

DR3v2	NM_003790.2	CGGGCCCTGC	GGGCGCGGGG	CTGAAGGCGG	AACCACGACG	GGCAGAGAGC	ACGGAGCCCG	GAAGCCCCTG	70
	DR3v2Δ6	-----	-----	-----	-----	-----GAGC	ACGGAGCCCG	GAAGCCCCTG	24
DR3v2	NM_003790.2	GGCGCCCGTC	GGAGGGCTAT	GGAGCAGCGG	CCGCGGGGCT	GCGCGGCGGT	GGCGGCGCGG	CTCTCTCTGG	140
	DR3v2Δ6	GGCGCCCGTC	GGAGGGCTAT	GGAGCAGCGG	CCGCGGGGCT	GCGCGGCGGT	GGCGGCGCGG	CTCTCTCTGG	94
DR3v2	NM_003790.2	TGCTGCTGGG	GGCCCCGGCC	CAGGGCGGCA	CTCGTAGCCC	CAGGTGTGAC	TGTGCCGGTG	ACTTCCACAA	210
	DR3v2Δ6	TGCTGCTGGG	GGCCCCGGCC	CAGGGCGGCA	CTCGTAGCCC	CAGGTGTGAC	TGTGCCGGTG	ACTTCCACAA	164
DR3v2	NM_003790.2	GAAGATTGGT	CTGTTTTGTT	GCAGAGGCTG	CCCAGCGGGG	CACTACCTGA	AGGCCCTTTG	CACGGAGCCC	280
	DR3v2Δ6	GAAGATTGGT	CTGTTTTGTT	GCAGAGGCTG	CCCAGCGGGG	CACTACCTGA	AGGCCCTTTG	CACGGAGCCC	234
DR3v2	NM_003790.2	TGCGGCAACT	CCACCTGCCT	TGTGTGTCCC	CAAGACACCT	TCTTGGCCTG	GGAGAACCAC	CATAAATTCTG	350
	DR3v2Δ6	TGCGGCAACT	CCACCTGCCT	TGTGTGTCCC	CAAGACACCT	TCTTGGCCTG	GGAGAACCAC	CATAAATTCTG	304
DR3v2	NM_003790.2	AATGTGCCCC	CTGCCAGGCC	TGTGATGAGC	AGGCCTCCCA	GGTGGCGCTG	GAGAACTGTT	CAGCAGTGCC	420
	DR3v2Δ6	AATGTGCCCC	CTGCCAGGCC	TGTGATGAGC	AGGCCTCCCA	GGTGGCGCTG	GAGAACTGTT	CAGCAGTGCC	374
DR3v2	NM_003790.2	CGACACCCGC	TGTGGCTGTA	AGCCAGGCTG	GTTTGTGGAG	TGCCAGGTCA	GCCAAATGTT	CAGCAGTTCA	490
	DR3v2Δ6	CGACACCCGC	TGTGGCTGTA	AGCCAGGCTG	GTTTGTGGAG	TGCCAGGTCA	GCCAAATGTT	CAGCAGTTCA	444
DR3v2	NM_003790.2	CCCTTCTACT	GCCAACCATG	CCTAGACTGC	GGGGCCCTGC	ACCGCCACAC	ACGGCTACTC	TGTTCCCGCA	560
	DR3v2Δ6	CCCTTCTACT	GCCAACCATG	CCTAGACTGC	GGGGCCCTGC	ACCGCCACAC	ACGGCTACTC	TGTTCCCGCA	514
DR3v2	NM_003790.2	GAGATACTGA	CTGTGGGACC	TGCCTGCCTG	GCTTCTATGA	ACATGGCGAT	GGCTGCGTGT	CCTGCCCCAC	630
	DR3v2Δ6	GAGATACTGA	CTGTGGGACC	TGCCTGCCTG	GCTTCTATGA	ACATGGCGAT	GGCTGCGTGT	CCTGCCCCAC	584
DR3v2	NM_003790.2	GAGCACCCCTG	GGGAGCTGTC	CAGAGCGCTG	TGCCGCTGTC	TGTGGCTGGA	GGCAGATGTT	CTGGGTCCAG	700
	DR3v2Δ6	-----	-----	-----	-----	-----	-----TGTT	CTGGGTCCAG	598
DR3v2	NM_003790.2	GTGCTCCTGG	CTGGCCTTGT	GGTCCCCCTC	CTGCTTGGGG	CCACCTGAC	CTACACATAC	CGCCACTGCT	770
	DR3v2Δ6	GTGCTCCTGG	CTGGCCTTGT	GGTCCCCCTC	CTGCTTGGGG	CCACCTGAC	CTACACATAC	CGCCACTGCT	668
DR3v2	NM_003790.2	GGCCTCACAA	GCCCCTGGTT	ACTGCAGATG	AAGCTGGGAT	GGAGGCTCTG	ACCCACCCAC	CGGCCACCCA	840
	DR3v2Δ6	GGCCTCACAA	GCCCCTGGTT	ACTGCAGATG	AAGCTGGGAT	GGAGGCTCTG	ACCCACCCAC	CGGCCACCCA	738
