

CD8 KIAA0408 MUT - specific CDR3 Vβ				
CDR3 nucleotide sequence	Vβ	TCR reads	Primary	Metastasis
TGTGCCAGTAGTATAGGTGGCAGCTCTACAATGAGCAGTTCTTC	TRBV19*01	15		1
TGCAGTGCTAGCAGACTAGCAGGGGACAATGAGCAGTTCTTC	TRBV20-1*01	1		1
TGTGCCAGCAGTTACCAGGGGCCACTTATGGCTACACCTTC	TRBV6-6*01	1		1
TGTGCCAGCAGCGTGGGACAGGGGTCCACGAGCAGTACTTC	TRBV9*01	1		1
TGTGCCAGCAGCCAGGGGTGGGCTACACCTTC	TRBV27*01	1	1	
TGTGCCAGCAGTCCGTGGGCGGGAGCAGATACGCAGTATTTT	TRBV12-3*01,TRBV12-4*01	1	1	
TGCGCCAGCAGCCAAAGATCGAAGGGGGGAACTACGAGCAGTACTTC	TRBV4-3*01	1	1	
TGCGCCAGCAGCGGTGGGCGGGTGGAGACCCAGTACTTC	TRBV5-1*01	1	7	2
TGTGCCAGCAGTTATTCGGGACTAGCGGGATTGCCGATGAGCAGTTCTTC	TRBV6-6*01	27		
TGTGCCAGCAGCCAAAGATTGGGTAGAGCAGTTCTTC	TRBV14*01	13		
TGTGCCAGCTACTAGACCTGAACACTGAAGCTTTCTTT	TRBV18*01	5		
TGTGCCAGCAGCTTCCGTGGGACAGGTCAAAGCGAGCAGTACTTC	TRBV13*01	5		
TGTGCCAGCAGCGCGGGACAGGGCCTACGAGCAGTACTTC	TRBV2*01	4		
TGTGCCAGTAGTTGGGACCGATTAACTGAAGCTTTCTTT	TRBV19*01	4		
TGCAGTGCTAGATCTCTGGGAGCAATAGTGGAGCAGTACTTC	TRBV20-1*01	4		
TGTGCCAGCAGCCTTGTGTCGAGCAGTACTTC	TRBV11-2*01	3		
TGTGCCAGCAGCTTTCGGGGACAGGGATGTCTACGAGCAGTACTTC	TRBV5-6*01	2		
TGCAGTGCTAGAGATTGAGACTAGCGATCTCTACGAGCAGTACTTC	TRBV20-1*01	2		
TGTGCCAGCAGCCCCGTTAGCGAAAATGAGCAGTTCTTC	TRBV7-2*01	2		
TGTGCCAGCAGCGGGAGCGGGAGTGGCGGGGAGCTGTTTTTT	TRBV7-8*01	2		
TGTGCCAGCAGCTTAGCGGGAGGGCCCTCCGCTACAATGAGCAGTTCTTC	TRBV7-2*01	2		
TGCAGCGTTGAACTAGCGGGAATACCGGGGAGCTGTTTTTT	TRBV29-1*01	2		
TGCAGTGCCCATCGACTAGAGCAGTTCTTC	TRBV20-1*01	2		
TGCAGTGGCGACAGTGGGACCGGGACAGGGGTTCAACCCTACGAGCAGTACTTC	TRBV20-1*01	2		
TGCAGTGCTAGAGATGGAAACAGGGGGCGGTGAGCAGTTCTTC	TRBV20-1*01	2		
TGCAGTGCTAGAGCAGTCACTCTACGAGCAGTACTTC	TRBV20-1*01	2		
TGTGCCAGCAGCCAAAGGGGAACTGAAGCTTTCTTT	TRBV16*01	2		
TGTGCCAGCAGTTATTCGGGACTAGCGGGATTGCCGATGAGCAGTTCTTC	TRBV6-9*01	2		
TGTGCCAGCAGTTGGCTAGTTATACAGATACGCAGTATTTT	TRBV28*01	2		
TGTGCCAGCGCAGGAGCTCTACAATGAGCAGTTCTTC	TRBV7-2*01	2		
TGTGCCAGCAGCCAAACCCCAAGGGAGGGACCGGGGAGCTGTTTTTT	TRBV6-5*01	2		
TGTGCCAGCAGCAGATAGCGGGAGAAAATCTACCGGGGAGCTGTTTTTT	TRBV2*01	2		
TGTGCCAGCAGCCCCGAGCAGGGGGAGGCCAGTACTTC	TRBV3-1*01	2		
TGCAGCGTTGAATTAGTGGGAGCTCTACGAGCAGTACTTC	TRBV29-1*01	2		
TGTGCCAGCAGCCGTTAGCGGTCTTGGCAGCCCCAGCATTTC	TRBV7-3*01	2		
TGTGCCAGCAGTTACGAAGCCCGGCAATCGATGAGCTGTTTTTT	TRBV6-5*01	2		
TGCGCCAGCAGCTTGGCAGGTCAACACCGGGGAGCTGTTTTTT	TRBV5-1*01	1		
TGCAGCACTGGGGGGGGAACACCGGGGAGCTGTTTTTT	TRBV29-1*01	1		
TGTGCCAGCAGTACGGTACGGGGGAGCAGTACTTC	TRBV3-1*01	1		
TGTGCCAGCAGTGATTATGATACCAAGCAATCAGCCCCAGCATTTC	TRBV6-1*01	1		
TGCAGTGCTAGAAAGGGGGCCCAAAAACATTGACTACTTC	TRBV20-1*01	1		
TGCGCCAGCAGCATAGGGTTAACCGGGGAGCTGTTTTTT	TRBV5-1*01	1		
TGCGCCAGCAGCCATGAGCTAGCCCCAGATACGCAGTATTTT	TRBV5-1*01	1		
TGTGCCAGTAGTATTTGGAGCGGGATTACAATGAGCAGTTCTTC	TRBV19*01	1		
TGTGCCAGCAGCCAAAGCCGGGACAGGATCCACGAGCAGTACTTC	TRBV3-1*01	1		
TGCGCCAGCAGCCAAAGATGCCAGTACCTCTACAATGAGCAGTTCTTC	TRBV4-3*01	1		
TGCGCCAGCAGCCCCCATGGGGAGGGCGGAAAATGAGCAGTTCTTC	TRBV4-1*01	1		
TGCGCCAGCAGCTCGGACGCAACTCTACGAGCAGTACTTC	TRBV5-1*01	1		
TGCGCCGGGAGCGATCCGAACACCGGGGAGCTGTTTTTT	TRBV5-1*01	1		
TGTGCCAGCAGCCACCGCTGGGTAGTAGCTCTACAATGAGCAGTTCTTC	TRBV7-9*01	1		
TGTGCCAGCAGCGTAGCTACCGGGGAGCTGTTTTTT	TRBV9*01	1		
TGTGCCAGCAGCTTATCGGGGGGCCCCAAGAGACCCAGTACTTC	TRBV11-3*01	1		
TGCAGCGTAGCGGGAGTGGCTACGAGCAGTACTTC	TRBV29-1*01	1		
TGCAGCGTCTTAAACAGGAACCTTAACATGGCTACACCTTC	TRBV29-1*01	1		
