Table 3. Fetal liver heavy chain CDR3 lengths in pol λ-, pol μ- and pol μ/ pol λ -double-deficient mice

| Embryos [a] | CDR3 length in bp (number of sequences) | | | CDR3 size range in bp |
	Total sequences	In frame sequences	Out of frame sequences	Total sequences
+/+	32.8 (67)	32.0 (24)	33.2 (43)	12-42
pol μ -/-	33.3 (53)	32.9 (23)	33.7 (30)	18-45
pol λ -/-	27.8 (68)	28.5 (26)	27.1 (42)	12-47
pol μ/pol λ -/-	29.0 (44)	28.3 (19)	30.1 (25)	12-42

a) V_HDJ_{H4} rearrangements were amplified from at least three embryos for each genotype.