

**Supplemental Table 1. Probe sequences used in Ching et al., Genes & Immunity 6, 186-193, 2005**

Group 1		Group 2			
		Group 2a		Group 2b	
Name	Probe sequence, 5' to 3'	Name	Probe sequence, 5' to 3'	Name	Probe sequence, 5' to 3'
VH-1-2	Cy5- AAAGAGATTGAGTCCAGTCC	VH-3-29P	Cy5- TATAGAGGAGGAGGGAGGGCC	VH-3-53	Cy5- CACTTCATCTGCACGGAGCCCG
VH-3-2.1P	Cy5- AATTTACAGCCCAGTGGTGAG	VH-3-32P	Cy5- CAGGCCCTCCCTCTCTCTGG.	VH-4-53.1P	Cy5- CCGAGCTAAGTGAGTCACAAGCA
VH-1-3	Cy5- TGAGGAGGAAGGGATCGAAG	VH-3-30&33	Cy5- ATAACGGTGTCTCTCTGTTT	VH-3-54P	Cy5- GAACCTTCTGACCAGGACAC
VH-3-5.1P	Cy5- AAGGACACGTCACTACATTA	VH-3-33.2P	Cy5- CAGACACAGTTTCTGCAATG	VH-4-55P	Cy5- AACTGGTGAATCTGGGTCCG
VH-3-5.2P	Cy5- CCAGATTCTGGAAAACCAC	VH-3-30.2P	Cy5- ACAGACACAGATTTCTGCA	VH-7-56P	Cy5- GCACGGCGGATCTGCAGACT
VH-3-6P	Cy5- TTCTGCTTGGTCAACACTGAA	VH-4-30.1P	Cy5- TACCCAAGCCTCCCTGTGGG	VH-3-57P	Cy5- GAGGCAGAAAATTGTACAGCCC
VH-3-7	Cy5- CCAACATAAAGCAAGATGGA	VH-4-31	Cy5- CGGACTCGCTATGTTTCTGGAT	VH-1-58	Cy5- CTCGTGGACAACGCCTTGAG
VH-1-8	Cy5- AGTAAGGGGCTTCTAGTCTC	VH-4-31.1P	Cy5- TCAACACTGTGTTTTCCAGT	VH-4-59	Cy5- GGCCCATATAGCTTACAGA
VH-3-9	Cy5- TGGACATGAGTGGATTTGTG	VH-4-33.1P	Cy5- CCCAAGCCTCCCTGTGGG	VH-3-60P	Cy5- CTGTCTCCACTTGGTCATGA
VH-3-13	Cy5- CTGGTGACACATACTATCCAGG	VH-4-34	Cy5- CTATGGTGGGTCCTTCAAGTGG	VH-4-60.1P	Cy5- TCACAAACAACCTGTGTTTCTT
VH-3-15	Cy5- GTATTAAGCAAAACTGAT	VH-3-42P	Cy5- ACTAGAGACAGTGAGGGGGA	VH-4-61	Cy5- GACCCATTATAACTTATCGA
VH-3-16	Cy5- CAGCGCAGCTGACCCTGCA	VH-3-43	Cy5- CTCTTATTAGTTGGGATGGT	VH-3-62P	Cy5- ATACAATTATTCCAATCTGT
VH-3-16.1P	Cy5- ACAATTTTCCCCCTGACTCT	VH-4-43.1P	Cy5- CATTTAGGTTTAAATTCTGG	VH-4-62.1P	Cy5- CAGAAATGTATGTAGTTGGG
VH-1-18	Cy5- GATCAGCGCTTACAATGGTA	VH-3-44P	Cy5- GCAGCTGGTAGACTCTTTTTT	VH-3-63P	Cy5- GCTGGGTCAATGAGACTCTA
VH-3-19P	Cy5- GTGGCGTTTTCTTAACTCAC	VH-4-44.1P	Cy5- TTCATACGTGGGAGACACG	VH-3-64	Cy5- AGAAGAGTCATGATGGAGTT
VH-3-20	Cy5- TGAAAACCACCCAGCCCTGA	VH-4-44.2P	Cy5- CACACGATGATCCATCTCTGC	VH-3-65P	Cy5- GTCAAAACCCACTGAGGACA
VH-3-21	Cy5- ATCATGAAAAGTAGAGAGATTT	VH-1-45	Cy5- CCGTCAGAAAACCCAAAGGAG	VH-4-65.1P	Cy5- AAAAACAGTGTTCCTGTAGCTG
VH-3-22.2P	Cy5- GAGCTGGGTTTTCTTTGTTGCT	VH-1-46	Cy5- CTGAGGAAGGGATTTTTTCC	VH-3-66	Cy5- GACTTCATCTGCACTGGGCCCG
VH-3-25P	Cy5- GATATGAGTGAAAACAGTG	VH-4-46.1P	Cy5- ATGGGTCTATCGGGTTAGGA	VH-1-67P	Cy5- CCAGGACTAAGGTGACTGGG