TGCAGCGTGAGGCTGGCCGGGACTAGCGGGAGGAGTGTAGCAGTTCTTC	TRBV29-1*01	1		
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TGCAGCGTTGAACGGGGGCTATGAGCAGTTCTTC	TRBV29-1*01	1		
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TGCAGCGTTGAGGATCGGGAACCTCTGGAACACCATATATTTT	TRBV29-1*01	1		
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TGCAGCGTTTCTTAGCGGGGGCCGTAGCACAGATACGCAGTATTTT	TRBV29-1*01	1		
TGCAGTGCCCTCCGACAGTGGCTACGAGCAGTACTTC	TRBV20-1*01	1		
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TGCAGTGCTAGAGCTGACTAGCGGCAAGAGACCCAGTACTTC	TRBV20-1*01	1		
TGCAGTGCTAGAGTGGCAGGGAGCTCTACGAGCAGTACTTC	TRBV20-1*01	1		
TGCAGTGCTAGTCTGGACTAGCTGACGAGCAGTACTTC	TRBV20-1*01	1		
TGCAGTGCTCCCGGACTAGCGGAGGGAGTTGACTACAATGAGCAGTTCTTC	TRBV20-1*01	1		

TGCAGTGCCTTATTAGATACGCAGTATTTT	TRBV20-1*01	1		
TGCGCCAGAGGAAACACCATATATTTT	TRBV10-1*01	1		
TGCGCCAGCACCGACAGGGGAAAGGGGAAAAACATTCACTTTC	TRBV5-1*01	1		
TGCGCCAGCAGCAGTACAGGGAAATGCCTACGAGCAGTACTTC	TRBV5-1*01	1		
TGCGCCAGCAGCCAAAGATGGGCATTGAACACTGAAGCTTTCTTT	TRBV4-1*01	1		
TGCGCCAGCAGCCAAGAGCCCAGCACAGATACGCAGTATTTT	TRBV4-1*01	1		
TGCGCCAGCAGCCAAGAGGGGACTAGCGCTACGAGCAGTACTTC	TRBV4-1*01	1		
TGCGCCAGCAGCCAAGATCGGGGACTCTGGGCTGGTGGGCTACACCTTC	TRBV4-3*01	1		
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TGCGCCAGCAGCCTGACAGGGGGCACCGGGGAGCTGTTTTTTT	TRBV4-1*01	1		
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TGCGCCAGCAGCTCAATCCGGGGCTCCACAGCAGTACTTC	TRBV4-3*01	1		
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TGTGCCACCAGCAGGACAGGGGAAACACCATATATTTT	TRBV15*01	1		
TGTGCCACCAGTGCAGAGGGTACAATGAGCAGTTCTTC	TRBV24-1*01	1		
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TGTGCCACCAGTGAATCCGACGGGAACTGAAGCTTTCTTT	TRBV24-1*01	1		
TGTGCCACCAGTGATTCGAGAACCAGGGAGCTGTTTTTTT	TRBV24-1*01	1		
TGTGCCACCAGTGATTTGGGGGAAAGCAGTACTTC	TRBV24-1*01	1		
TGTGCCACCAACTACTAGCACAGATACGCAGTATTTT	TRBV2*01	1		
TGTGCCAGCAAATCAGACGGGGGCCGGGAGCTGTTTTTTT	TRBV6-1*01	1		
TGTGCCACAATCAGGAGCTGGAGCTGAAGCTTTCTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCACCCAGACAGGGGGCGGTGAGCAGTTCTTC	TRBV28*01	1		
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TGTGCCAGCAGCCATCCGGGACTAGTTAATGAGCAGTTCTTC	TRBV7-8*01	1		
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TGTGCCAGCAGCCCCAGATTATACCGGGGAGCTGTTTTTTT	TRBV12-3*01,TRBV12-4*01	1		
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TGTGCCAGCAGCCGCTAGCGGGTGGAAATGAGCAGTTCTTC	TRBV7-9*01	1		
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TGTGCCAGCAGCCTCAGAAAGCGGGGACAGGGTCACAATGAGCAGTTCTTC	TRBV7-9*01	1		
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TGTGCCAGCCCTCAGAGGGGCCAGGTCCTACAATGAGCAGTTCTTC	TRBV2*01	1		
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TGTGCCAGTAGTCAGGGACTAGCGGTTTACGAGCAGTACTTC	TRBV19*01	1		
TGTGCCAGTAGTGCCGAAATAGCAATCAGCCCCAGCATTTT	TRBV19*01	1		
TGTGCCAGTCTACTAGCGGCAATGAGCAGTTCTTC	TRBV6-1*01	1		
TGTGCCAGTGGGACAGGGGCTACGAGCAGTACTTC	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCATCAGGGACAATATGGCTACACCTTC	TRBV10-3*01	1		
TGTGCCATCAGTGAGACAGGGACCCAGCAGCAGTACTTC	TRBV10-3*01	1		
TGTGCCATCAGTGAGACCATCAGGGGCGCTACAATGAGCAGTTCTTC	TRBV10-3*01	1		
TGTGCCAGTACCCCGGACTAGCAACGAGCAGTACTTC	TRBV30*01	1		
TGTGCCTGGAGTGTAGCTAGCGGATTCAGCTCTACAATGAGCAGTTCTTC	TRBV30*01	1		
TGTGCCTGGAGTGTGGTGAAGCTTTCTTT	TRBV30*01	1		
