AACCTGCTTA GGCCACCCG TGTGGCCTT CGAGGAGCCC ACTTCTGCCT GAAGCTCTT
1961 CGGGCTGAGG ACTTACCACA GATGGACGAT GCTGTGATGG ACAACGTCAA GCAGATCTTT GGCTTTGACA
2031 GCAACAAGAA GAACTTGGTG GATCCCTTTG TGAAGTCAG CTTTGCTGGG AAAATGCTCT GCAGCAAGAT
2101 CCTGGAGAAG ACAGCTAACC CTCAGTGAA CCAGAACATC ACCTTGCTG CGATGTTTCC CTCTATGTGT
2171 GAAAAAATAA GGATTCGTGT CATGGACTGG GACCCTCA CTCACAATGA CACTGTGGCC ACCACCTACT
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7631 AGAAGAATAT CCTGATTGAG GTGAAAATAT TGTGATGCG CTGGCAGTGT TCCTGCGCCG GTTGCATTCG
7701 ATTCTGTTT GTAATTGTCC TTTTAAACAGC GATCGCTPAT TTCGTCTCGC TCAGGCGCAA TCACGAATGA
7771 ATAACGGTTT GGTGATGCG AGTGATTTTG ATGACGAGCG TAATGGCTGG CCTGTTGAA CAGTCTGGAA
7841 AGAAATGCAT AAACTTTTGC CATTCTCACC GGATTCAGTC GTCACTCATG GTGATTTCTC ACTTGATAAC
7911 CTTATTTTTG ACGAGGGGAA ATTAATAGGT TGTATTGATG TTGGACGAGT CGGAATCGCA GACCGATACC
7981 AGGATCTTGC CATCCTATGG AACTGCCTCG GTGAGTTTTT TCCTTCATTA CAGAAAACGGC TTTTTCAAAA
8051 ATATGGTATT GATAATCCTG ATATGAATAA ATTCGAGTTT CATTTGATGC TCGATGAGTT TTTCTAATCA
8121 GAATTGGTTA ATTGGTTGTA ACACTGGCAG AGCATTACGC TGACTTGACG GGACGGCGCA AGCTCATGAC
8191 CAAAATCCCT TAACGTGAGT TACGCGTCGT TCCACTGAGC GTCAGACCCC GTAGAAAAGA TCAAAGGATC
8261 TTCTTGAGAT CCTTTTTTTC TGCGCGTAAT CTGCTGCTTG CAAACAAAAA AACCACCGCT ACCAGCGGTG
8331 GTTTGTTTGC CGGATCAAGA GCTACCAACT CTTTTTCCGA AGGTAAGTGG CTTTCAGCAGA GCGCAGATAC
8401 CAAATACTGT TCTTCTAGTG TAGCCGTAGT TAGGCCACCA CTTCAAGAAC TCTGTAGCAC CGCCTACATA
8471 CCTCGCTCTG CTAATCCTGT TACCAGTGGC TGCTGCCAGT GGCGATAAGT CGTGTCTTAC CGGGTTGGAC
8541 TCAAGACGAT AGTTACCGGA TAAGGCGCAG CGGTCGGGCT GAACGGGGGG TTCGTGCACA CAGCCCAGCT
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8751 GGGGGAAACG CCTGGTATCT TTATAGTCTT GTCGGGTTTC GCCACCTCTG ACTTGAGCGT CGATTTTTGT
8821 GATGCTCGTC AGGGGGGCGG AGCCTATGGA AAAACGCCAG CAACGCGGCC TTTTACGGT TCCTGGCCTT
8891 TTGCTGGCCT TTTGCTCACA TGTT

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