PCR (TCRγ)				Flow cytometry
Patient	# PCR products	# T-cell clones Minimum (m) Maximum (M)		Vβ Spectratyping (CD8)**
Α	3	2	3	Clonal (Vβ8; 86%)
В	2	1	2	Clonal (Vβ17; 98%)
D^{\dagger}	3	2	3	Oligoclonal [‡] (Vβ14; 46%)
F	2	1	2	Oligoclonal [‡] (Vβ2; 68%)
G	7	4	7	Oligoclonal (Vβ16; 16%)
Н	2	1	2	Oligoclonal (Vβ8; 16%)
I	3	2	3	NA [§]

^{*} The maximum number of T-cell clones (M) assumes rearrangement of one TCRγ allele in each clone, whereas the minimum number (m) assumes that both TCRγ alleles are rearranged in all clones (if m is even) or in m-1 clones (if m is odd)

^{**} The dominant Vβ clone and the percent of all CD8+ cells expressing this clone in the cell population tested are indicated in parentheses

[†] The cell population tested for this patient contained a mixture of two limiting dilution T-cell products

[‡]Oligoclonal pattern including one prominent clone and several smaller clones

[§] Uninterpretable (high background)