VH-3-25.1P	Cy5- ATTTCTCTGTCTCCCCGGT	VH-3-47P	Cy5- TCGCCGGGCTCCAGGAAAGGTCC	VH-4-67.1P	Cy5- TCACACACTGACCTCACCA
VH-3-26.1P	Cy5- CTGGTTTTCGGGTTTACTGGGT				
VH-4-28.1P	Cy5- ATAAAAATGTGTGGTTTTGGG				
VH-6-1	Cy3- TCATCTTCTGCCCGTGCTG	VH-7-34.1P	Cy3- AGCAGCGGTGTCTTCAGATC	VH-3-67.2P	Cy3- GGTTTTCTCAGTTTTATTGGT
VH-4-1.1P	Cy3- GAAGGCGCTGGGACACCT	VH-3-35	Cy3- CAGAGCTCGCTGTGAAATA	VH-3-67.3P	Cy3- TGATAAATGGTGCTTAGGACA
VH-4-4	Cy3- GGAGGTCCCGAATACTACATT	VH-3-36P	Cy3- GTCAGGTCTCTGAACACAGA	VH-3-67.4P	Cy3- TATGAAGTTCAGTGGTGGAG
VH-2-5	Cy3- CACAGGTCCAGCTCAGTGAC	VH-3-37P	Cy3- GGGGTGTCTCTGAGACTCTCAT	VH-1-68P	Cy3- ACCCTCCAATGGGTGTCTGC
VH-2-10P	Cy3- GAGCTTTACTCCAGCTTCTC	VH-3-38	Cy3- TTTTAGCACTCTGGGAGTGG	VH-1-69	Cy3- CCCATCTTTGGTACAGCAAA
VH-3-11	Cy3- CTAGAGACATTGAGTGGACGT	VH-3-38.1P	Cy3- AGTGAGATCAGACAGGAAAAT	VH-2-70	Cy3- CTCTGCTACTGACTGTCCCG
VH-3-11.1P	Cy3- CTGCTGCTCATGGGCTGCT	VH-4-39	Cy3- CCAACTCATAAGGGAAATGC	VH-3-71P	Cy3- TTGTTCTTCTGTATTTTTGT
VH-1-12P	Cy3- TCATGGGCTGGACCTGCAAGAT	VH-7-40P	Cy3- ATATGTTTTTCAACACCAT	VH-3-72	Cy3- TACACCACAGAATACGCCGCG
VH-3-13.1P	Cy3- TGTGGCTGTTGCCTCTGTGTT	VH-4-40.1P	Cy3- TGTATCATCCAGAACTTGTG	VH-3-73	Cy3- TAACAGTTACGCGACAGCATAT
VH-1-14P	Cy3- CTGATGACTCTCTCTGTTGCC	VH-3-41P	Cy3- TGAGTGGCACATATCTGGAA	VH-3-74	Cy3- ATAGTGATGGGAGTAGCACAA
VH-4-15.1P	Cy3- CAGTGCACATGTTCTAGT	VH-3-47.1P	Cy3- GCTCCAGTGGGGGCTGTT	VH-4-74.1P	Cy3- TTATTCATAACTCCAGGTGCC
VH-1-17P	Cy3- CCTCCTCTATAGAAGCCCTG	VH-3-48	Cy3- TGAATATTTAAATGAAATA	VH-3-75P	Cy3- AATTTCAAGGTTTTTCCACAT
VH-4-20.1P	Cy3- TGATGGAGCTGGATTCAGGG	VH-3-49	Cy3- AGGGACCGCTGAGTACGGG	VH-3-76P	Cy3- TGAAGACATCCCATGGAGTT
VH-4-22.1P	Cy3- ATCCTCCCCCTGGCCTGTGG	VH-4-49.1P	Cy3- AACACAACGCAACCCACGCCCT	VH-3-76.1P	Cy3- TGGCTGCTTCTCAGCCTGTT
VH-3-22P	Cy3- TAGCGCCCCCTGGTGGC	VH-3-50P	Cy3- GACTCAGATGTCTCTGTGCT	VH-5-78P	Cy3- TCATCCTTTCCCTCGTCTG
VH-3-23	Cy3- CCAGCCCTGGGATTTTCAG	VH-5-51	Cy3- CATTTTCTGTCTCCACCATC	VH-4-78.1P	Cy3- GAGGTCTTCTGCTGAAGCT
VH-1-24	Cy3- GGCAAGAGAATCTGAGTTC	VH-3-51.1P	Cy3- CGGATGTTTGTGAAGAAAT	VH-3-79P	Cy3- CCCGTGATAACGAGGTCTTC
VH-2-26	Cy3- TGGTCAAGGACTTCTCTCAGA	VH-4-51.2P	Cy3- AATGCCACAACCTTGTATTAC	VH-4-80P	Cy3- GGCCTTCTGCTTACACGGGA
VH-4-26.2P	Cy3- TTACCATGGATATTATTGA	VH-3-52P	Cy3- ATTTAGTCTTCAGGGCTGAG	VH-7-81	Cy3- GTATCTGTACCCTGAAAGCA
VH-7-27	Cy3- CCACAGCACCTCAACATGTA			VH-3-82P	Cy3- AAGTCCTGTTGAAGCTTACT
VH-4-28	Cy3- AGTAACTGGTGGGCTGGAT				