DR3v2	NM_003790.2	TCTGTACACC	TTGGACAGCG	CCCACACCCT	TCTAGCACCT	CCTGACAGCA	GTGAGAAGAT	CTGCACCCTC	910
	DR3v2Δ6	TCTGTACACC	TTGGACAGCG	CCCACACCCT	TCTAGCACCC	CCTGACAGCA	GTGAGAAGAT	CTGCACCCTC	808
DR3v2	NM_003790.2	CAGTTGGTGG	GTAACAGCTG	GACCCCTGGC	TACCCCGAGA	CCCAGGAGGC	GCTCTGCCCG	CAGGTGACAT	980
	DR3v2Δ6	CAGTTGGTGG	GTAACAGCTG	GACCCCTGGC	TACCCCGAGA	CCCAGGAGGC	GCTCTGCCCG	CAGGTGACAT	878
DR3v2	NM_003790.2	GGTCCTGGGA	CCAGTTGCCC	AGCAGAGCTC	TTGGCCCCGC	TGCTGCGCCC	ACACTCTCGC	CAGAGTCCCC	1050
	DR3v2Δ6	GGTCCTGGGA	CCAGTTGCCC	AGCAGAGCTC	TTGGCCCCGC	TGCTGCGCCC	ACACTCTCGC	CAGAGTCCCC	948
DR3v2	NM_003790.2	AGCCGGCTCG	CCAGCCATGA	TGCTGCAGCC	GGGCCCGCAG	CTCTACGACG	TGATGGACGC	GGTCCCAGCG	1120
	DR3v2Δ6	AGCCGGCTCG	CCAGCCATGA	TGCTGCAGCC	GGGCCCGCAG	CTCTACGACG	TGATGGACGC	GGTCCCAGCG	1018
DR3v2	NM_003790.2	CGGCGCTGGA	AGGAGTTCGT	GCGCACGCTG	GGGCTGCGCG	AGGCAGAGAT	CGAAGCCGTG	GAGGTGGAGA	1190
	DR3v2Δ6	CGGCGCTGGA	AGGAGTTCGT	GCGCACGCTG	GGGCTGCGCG	AGGCAGAGAT	CGAAGCCGTG	GAGGTGGAGA	1088
DR3v2	NM_003790.2	TCGGCCGCTT	CCGAGACCAG	CAGTACGAGA	TGCTCAAGCG	CTGGCCGACG	CAGCAGCCCG	CGGGCCTCGG	1260
	DR3v2Δ6	TCGGCCGCTT	CCGAGACCAG	CAGTACGAGA	TGCTCAAGCG	CTGGCCGACG	CAGCAGCCCG	CGGGCCTCGG	1158
DR3v2	NM_003790.2	AGCCGTTTAC	GCGGCCCTGG	AGCGCATGGG	GCTGGACGGC	TGCGTGAAG	ACTTGGCCAG	CCGCCTGCAG	1330
	DR3v2Δ6	AGCCGTTTAC	GCGGCCCTGG	AGCGCATGGG	GCTGGACGGC	TGCGTGAAG	ACTTGGCCAG	CCGCCTGCAG	1228
DR3v2	NM_003790.2	CGCGGCCGT	GACACGGCGC	CCACTTGCCA	CCTAGGCGCT	CTGGTGGCCC	TTGCAAGAAG	CCTAAGTACG	1400
	DR3v2Δ6	CGCGGCCGT	GACACGGCGC	CCACTTGCCA	CCTAGGCGCT	CTGGTGGCCC	TTGCAAGAAG	CCTAAGTACG	1298
DR3v2	NM_003790.2	GTTACTTATG	CGTGTAGACA	TTTTATGTCA	CTTATTAAGC	CGCTGGCAGG	GCCCTGCGTA	GCAGCACCAG	1470
	DR3v2Δ6	GTTACTTATG	CGTGTAGACA	TTTTATGTCA	CTTATTAAGC	CGCTGGCAGG	GCCCTGCGTA	GCAGCACCAG	1368
DR3v2	NM_003790.2	CCGGCCCCAC	CCCTGCTCGC	CCCTATCGCT	CCAGCCAAGG	CGAAGAAGCA	CGAACGAATG	TCGAGAGGGG	1540
	DR3v2Δ6	CCGGCTCCAC	CCCTGCTCGC	CCCTATCGCT	CCAGCCAAGG	CGAAGAAGCA	CGAAC-----	-----	1423
DR3v2	NM_003790.2	GTGAAGACAT	TTCTCAACTT	CTCGCCGGA	GTTTGGCTGA	GATCGCGGTA	TAAATCTGT	GAAAGAAAAC	1610
	DR3v2Δ6	-----	-----	-----	-----	-----	-----	-----	1423
DR3v2	NM_003790.2	AAAAAAAAC	AAAAAAAAAA	AAAAAAAAAA	1638				
	DR3v2Δ6	-----	-----	-----	1423				