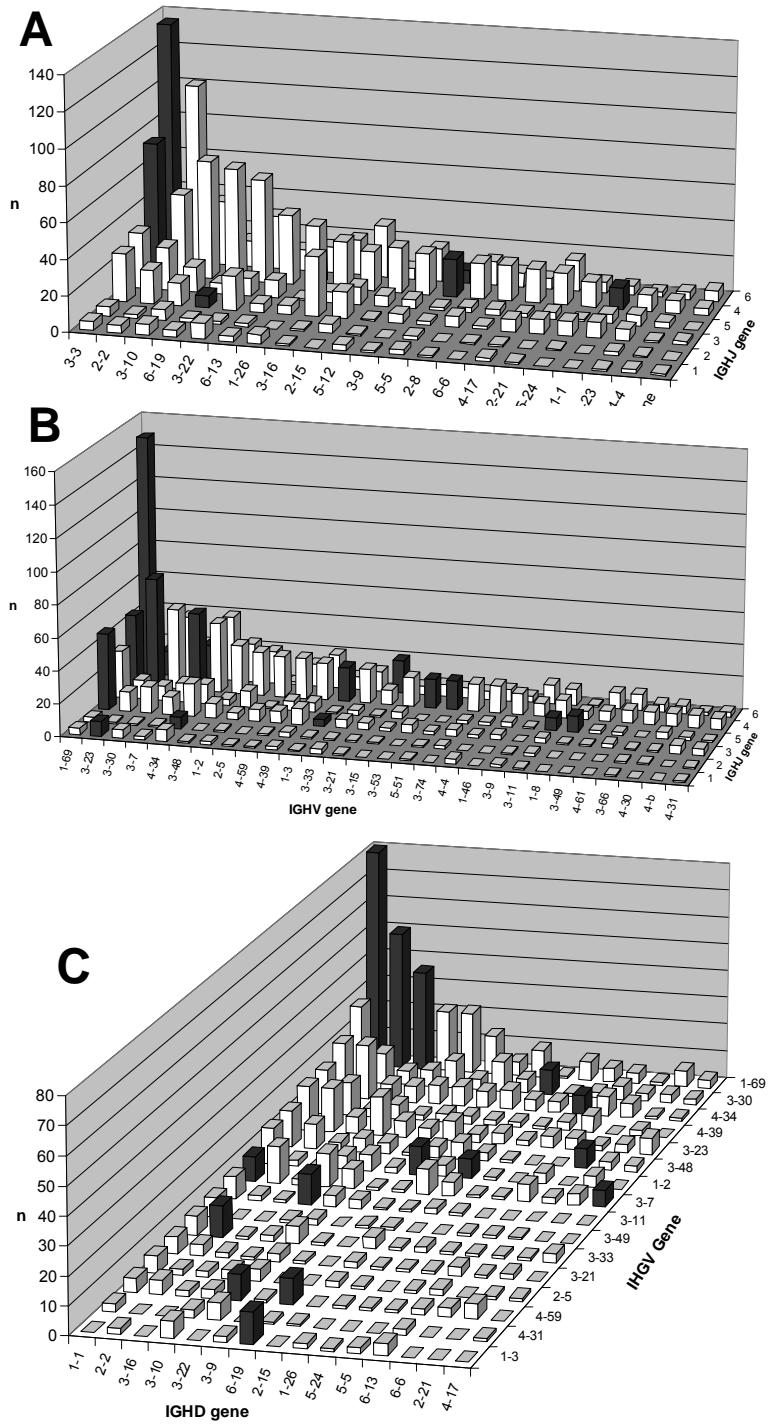


Supplemental Table I. Statistical evalution of V-D, V-J, and J-D association frequencies.

IGHV	IGD	n _{VD} (n _V) (n _D)	p value	IGHV	IGHJ	n _{VJ} (n _V) (n _J)	p value	IGHJ	IGD	n _{DJ} (n _D) (n _J)	p value
1-2	1-26	7 (64) (60)	0.006	1-3	J4	21 (36) (629)	0.024	1-01	4	12 (31) (629)	0.019
1-2	6-19	10 (64) (97)	0.003	1-69	J6	149 (276) (505)	<0.001	3-03	6	135 (277) (505)	0.007
1-3	6-19	11 (36) (97)	<0.001	1-69	J4	44 (276) (629)	<0.001	3-03	4	74 (277) (629)	<0.001
1-69	2-2	50 (276) (196)	0.002	1-69	J3	47 (276) (204)	0.029	4-4	4	7 (18) (629)	0.010
1-69	3-16	36 (276) (87)	<0.001	1-8	J5	10 (17) (152)	<0.001	5-5	6	7 (37) (505)	<0.001
1-69	3-3	80 (276) (277)	<0.001	2-05	J4	25 (36) (629)	<0.001	5-5	4	21 (37) (629)	<0.001
3-11	3-10	11 (35) (141)	<0.001	3-07	J6	21 (95) (505)	0.030	6-19	3	7 (97) (204)	0.027
3-30	1-26	9 (119) (60)	0.041	3-07	J4	48 (95) (629)	0.030				
3-33	2-2	11 (45) (196)	0.020	3-11	J5	8 (35) (152)	0.014				
3-48	6-6	7 (68) (34)	<0.001	3-15	J4	18 (30) (629)	0.025				
3-49	3-3	7 (19) (277)	0.041	3-21	J6	20 (36) (505)	0.003				
3-7	3-3	9 (95) (277)	0.026	3-23	J6	15 (117) (505)	<0.001				
3-7	4-17	6 (95) (37)	0.019	3-23	J4	68 (117) (629)	<0.001				
4-31	3-22	6 (17) (128)	<0.001	3-23	J1	9 (117) (52)	0.011				
4-34	5-5	7 (129) (37)	0.029	3-53	J4	18 (30) (629)	0.025				
4-59	3-22	9 (54) (128)	0.032	4-34	J2	7 (129) (35)	0.019				
4-59	6-19	9 (54) (97)	0.002	5-51	J4	18 (28) (629)	0.009				

Supplemental Table II. Light chain gene use among several cluster pairs.

cluster	V	J	% mutation
44	κ1-33	κ4	0.0%
	κ1-33	κ4	0.0%
47	κ1-5	κ2	3.6%
	κ1-39	κ2	6.8%
48	κ3-20	κ2	3.6%
	κ3-20	κ2	4.1%
8	κ2-28	κ5	0.0%
	κ2-28	κ1	0.5%
59	κ2-28	κ2	0.0%
	κ2-28	κ4	0.0%
62	λ3-21	λ3	1.4%
	λ3-21	λ3	1.4%
65	λ1-51	λ3	0.0%
	λ1-51	λ3	0.5%
67	κ3-20	κ4	0.0%
	κ3-20	κ4	0.0%



Supplemental Figure 1. Pairwise association of V, D, and J genes. The number of occurrences of each D-J (A), V-J (B), and V-D (C) pair is shown. The dark bars indicate statistically significant values, as determined by Chi squared statistics.