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Please note that an error appeared in the online early release version of this article. The final html, pdf, and print versions have been corrected. For reference, the correction appears below.

All data in the bottom row of Table II should have been shifted one column to the right. The corrected table appears below.

Table II. Mutations in VH3-23-C γ transcripts from PBL from ATRD patients and controls^a

Age (yr)	Clones			Mutations in V region		Clustering ratio	R/S	
	All	Distinct	Unmutated	Total	% bp		CDRs	FRs
ATRD patients								
ATRD1	25	15	13	0	302	8.6	0.73	3.4
ATRD2	13	15	14	2	258	6.8	0.73	3.7
ATRD3	15	6	6	0	194	11.9	0.62	3.9
Total		36	33	2	754	8.4	0.70	3.6
Controls								
C1	14	12	11	1	100	3.4	1.22	5.9
C2	adult	11	11	0	275	9.3	0.73	4.5
C3	28	6	6	0	147	9.0	0.69	5.7
C4	24	8	8	0	178	8.2	0.51	14.0
C5	30	4	4	0	49	4.5	0.53	— ^c
Total		41	40	1	749	6.9	0.70	6.2

^aTotal RNA was prepared from PBL from all the patients and controls, except for donor C2; buffycoat was obtained and CD27 positive B cells were used.

^bFR, framework region.

^cToo few mutations to give a reliable ratio.