

Supplementary Table 1: Characteristics of EATL patients

	Total (N=19)	<i>de novo</i> EATL (N=8)	RCDII-EATL (N=11)	Stat
<b>Clinical features</b>				
Age, mean in years (range)	55.9 (38-73)	57.4 (38-73)	55 (37-71)	ns
Gender (F, %)	12 (63%)	4 (50%)	8 (73%)	ns
<b>Time from CD diagnosis and molecular evaluation, mean in years (range)</b>	4.7 (0-21.7)	1.7 (0-10.9)	6.8 (0.6-21.7)	<b>p=0.0064</b>
<b>Biological features</b>				
<b>TCR status (frozen biopsies)</b>				
Gamma clonal	19 (100%)	8 (100%)	11 (100%)	ns
VDJ Beta clonal	4/18 (22%)	3/7 (43%)	1 (9%)	ns
<b>Mutational status (frozen biopsies)</b>				
<i>JAK1/STAT3</i> mutations				
<i>JAK1</i>	14 (74%)	8 (100%)	6 (55%)	<b>p=0.045</b>
<i>STAT3</i>	10 (50%)	2 (25%)	7 (64%)	ns
<i>TNFAIP3</i> mutations	5/18 (28%)	0	5/10 (50%)	<b>p=0.036</b>
<i>TNIP3</i> mutations	1/18 (6%)	1 (13%)	0/10	ns
<i>DDX3X</i> mutations	6 (32%)	2 (25%)	4 (36%)	Ns