TGTGCCAGTAGTATCTTAGGAGGGGCGGGGAGCTGTTTTT	TRBV19*01	1		
TGCAGCGTCAAACCCGACCGCATCTAAACACTGAAGCTTTCTTT	TRBV29-1*01	1		
TGCAGCGTTGAAGGGGTGACTAGCGGGAGGAGCTCTACAATGAGCAGTTCTTC	TRBV29-1*01	1		
TGCAGTGTCTAGAGACCCGTTGGAGCACTCAATCAGCCCCAGCATTTT	TRBV20-1*01	1		
TGCAGTGTAGGGGGCAGGGTTGGGCACTGAAGCTTTCTTT	TRBV20-1*01	1		
TGCCAGCAGAGAAGGAAACGGGGAACTGAAGCTTTCTTT	TRBV10-1*01	1		
TGCCAGCAGCCGGAACAGGGGACTGGAACACCATATATTTT	TRBV5-1*01	1		
TGCCAGCAGCTTGGCCGGGGAGGGCAACAATGAGCAGTTCTTC	TRBV5-1*01	1		
TGCCAGCAGTGTGACAGACGAGCAGTACTTC	TRBV10-2*01	1		
TGTGCCACAGAGGACGATTAGCTGGAGCAGATACGCAGTATTTT	TRBV24-1*01	1		
TGTGCCACAGCAGAGACACAGCAATCAGCCCCAGCATTTT	TRBV15*01	1		
TGTGCCACAGTACGGGATGTTGGTGGCGGAGACCCAGTACTTC	TRBV24-1*01	1		
TGTGCCAGCAAGGGCGCAGTTCGGCTCGGAGACCCAGTACTTC	TRBV6-6*01	1		
TGTGCCAGCAATTTCGGCAGGCACAGATACGCAGTATTTT	TRBV28*01	1		
TGTGCCAGCAGCAAATTCAGTATGAGGGGTTACCAATCAGCCCCAGCATTTT	TRBV7-4*01	1		
TGTGCCAGCAGCAAGATCGGGGTTAGGACCCGGGAGCTGTTTTT	TRBV4-2*01	1		
TGTGCCAGCAGCCAAAGCGACAGGGAGCTCTACGAGCAGTACTTC	TRBV4-2*01	1		
TGTGCCAGCAGCCAAAGGACTAGCGGGAGAACCACAGATACGCAGTATTTT	TRBV14*01	1		
TGTGCCAGCAGCCAAAGGGGACATTTGAACATTGAGTACTTC	TRBV3-1*01	1		
TGTGCCAGCAGCCAGACTAGCGGGCATACTACGAGCAGTACTTC	TRBV3-1*01	1		
TGTGCCAGCAGCCGAGCGGACTAGCTCTACCCGGGAGCTGTTTTT	TRBV7-2*01	1		
TGTGCCAGCAGCCTGAACCATATGAACACTGAAGCTTTCTTT	TRBV7-9*01	1		
TGTGCCAGCAGCGCAGGACTTGGGGCTTTCTTT	TRBV9*01	1		
TGTGCCAGCAGCGGCCAACTATCATACGAGCAGTACTTC	TRBV9*01	1		
TGTGCCAGCAGCTTAGCGTTGCTAGCGGTTCTTTCAATGAGCAGTTCTTC	TRBV7-8*01	1		
TGTGCCAGCAGCTTCCAGCTAGCAGATACGAGCAGTACTTC	TRBV13*01	1		
TGTGCCAGCAGGAGGGGAGGGGAGGAGACCCAGTACTTC	TRBV7-9*01	1		
TGTGCCAGCAGGGGGGACGGTTCTACGAGCAGTACTTC	TRBV11-3*01	1		
TGTGCCAGCAGTCTCACTCTCGGGCAATGAGCAGTTCTTC	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGTTCCGCTAGCGTCCAAGAGACCCAGTACTTC	TRBV6-6*01	1		
TGTGCCAGCATCGGCGAGAATCGGGCGAGCAGTACTTC	TRBV2*01	1		
TGTGCCAGCGATATGACAGGTGGATACACTGAAGCTTTCTTT	TRBV2*01	1		
TGTGCCAGTAGACGGAACATGAACACTGAAGCTTTCTTT	TRBV19*01	1		
TGTGCCAGTAGTATGGGGATCAGGGAGACCCAGTACTTC	TRBV19*01	1		
TGTGCTAGTTTACGGGGAGGTCAGCCCCAGCATTTT	TRBV12-5*01	1		
TGCAGTATGGGACAGGGGGTTACGAGCAGTACTTC	TRBV20-1*01	1		
TGTGCCAGCAGCTTAGGACAGAACCCGGGAGCTGTTTTT	TRBV11-3*01	1		
TGCAGTGCCTCCCCGAGGCCACAGATACGCAGTATTTT	TRBV20-1*01	1		
TGCCAGCAGCGAAGCGACAGGGAACCCGGGAGCTGTTTTT	TRBV5-1*01	1		
TGCCAGCAGCTCGGAAGCGGGGGAATCAATGAGCAGTTCTTC	TRBV5-1*01	1		
TGCCAGCAGTGTAGGGCAGGTCGGTCAGCCCCAGCATTTT	TRBV10-2*01	1		
TGTGCCACCGGAACGAGCAATCAGCCCCAGCATTTT	TRBV6-6*01	1		
TGTGCCAGCAGCCAAAGATTGGGCTAGCGGGGAGCTGTTTTT	TRBV3-1*01	1		
TGTGCCAGCAGCGCTCCACAGATACGCAGTATTTT	TRBV9*01	1		
TGTGCCAGCAGCTTAGCGGGGGGATGAACACTGAAGCTTTCTTT	TRBV7-2*01	1		
TGTGCCTGGAGTGTGACAGCGAGCTACGAGCAGTACTTC	TRBV30*01	1		
TGTGCCAGCAGCAATAACGAGCAGTACTTC	TRBV2*01	1		
TGTGCCAGCAGCTCCGGGGGGCGGGAGATACGCAGTATTTT	TRBV7-3*01	1		
TGTGCCAGTAGCGCCAGGGGACGGCTCAATCAGCCCCAGCATTTT	TRBV19*01	1		
TGTGCCAGCACCGCTCTGAGCAGTTCTTC	TRBV11-2*01	1		

TGTGCCAGCAGCCAAGTCCCTGGACAGGGGAATAATTCACCCCTCCACTT	TRBV4-2*01	1		
TGTGCCAGCAGCAGCTACATTACAGGCCGGATCCTAGCGGGAGGGTCGACAATG	TRBV13*01	1		
TGTGCCAGCAGCTTGACAGGGATCCTACGAGCAGTACTTC	TRBV13*01	1		
TGCAGCGTGCCGGGATTAGCGGACTTC	TRBV29-1*01	1		
TGTGCCAGCAGCCAATCAATAAACAGGGTGACGAGCAGTACTTC	TRBV7-2*01	1		
TGTGCCAGCAGCCAGGCTGGGGCCAGGGCGCTCCACTTT	TRBV14*01	1		
