

CR1 Wild-type

GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACTCAAGTGGGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	WT
CR2 genome and transgene targeting deletions (19)	
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGCA- CAAAGTGGGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-2
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGA-- AGTGGGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-4
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGAA-- -AGTGGGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-8
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGA-- -CTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-10
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGA-- -ACTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-16
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACT----- CAGTCAGAGTTGTGCACATGGCTTAGTTT	-17
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACT----- ACCAGTCAGAGTTGTGCACATGGCTTAGTTT	-23
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACT----- ACCAGTCAGAGTTGTGCACATGGCTTAGTTT	-23
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAG----- TTCTTAGTTT	-68
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAG----- TTT	-108
CR1 genome targeting deletions (15)	x5
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACTG- GTGGGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-2
GCTACCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACTC- GGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT	-4
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACT----- GGTGACCCAGTCAGAGTTGTGCACATGGCTTAGTTT	-10
GCTCCCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGACACGGACT----- CCAGTCAGAGTTGTGCACATGGCTTAGTTT	-23
GCTACCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGAC----- CCAGTCAGAGTTGTGCACATGGCTTAGTTT	-23
GCTACCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTGAC----- CCAGTCAGAGTTGTGCACATGGCTTAGTTT	-40
GCTACCGAGGAGCAAGCTCAGTTACACCGATCCACTGGGAGCAGGAATATCTGTGGGCTTGTG----- CCATGGCTTAGTTT	-40
GCT----- GGTGACCCAGTCAGAGTTGTGCACATGGCTTAGTTT	-84
TTGGTTTTGGGCCACATGCTGGTACATCCTCATCCTGATAAACTGCAAAGGCTGAAGAGCATGACTGACATCACCTGGCTACCTGCAACCTGGCATCTGACCTGTTTTCCA-----	
-----	-839
-----	-----
-----	GGCTGGTGAACAGTCAGAGTTGTGCACATGGCTTAGTTT

CR2 Wild-type

GGGGCTGTCTGGGCTGCTTG- TCATGGTACCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	WT
CR2N genome and transgene targeting deletions (8)	
GGGGCTGTCTGGGCTGCT----- TGTCATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-5
GGGGCTGTCTGGGCT----- GCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-17
GGGGCTGTCTGGGCT----- ACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-20
GG----- CATGGTCATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-21
GGGGCTGTCTGGGCT----- GGGAAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-24
CR2 genome and transgene targeting insertion (1)	
GGGGCTGTCTGGGCTGCTTG- TCATGGTACCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	1
CR2 genome targeting deletions (16)	
GGGGCTGTCTGGGCTGCT----- CATGGTCATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-5
GGGGCTGTCTGGGCTGCT----- TGTCATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-5
GGGGCTGTCTGGGCTGCT----- ATGGTCATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-8
GGGGCTGTCTGGGCTGCT----- TCTAGTCATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-10
GGGGCTGTCTGGGCTGCT----- CACTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-11
GGGGCTGTCTGGGCTGCT----- TCACTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-16
GGGGCTGTCTGGGCTGCT----- CTAGGGAAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-17
GGGGCTGTCTGGGCTGCT----- CTGGAAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-20
GGGGCTGTCTGGGCTGCT----- ATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-21
GGGGCTGTCTGGGCTGCT----- TACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-23
GGGGCTGTCTGGGCTGCT----- AAACCTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-30
GGGGCTGTCTGGGCTGCT----- CTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	-42
GGGGCTGTCTGGGCTGCTGGT----- CCATCATG	-88
ACTCACTGGTGTACATTTGGTTTGTGGCAACATGCTGGTACATCCTCATCCTGATAAACTGCAAAGGCTGAAGAGCGTGAACATCACCTGCTAAC-----	
-----	-426
-----	-----
-----	CTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG
CR2 genome targeting insertion (1)	
GGGGCTGTCTGGGCTGCTTG- TTCACTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	1
CR2 genome targeting deletion and insertion (2)	
GGGGCTGTCTGGGCTGCTTG- A-CATGG- CATCTGCTACTCGGAATCCTAAAAACTCTGCTTCGGTGTGAAATGAGAAGAGAGGACAGGGCTGTGAGGCTTATCTCACCATCATG	(+1)(-1)
GGGGCTGTCTGGGCTGCTTG----- CCATCATG	(+2)(-2)

CR3 Wild-type

GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC-- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	WT
CR3 genome and transgene targeting deletions (13)	
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC-- TCGTAGGCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-2
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC----- GGAGGCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-4
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC----- GGAGGCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-7
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT----- CGGGAGCCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-8
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGGT----- AGCCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-8
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC----- GCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-19
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC----- CTGGAAAATAGAACAGCATTTGCGAAAGCGCTTGGCAATGTGC	-25
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC----- TGCGAGGCTCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-23
CR3 genome and transgene targeting insertion (6)	
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	1
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	1
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	1
CR3 genome targeting deletions (11)	
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC----- GGAGGCTCTTGTGCGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-4
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCAAGGCTA----- GTAGCTCTTGTGCGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-4
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT----- CGGGAGCCTCTTGTGCGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-8
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT----- GGAGGCTCTTGTGCGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-10
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT----- CTGAAAATAGAACAGCATTTGCGAAAGCGCTTGGCAATGTGC	-19
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT----- CTCTGCGAAAATAGAACAGCATTTGCGAAAGCGCTTGGCAATGTGC	-17
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT----- TAGCCTCTTGTGCGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	-27
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCT-----	-199
----- ACTGCAATTATTCAGGCAAAAGATTCTGGAAAGGTTTGTAGGAGAAGGACAATGTTGTAGGGAGCCAGAAGAGAAAATAAACACATGATGGTGAAGATAAGCCTCACAG	
CR3 genome targeting insertion (8)	
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	1
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	1
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	1
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAACT-- GAGCTTGCTC- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	2
CR3 genome targeting deletion and insertion (1)	
GTCACAAAGCCACAGATATTTCTGCTCCCCAGTGGATCGGGTGTAAAC- AAGAG----- GCTCGGGAGCCTTGTGCTGGAAAATAGAACAGCATTTGCGAAAGCGTTGGCAATGTGC	(-1)/2(-7)