

PRIMERS

Subdivision of *IL1RAPL1* intron 5

1st subdivision

Int5QtF	TGACATGCATAATACAGAATCAGAG
Int5QtR	ATCAGCACCTTGATCTTGGACTTTC
Int5HfF	GGAAGTAATTTGACAGTCAATCATC
Int5HfR	TCCTTAGGTGCTATTGTATAATGTG
Int53qtF	GTATCAGTGTCAGCTGTTGGTCATC
Int53qtR	CGCTTTGAAGCAACTCTCATCTTAC

2nd subdivision

Int5STSaF	GATTCAGCCATGTGAATTAAGATGG
Int5STSaR	GCACTTCACATGAAAGTTTATCTTC
Int5STSBfF	CACACCTGCTCCTTTTACAGACTTC
Int5STSBfR	TTCTGACCAGGGACTAGTGATGATC
Int5STScF	GAAGTCAAGCATTATAGACATTGCC
Int5STScR	CTTTGATCTCTTAACTCCTTGGTCG

3rd subdivision

Int5STS1F	GAAAGGAGTACAGGTTTCAATGCAG
Int5STS1R	AGAATCAGATAAGTTAGCCATCTTC
Int5STS2F	GCAAACATCTTGCTTTTTGGAAATC
Int5STS2R	ACAGCATTAGTAGATGTTCAACAC
Int5STS3F	AGAGGTTGTTGAGGTAAACTGTAAC
Int5STS3R	CATGACCTCACTTATATGTGGAATC

Other primers

5'MiddF	CCTTGGACCATGTCAAATCACATTG
Int5STSdR	TAGTTCACACCTGTAATCCCAGCAC

Subdivision of dystrophin intron 51

1st subdivision

Int51QtF	TGTCACAAACCCAGCAGAAAGTGGC
Int51QtR	AAGATGACTGTCACCCAACCTCGTG

Int51HfF TGATACTATGTGAAGTGGCAAGCAG
Int51HfR AAGATCCTATTCTAGACATCTGGCC

Int513QtF ACACAGGAATAAGAAAGTGGAAACGG
Int513QtR CTTAGATGTTAGTTTGATCTTCTGG

2nd subdivision

Int51STSaF CTAGCCTTTCCTGCAGCCTTCACTC
Int51STSaR CAACCAGTCATACCAGGCTCCTGTG

Int51STSBf AGCTCGGTTAGGAACACAACATTAC
Int51STSBr AGACATACTTGTCTATTCCCTAGGC

Int51STScF CATCCATGAGGCACATTCCCTACTG
Int51STScR CCCTATTCAGAGGAATCTATACAGC

Int51STSDf CTGGGGGCAATGTCATGGTCCACTC
Int51STSDr CTGTGACTCTGGTTCTCATGCCACC

Subdivision of dystrophin intron 52

1st subdivision

Int52QtF AAAGTGCTGGGATTACAGGCGTGAG
Int52QtR TTCTACTTCTCCTAGGGACAACAGG

Int52HfF TTGTGTTCCGCTGAACTAGAGAGAC
Int52HfR CTCTTGA ACTCTGTATCCATAGAGC

Int523QtF GGAGCTTCAATACATCTGGCAGCAG
Int523QtR AGTTCTCTCAGCATTCACCACAAGC

2nd subdivision

Int52STSaF TGGCTCACTCCTGTAATTCCAGCAC
Int52STSaR GTTGCCCAGGTTAGAGTGCAATGGC

Int52STSBf CATCTGTAAGTTACAAATGCCATAG
Int52STSBr GTAATTCCAGTGATGACTCATGGTC

Int52STScF ACTCGATTGATTGATTGATTGATTG
Int52STScR CTTTCTAGAAGTTCTGCATATGAGC

3rd subdivision

Int52STSGf TACACAGCATAAGTCCTGATTATTC
Int52STSGr TTGCTTTACCATAACCACTGTCCTAC

Int52STShF ATCCAATCTCTTGTCAGCCATGTCC
Int52STShR TGGAATAGATGAAATAGGTGAGAGG

Subdivision of dystrophin-*FTHL17* intergenic region

1st subdivision

QuartF TCTTGCTTACATCACAGTCTGAGGC
QuartR TCAGCTTTTCTCCTGCACTTCCCAG

HalfF CCTCAACAGTTAATACATGAGCAAC
HalfR AAGTACAAGGCTGCATCTCGATGTC

3quartF TATGGAGTGAAGAACCTCTGTTATG
3quartR CAATCTAACTGCAAGGAAGGCTAGG

2nd subdivision

BkPtQtF GAACAACTGGTCATCACCTTCAATG
BkPtQtR TGAATGGCTGAACTAGGGAGGAGAC

BkPtHfF GTGGCTGGGGTGTATCTACAAGCTC
BkPtHfR CTA ACTGGCAAGCTCCAGCTCTTAG

BkPt3qF TGTAAGGAGGCATGATGTCTACAAC
BkPt3qR AAAGGTTACCTACAGTAACTTGAGG

Other primers

BkPt1F TTCATGACCATCGTACGCTGCAGGG
3'MiddR GTTAAAAGGGGATCTGTGTGTCCAG

Amplification of hybrid transcript

Outer primers

IL1RAPL14F CCTGCTCTGCACCTGGATTG
DMD8b CTCAGGAGGCAGCTCTCTGG

Inner primers

IL1RAPL16F CTGAATACGCATTGTGATGACCAGC
DMD8d CTCCTGGTAGAGTTTCTCTAG

3'RACE

ex46F: TAGTATCCCCTTGAACCTG
ex49F: CCAGCCACTCAGCCAGTGAAG