TGTGCCAGCAGCTTATCAGTAATGAACACTGAAGCTTTCITT	TRBV7-2*01	1		
TGTGCCAGCAGAGGGTCTCAGGGAAAAGAGACCCAGTACTTC	TRBV15*01	1		
TGCGCCAGCAGCCAAGTGAAGGCCCTTTAAACACTGAAGCTTTCITT	TRBV4-3*01	1		
TGTGCCAGCAGCCAAGATGGAGGAAACACTGAAGCTTTCITT	TRBV4-2*01	1		
TGTGCCAGCAGTTCCTCAGGGCGGGACCCAGTACTTC	TRBV6-5*01	1		

CD4 KIAA0408 MUT - specific CDR3 Vβ				
CDR3 nucleotide sequence	Vβ	TCR reads	Primary	Metastasis
TGTGCCAGCTCACCCCCACGGGGCGAACCCGGGAGCTGTTTTTT	TRBV18*01	4		1
TGTGCCAGCAGTACCTGTCGGGGCACTAATGAAAACTGTTTTTT	TRBV6-5*01	2		1
TGTGCCAGCAGCAAGATGCTAGCGGGGTCTACAATGAGCAGTTCTTC	TRBV4-2*01	1		1
TGCGCCAGCAGCTTACAGGGGACCCGAAACCCGGGAGCTGTTTTTT	TRBV5-1*01	1		1
TGCGCCAGCAGCTTGGACACCTTGGCTCTACGAGCAGTACTTC	TRBV5-1*01	1		1
TGTGCCAGCAGCAAGATGGGCGGGGGCCCCACAGATACGCAGTATTT	TRBV3-1*01	1		1
TGTGCCAGCAGCAAGGGGATTCACCCCTCCACTT	TRBV7-6*01	1		1
TGTGCCAGCAGCCCGGACGGGACTAGCGGGAGTGAGACCCAGTACTTC	TRBV11-2*01	1		1
TGTGCCAGCAGCCCTAGTGGGAGAGAAAGACCCAGTACTTC	TRBV11-3*01	1		1
TGTGCCAGCAGCTAGGGGTAGCGGGAGTGAGACCCAGTACTTC	TRBV11-3*01	1		1
TGTGCCAGCAGTAGCAGGAGGGTCAACTAATGAAAACTGTTTTTT	TRBV27*01	1		1
TGTGCCAGCAGGTATCCGGGACAGCGAACACTGAAGCTTCTTT	TRBV27*01	1		1
TGTGCCCTCAAGATGGGGCCAGCACTGAAGCTTCTTT	TRBV10-3*01	1		1
TGCGCCAGCAGCTAGCAGGATGAACACTGAAGCTTCTTT	TRBV5-1*01	1	1	
TGTGCCAGCAGCTTAGACAGGGCAACCCCTGGCTACACCTTC	TRBV7-2*01	37		
TGTGCCAGCAGCACTCGACAGGATCGGGAAACCATATATTTT	TRBV7-2*01	34		
TGCGCCAGCAGCCCCCAGCCGACAGTCTGAGAGACCCAGTACTTC	TRBV5-1*01	26		
TGTGCCAGCAGCTTTTGGGAATGAGCAGTTCTTC	TRBV5-5*01	18		
TGTGCCAGCAGCCACCCGGGACAGGGTTTGACACTGAAGCTTCTTT	TRBV28*01	11		
TGCGCCAGCAGCGCAAAACACAGGGGGCTCGAGCACTGAAGCTTCTTT	TRBV5-1*01	10		
TGTGCCAGCACAATATTACCGGGGGGGAGGCTACACCTTC	TRBV2*01	10		
TGTGCCAGCTCCAATAAATTCACCCCTCCACTT	TRBV18*01	10		
TGTGCCAGCAGTACGGGGGGCGGTATGGGAGACCCAGTACTTC	TRBV6-6*01	9		
TGCAGTGCTAGAGTACCGGGGATTCGCAAAACATTAGTACTTC	TRBV20-1*01	8		
TGTGCCAGCTCAGGGCTACTAGCGGTTCTAGCAGATACGCAGTATTT	TRBV18*01	8		
TGTGCCAGCAGCAATCGGGGACTAGCGGGGAGAAAGTTCTTC	TRBV7-2*01	7		
TGTGCCAGCTCACCATCAAGTTTGAAGCTTCTTT	TRBV18*01	5		
TGCAGTGCTAGGGACAGAGGCCAGGGACCCAGTACTTC	TRBV20-1*01	4		
TGCGCCAGCAGCTCTAGGGACAGGGGATCTCCGTGATGGCTACACCTTC	TRBV5-1*01	4		
TGTGCCAGCAGCTAGCCCTAGCGGGCTCTAGCAGATACGCAGTATTT	TRBV11-2*01	4		
TGCAGTGCGACAGAGGGCGGAATAGCAATCAGCCCCAGCATTTT	TRBV20-1*01	3		
TGCAGTGCTAGGGGGCACCTAATCAGCCCCAGCATTT	TRBV20-1*01	3		
TGTGCCAGCAAGTTAGGAAAAGCTACGAGCAGTACTTC	TRBV28*01	3		
TGTGCCAGCAGACCCGGACAGGGGATGGAGCACTGAAGCTTCTTT	TRBV12-3*01,TRBV12-4*01	3		
TGTGCCAGCAGCCCGGACAGGGGTTGAACACTGAAGCTTCTTT	TRBV2*01	3		
TGTGCCAGCAGCTCGGGACTAGCGGGGGAGACCCAGTACTTC	TRBV5-6*01	3		
TGTGCCAGCAGTACTTTCGGGGCCCTACGAGCAGTACTTC	TRBV6-6*01	3		
TGTGCCAGTAGTATGCCCGGGTACCACTGAAGCTTCTTT	TRBV19*01	3		
TGTGCCAGCTCACCCGATAAGCGGCACTGAAGCTTCTTT	TRBV18*01	2		
TGCGCCAGCAGCTTCCGGCTCTGAGGACAGAGACCCAGTACTTC	TRBV5-1*01	2		
TGCAGCGCAGTCCGCTCAAGGCATGAACACTGAAGCTTCTTT	TRBV29-1*01	2		
TGCAGCTTGATCTGGGGGGGAGCGGCACTGAAGCTTCTTT	TRBV29-1*01	2		
TGCAGTGCTAGAGAGACAGGGTCTTATGGCTACACCTTC	TRBV20-1*01	2		
TGCGCCAGCAGAGAAGGGTGGACTGAAGCTTCTTT	TRBV5-1*01	2		
TGTGCCAGCAGCACTTAGCGGGGGCACAGATACGCAGTATTT	TRBV7-9*01	2		
TGTGCCAGCAGCTTAAGCGGGACCTACGAGCAGTACTTC	TRBV7-2*01	2		
TGTGCCAGCAGCTTAGACAGGGCAACCCCTGGCTACACCTTC	TRBV11-3*01	2		
TGTGCCAGCAGCTATATTTACAGCCCCAGCATTT	TRBV7-2*01	2		
TGTGCCAGCAGCTTGGGGGAAATCAGGGGAGCGTTTACGAGCAGTACTTC	TRBV5-4*01	2		
TGTGCCAGCAGTGGACAGGTTCCAGTACAGCCCCAGCATTT	TRBV2*01	2		
TGTGCCAGCAGGGACAGGCATATGGCTACACCTTC	TRBV25-1*01	2		
TGTGCCAGCTCTGCCAGGGAGGAAATTCACCCCTCCACTT	TRBV12-3*01,TRBV12-4*01	2		
TGTGCCAGCAGCTAGTGGGATCCCTAGCAGATACGCAGTATTT	TRBV7-8*01	1		
TGTGCCAGCAGTACCCCGGGGAGGGGTGACCCAGTACTTC	TRBV6-5*01	1		
TGTGCCAGCAGTTATTCGGGACTAGCGGGATTGCCGATGAGCAGTCTTC	TRBV6-6*01	1		
TGCAGTGCTCAGAACCGGGTGAACACCCGGGAGCTGTTTTTT	TRBV20-1*01	1		
TGTGCCAGCAGTACTTGGGGAGCACTGAAGCTTCTTT	TRBV6-2*01,TRBV6-3*01	1		
TGCAGTCCGCTTGGCGGGACAGGGGACACTGAAGCTTCTTT	TRBV20-1*01	1		
TGTGCCAGCACTCCCGGGGTAGCGGGTCTAGGCCAGTACTTC	TRBV6-5*01	1		
TGCGCCAGCAGCTTGGCCGACTAGCGGGAGCCGGAGCTCTACAATGAGCAGTCTTC	TRBV5-1*01	1		
TGTGCCAGCAGAGTGGGACAGGTAATCTACGAGCAGTACTTC	TRBV6-1*01	1		
TGTGCCAGCAGCGGACGAGCGGGGGCCATAAATGAGCAGTCTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGGACAGGGGGCAAGCGGGGAGCTGTTTTTT	TRBV7-6*01	1		
TGTGCCAGTAGTATAGATGAGGGTCAACCCCTCCACTT	TRBV19*01	1		
TGTGCCAGCACCGGATGGACAGGGTCCCATTTGAATTCACCCCTCCACTT	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCAGCAGCGCCGCGGACAGGAATATTCGGCTACACCTTC	TRBV7-9*01	1		
TGTGCCAGCAGTCTAGGGTGGCACTGAAGCTTCTTT	TRBV28*01	1		
TGCAGCACACCCGGGAAGCGGGGGTGAACACAGATACGCAGTATTT	TRBV29-1*01	1		
TGCAGCGCAGCTAGCGGGAGTGGGGAGACCCAGTACTTC	TRBV29-1*01	1		
TGCAGCTAAGGAGTCCCTACGAGCAGTACTTC	TRBV29-1*01	1		
TGCAGCTCCCGGACATAAACCAGGGGAGCTGTTTTTT	TRBV29-1*01	1		
TGCAGCTGCCTTAGGTCCACTACGAGCAGTACTTC	TRBV29-1*01	1		
TGCAGCTGGCACAGGGGACTCCGAAACCCGGGGAGCTGTTTTTT	TRBV29-1*01	1		
TGCAGCTGGCGGGGCTACAATGAGCAGTCTTC	TRBV29-1*01	1		
TGCAGCTTACGAATCAGGGAGTCTCAAAAACATTGAGTACTTC	TRBV29-1*01	1		
TGCAGCTTGAAGTACAGAGCTACGAGCAGTACTTC	TRBV29-1*01	1		
TGCAGCTTGAAGGGGGGGCCCCGGGGAGCTGTTTTTT	TRBV29-1*01	1		
TGCAGCTTGATAGGGGGTCCCAAGAGACCCAGTACTTC	TRBV29-1*01	1		
TGCAGCTTGATCCGGGGCACACAGATACGCAGTATTT	TRBV29-1*01	1		
TGCAGCTTGGTTACGGACCACTACGAGCAGTACTTC	TRBV29-1*01	1		
TGCAGTGCAGGGACCCCCCGGGAAGGGTACTTC	TRBV20-1*01	1		
TGCAGTCCCATCAGGGGACTAGCGGGGGGGCTTACAGATACGCAGTATTT	TRBV20-1*01	1		
TGCAGTGCCTCCGACACCTATGGCTACACCTTC	TRBV20-1*01	1		
TGCAGTGCCTACAAGACTAGCGGCTTCTGCTGGGAGTCTTC	TRBV20-1*01	1		

TGCAGTCCGGACAGGGATAGCTCCTACAATGAGCAGTTCTTC	TRBV20-1*01	1		
TGCAGTGTACACAGGGCCAAAGAGACCCAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGAAACAGCGGGAGGGCCGAGTCTACGAGCAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGAGATGTCATCGACCGGGGAGCTGTTTTTT	TRBV20-1*01	1		
TGCAGTGTAGAGATGTTGGAAACACCATATATTTTT	TRBV20-1*01	1		
TGCAGTGTAGAGCCCTCGACCGGAGAGACCCAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGAGCCGCTGGGGACAGAAAGGCTACGAGCAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGAGCTTCCGGGGGAGTCCAGCCCAAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGAGGGGACAGGGGACGACAGATACGAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGATATCCGGGACTAGATCAATGAGCAGTTCTTC	TRBV20-1*01	1		
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TGCAGTGTAGATCTCCGACGAACTGAAGCTTTCTTT	TRBV20-1*01	1		
TGCAGTGTAGTCCCGGGGACGGCCCTACGAGCAGTACTTC	TRBV20-1*01	1		
TGCAGTGTAGTGGAGCGGAAAGGCTACACCTTC	TRBV20-1*01	1		
TGCAGTGTGGCAGCGGAGGACAGATACGAGTACTTC	TRBV20-1*01	1		
TGCAGTGTGTACAGGGAGGGCTAATCAGCCCAAGTACTTC	TRBV20-1*01	1		
TGCAGTGTGAGCTGGGGGCTATGGCTACACCTTC	TRBV20-1*01	1		
TGCGCCAGCAGAGCGACAGGGGGGACTACGAGCAGTACTTC	TRBV5-1*01	1		
TGCGCCAGCAGCGGGACTAGCGGTAACCCGGGAGCTGTTTTTT	TRBV5-1*01	1		
TGCGCCAGCAGCAACTCGACCGTCTCGGACAGATACGAGTACTTC	TRBV4-1*01	1		
TGCGCCAGCAGCAAGATAGATCAAGCCAGTTCTTC	TRBV4-3*01	1		
TGCGCCAGCAGCAAGGAGTAGAGACTAGCGGATGAATACGAGTACTTC	TRBV4-1*01	1		
TGCGCCAGCAGCCCAAGCGGCGGCGGCGAATGAAAACCTGTTTTTT	TRBV4-1*01	1		
TGCGCCAGCAGCCCAAGGACAGACTTCGGCTACACCTTC	TRBV4-3*01	1		
TGCGCCAGCAGCCCGGACAAACAATCAGCCCAAGTACTTC	TRBV5-1*01	1		
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TGCGCCAGCAGCGTGGGACTACCTACGAGCAGTACTTC	TRBV5-1*01	1		
TGCGCCAGCAGCGGGACAGGGGACGTAAGCAGTACTTC	TRBV10-2*01	1		
TGCGCCAGCAGCGTGAACAGGGGTGGCTACACCTTC	TRBV5-1*01	1		
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TGCGCCAGCAGCTCTCAGGGGTCTACGAGCAGTACTTC	TRBV5-1*01	1		
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TGCGCCAGCAGCTTGGAAACCGCAGGGATTCACCCCTCCACTTT	TRBV5-1*01	1		
TGCGCCAGCAGCTTGGAAATCCGGCAGCAATCAGCCCAAGTACTTC	TRBV5-1*01	1		
TGCGCCAGCAGCTGGCCCCGGGCACTGAAGCTTTCTTT	TRBV5-1*01	1		
TGCGCCAGCAGCTTGGGACTAGCGGGAGGAATAGATACGAGTACTTC	TRBV5-1*01	1		
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TGCGCCAGCAGGAAAGGGAGCGGGGGGCCCCCTCTACAATGAGCAGTTCTTC	TRBV10-1*01	1		
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TGCGCCAGCCTCTCGTGGACTAGCGGGCAGACAGATACGAGTACTTC	TRBV10-2*01	1		
TGTGCCACACAGAGATCTGGGCCAGGGGACAGGGGGCGGGGAGCTGTTTTTT	TRBV15*01	1		
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TGTGCCAGCAAAGCCCCCTAGCTTGGTACTTC	TRBV6-5*01	1		
TGTGCCAGCAAAGTGGGGGCGGAGGGTCCCCCTCTACGAGCAGTACTTC	TRBV28*01	1		
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TGTGCCAGCACAGGAAAGACCCAGTACTTC	TRBV28*01	1		
TGTGCCAGCACAAAGACAGGGTGAAGCTTTCTTT	TRBV6-5*01	1		
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TGTGCCAGCACCGACCCCTGGGATGACTGAAGCTTTCTTT	TRBV2*01	1		
TGTGCCAGCACCGCAGACTAGGGAATGAGCAGTTCTTC	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCAGCACCGGGGGGGTGGCTCTACAATGAGCAGTTCTTC	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCAGCAGAATGACAGGGGTTGATCTACGAGCAGTACTTC	TRBV5-6*01	1		
TGTGCCAGCAGACTATGGGTTGGGACAGAAACCATATATTTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGACTAACAGGGTACCCTACAGATACGAGTACTTC	TRBV6-1*01	1		
TGTGCCAGCAGACTAGGACTAGCGGGGACAGATACGAGTACTTC	TRBV7-9*01	1		
TGTGCCAGCAGACTGGAGGCTCGAACACTGAAGCTTTCTTT	TRBV2*01	1		
TGTGCCAGCAGCAACTCTGAAGTTCTTC	TRBV7-9*01	1		
TGTGCCAGCAGCACTCGACAGGATCGGGAAACCATATATTTTT	TRBV11-2*01	1		
TGTGCCAGCAGCAACTCGGGGAGCTGTTTTTT	TRBV3-1*01	1		
TGTGCCAGCAGCAAGAGGGCTGTAGGGACCCAGTACTTC	TRBV3-1*01	1		
TGTGCCAGCAGCAAGAGTACGGGGCGGTACGCCAGCATTTC	TRBV3-1*01	1		
TGTGCCAGCAGCAAGATCGGGGGCCAGGGAGAGACCCAGTACTTC	TRBV14*01	1		
TGTGCCAGCAGCAAGGAAATAGCGGGGGATTCACTACTTC	TRBV3-1*01	1		
TGTGCCAGCAGCAAGGGCTAGCGGGAGGCGCGGATGAGCAGTTCTTC	TRBV14*01	1		
TGTGCCAGCAGCAAGTAGGGGACAGGGGAAACCCGGGAGCTGTTTTTT	TRBV4-2*01	1		
TGTGCCAGCAGCAAGTGGGATGAAAAAGAGACCCAGTACTTC	TRBV14*01	1		
TGTGCCAGCAGCAAGTCCGAGACTCTCGGGCAATGAGCAGTTCTTC	TRBV14*01	1		
TGTGCCAGCAGCCCAAGCGGGGAAACCCGGGAGCTGTTTTTT	TRBV7-2*01	1		
TGTGCCAGCAGCCCATAGTTACATCCAGGAGCAGTACTTC	TRBV27*01	1		
TGTGCCAGCAGCCGAGCGGACTAGCGGGCCGAAACCCGGGAGCTGTTTTTT	TRBV9*01	1		
TGTGCCAGCAGCCCTGGGACAGTTAATATGCTACACCTTC	TRBV7-8*01	1		
TGTGCCAGCAGCCGAGCGGGGAGGATGATGAGCAGTTCTTC	TRBV14*01	1		
TGTGCCAGCAGCCGAGTAACTCAACTGAAGCTTTCTTT	TRBV4-2*01	1		
TGTGCCAGCAGCCCTCGGACTAGCGGGGGCGTCAATGAGCAGTTCTTC	TRBV28*01	1		
TGTGCCAGCAGCCGGACAGGGGGCGGCTAGGCTACACCTTC	TRBV7-2*01	1		

TGTGCCAGCAGCCGGGACGTCAATCAGCCCCAGCATTTT	TRBV11-3*01	1		
TGTGCCAGCAGCCTAGAGGACCGACTGTACAGATACGAGTATTTT	TRBV6-9*01	1		
TGTGCCAGCAGCCTCGGACCCTAATGTAACCTTAATCAGCCCCAGCATTTT	TRBV7-8*01	1		
TGTGCCAGCAGCCTCTACGGGAGGACCTCTGAAACACCATATATTTT	TRBV6-6*01	1		
TGTGCCAGCAGCCTCGCCCTCGGGGGAAACACTGAAGCTTTCTTT	TRBV11-2*01	1		
TGTGCCAGCAGCCTGGACAGGTCTACTGAAGCTTTCTTT	TRBV2*01	1		
TGTGCCAGCAGCAGCCGGGGTACACCGGGGAGCTGTTTTT	TRBV7-2*01	1		
TGTGCCAGCAGCAGCGGTAGCGGATGGGGGCTATGAGCAGTTCTTC	TRBV3-1*01	1		
TGTGCCAGCAGCGGATGGTCTGGGGTGGAGCAGTTCTTC	TRBV7-2*01	1		
TGTGCCAGCAGCGCCAGGGGTTGGTGAGCAGTTCTTC	TRBV9*01	1		
TGTGCCAGCAGCGTACAGGGGATGGGGGGTGAGTACTTC	TRBV7-2*01	1		
TGTGCCAGCAGCGTAGAGGTATGGGAGACCCAGTACTTC	TRBV9*01	1		
TGTGCCAGCAGCGTAGCGACAGGGGGCAGGAGTCAAGGGGACATTTT	TRBV9*01	1		
TGTGCCAGCAGCGTAGGGGTAGCGGGAGTCTCTGAGCAGTTCTTC	TRBV9*01	1		
TGTGCCAGCAGCTCAGAACCCGGGACAGGGGGCGTGAATAACTGTTTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTCCACAGGGGGTGGGGAAGCTTTCTTT	TRBV11-2*01	1		
TGTGCCAGCAGCTCCCTCGGGGGACAGGGGCTCAAAAACTTCACTACTTC	TRBV9*01	1		
TGTGCCAGCAGCTCCGGGCTCTAGCGGGGGTCCGGGGAGCTGTTTTT	TRBV7-3*01	1		
TGTGCCAGCAGCTCGAGGGGGCGGGCCAGTACTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTCGCCACAGGGGAGAGCGGGGAGCTTTTTT	TRBV7-9*01	1		
TGTGCCAGCAGCTCGCCGGGACTAGCGGGAGGGCTCAAGGGGACATACGAGTATTTT	TRBV5-6*01	1		
TGTGCCAGCAGCTCGGACAGGTTGACAAATGAGCAGTTCTTC	TRBV7-9*01	1		
TGTGCCAGCAGCTCGGGACTAGCGGGAAATCAGTACTTC	TRBV7-7*01	1		
TGTGCCAGCAGCTTAACGTATCTCAGGAACACTGAAGCTTTCTTT	TRBV7-3*01	1		
TGTGCCAGCAGCTTAAGGCTTGGCGGGGGCGGTTGGTACGAGTATTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGGGGGAGGGTTGACGAGCAGTACTTC	TRBV7-6*01	1		
TGTGCCAGCAGCTACGTACAGGGAGGGCTCTGAAACACCATATATTTT	TRBV5-4*01	1		
TGTGCCAGCAGCTTACTAGCGGGCTACGAGCAGTACTTC	TRBV7-3*01	1		
TGTGCCAGCAGCTTAGAATCGGCCAGTACTTC	TRBV11-3*01	1		
TGTGCCAGCAGCTTAGACCTTACACAGATACGAGTATTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGATAAATCAGCCCCAGCATTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGCGCTCATAGCGGGACAGGGACCTACAATGAGCAGTTCTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGCGGGGGGATACGAGTATTTT	TRBV11-2*01	1		
TGTGCCAGCAGCTTAGGAAACCGACAGTCTACGAGCAGTACTTC	TRBV11-2*01	1		
TGTGCCAGCAGCTTAGGACAGGGTCTATAAATGAGCAGTTCTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGACTGAAGGCTACACCTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGAGGAGCGGGACCGTACGAGCAGTACTTC	TRBV7-9*01	1		
TGTGCCAGCAGCTTAGGGACAACCAAGAGACCCAGTACTTC	TRBV5-5*01	1		
TGTGCCAGCAGCTTAGGGCAGGGGATTAATGCTACACCTTC	TRBV11-2*01	1		
TGTGCCAGCAGCTTAGGGCAGGGGCTTACAGCAGTACTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGGCAGGGGCTTACGAGCAGTACTTC	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGTCAGGGGATGGACACTGAAGCTTTCTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTTATCAGGGGACAGTAGCAATCAGCCCCAGCATTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTTAGGGTCCACAGGGTGGGAATCACCCTCCACTTT	TRBV7-2*01	1		
TGTGCCAGCAGCTTCGGGGGACAGCGGGGCTATGGCTACACCTTC	TRBV11-3*01	1		
TGTGCCAGCAGCTTGCCATCGAAGAGACCCAGTACTTC	TRBV5-4*01	1		
TGTGCCAGCAGCTTGGGAGTGGGTTGCTAACTATGGCTACACCTTC	TRBV5-6*01	1		
TGTGCCAGCAGCTTGTTCGGTCACTAGCGGGCACAGATACGAGTATTTT	TRBV5-6*01	1		
TGTGCCAGCAGGAGGACTAGCGGGGAGGCAAGAGACCCAGTACTTC	TRBV7-2*01	1		
TGTGCCAGCAGGGCCAGCGGGGCAATCAAGAGACCCAGTACTTC	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCAGCAGGGGACGGGGGACTAGCGGGTTGAGTCTTC	TRBV25-1*01	1		
TGTGCCAGCAGGGGGTTCGGGGCGGGTGAAGCTTTCTTT	TRBV28*01	1		
TGTGCCAGCAGGTGGACAGCAGGAGAGAGACCCAGTACTTC	TRBV5-6*01	1		
TGTGCCAGCAGTAAACAGACAGGAGCTAACTATGGCTACACCTTC	TRBV6-5*01	1		
TGTGCCAGCAGTACGAGGACAGGAAACCCGGGGAGCTGTTTTT	TRBV11-2*01	1		
TGTGCCAGCAGTCCCGGACAGGGTTCGGGGGGGAGCTGTTTTT	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCAGCAGTCCGGGGACAGGGCTGAGAAACACAGATACGAGTATTTT	TRBV25-1*01	1		
TGTGCCAGCAGTCTCCGACACAGGGCGCGGGGAGCTGTTTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGTGAAGCAAGCTGACCTCGCCCGGAGTTACAATGAGCAGTTCTTC	TRBV6-1*01	1		
TGTGCCAGCAGTGAATCGCGAGCAGTACTTC	TRBV2*01	1		
TGTGCCAGCAGTGACAGACCACTGAAGCTTTCTTT	TRBV28*01	1		
TGTGCCAGCAGTGGCCGGGACGGAAACCCGGGAGCTGTTTTT	TRBV6-1*01	1		
TGTGCCAGCAGTGAAGGACAGGGGGCAGTGGAGCAGTTCTTC	TRBV25-1*01	1		
TGTGCCAGCAGTACGGGGATATCGCTACAATGAGCAGTTCTTC	TRBV6-6*01	1		
TGTGCCAGCAGTACTCGAATACCTACAATGAGCAGTTCTTC	TRBV6-6*01	1		
TGTGCCAGCAGTACTCGGGGGAAGTCCAATGAGCAGTTCTTC	TRBV6-2*01,TRBV6-3*01	1		
TGTGCCAGCAGTATCTGAGAGGGGGCGCTACGAGCAGTACTTC	TRBV6-5*01	1		
TGTGCCAGCAGTATCCCGGACAGCAAATCAGCCCCAGCATTTT	TRBV6-5*01	1		
TGTGCCAGCAGTCCAGACAGGGGCTTCTACAATGAGCAGTTCTTC	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGTCCGAGCGGACAGGGGAAATCACCCTCCACTTT	TRBV6-5*01	1		
TGTGCCAGCAGTTAACCACAATTAGTGAAGCTTTCTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGTTAAGCCGGGAGCAGTACTTC	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGTTAGGCCAGCCCCAGCATTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCAGTTGGTCTCAATGAGCAGTTCTTC	TRBV28*01	1		
TGTGCCAGCAGTTTCTGGGGAGCTCTGGAAACACCATATATTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGCATCTCAAGGACAGGAGGGGGCTACACCTTC	TRBV28*01	1		
TGTGCCAGCATTTCCGACAGGAGCTACGAGCAGTACTTC	TRBV6-5*01	1		
TGTGCCAGCAGTCCGGACACACTCTGAAACACCATATATTTT	TRBV3-1*01	1		
TGTGCCAGCGGGGGTCCGAAAGGAGACCCAGTACTTC	TRBV2*01	1		
TGTGCCAGCTCACCAGCGCCGGGGACGGCTACACCTTC	TRBV18*01	1		
TGTGCCAGCTCACCAGGGGACAAAGCGCTGGAGAGCAGTTCTTC	TRBV18*01	1		
TGTGCCAGCTCACCAGCTAGGACGGCTACACCTTC	TRBV18*01	1		
TGTGCCAGCTCACCAGGGGGCAGATACGAGTATTTT	TRBV18*01	1		
TGTGCCAGCTCACCAGGGGGACCGCTGGGATGGCTACACCTTC	TRBV18*01	1		
TGTGCCAGCTCACCGGGACAGGGGGCACTTGGGAGACCCAGTACTTC	TRBV18*01	1		
TGTGCCAGCTCAGAGATAGGAGACTAGCATTAGGGAGACCCAGTACTTC	TRBV7-9*01	1		
TGTGCCAGCTCGCCAAACCCGGACATCCAACACTGAAGCTTTCTTT	TRBV18*01	1		
TGTGCCAGTAGTACACAGGGTATTAATCACCCTCCACTTT	TRBV19*01	1		

TGTGCCAGTAGTATAGATCTATCGTCACAGGAGGATTTT	TRBV19*01	1		
TGTGCCAGTAGTATCCGACTCCGGGACATAAGCAATCAGCCCCAGCATTTT	TRBV19*01	1		
TGTGCCAGTAGTATCGGAGTGGGGATGGCCTAAATCAGCCCCAGCATTTT	TRBV19*01	1		
TGTGCCAGTGCCAGGACAGGGGGCAACGAGCAGTACTTC	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCAGTCCGCGAGCGGGATCGCACAGATACGCAGTATTTT	TRBV19*01	1		
TGTGCCATCAGAGGAGATACCACAGATACGCAGTATTTT	TRBV10-3*01	1		
TGTGCCATCAGCGCTAGCGGGACGAACACAGATACGCAGTATTTT	TRBV10-3*01	1		
TGTGCCATCAGTGAGCAGGGCGGGAGCTATGGCTACACCTTC	TRBV10-3*01	1		
TGTGCCATCAGTGAGTCTGTGGGAGGGCCCTCCATGAGCAGTCTTC	TRBV10-3*01	1		
TGTGCCATCAGTGGTCTGTGGACCCGGTCTACGAGCAGTACTTC	TRBV10-3*01	1		
TGTGCCATTAGCGCCCCACAGATACGCAGTATTTT	TRBV12-3*01,TRBV12-4*01	1		
TGTGCCCGGGGGGGGATCTATGGCTACACCTTC	TRBV6-5*01	1		
TGTGCCTGGACCAACAGGTCTAGCGGCGGGAAACAGATACGCAGTATTTT	TRBV30*01	1		