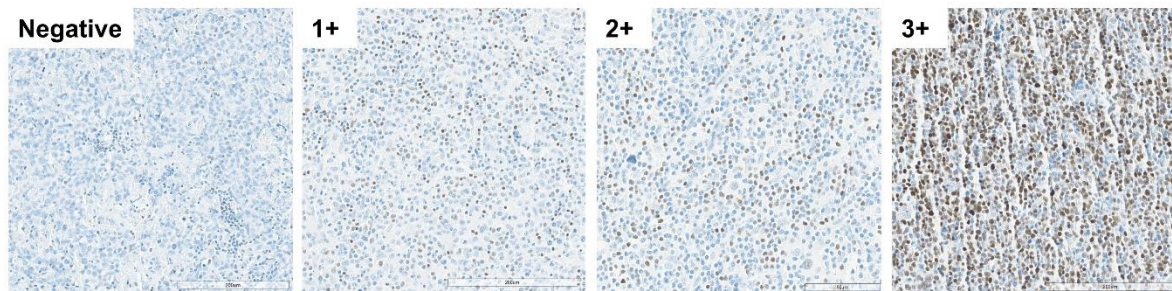
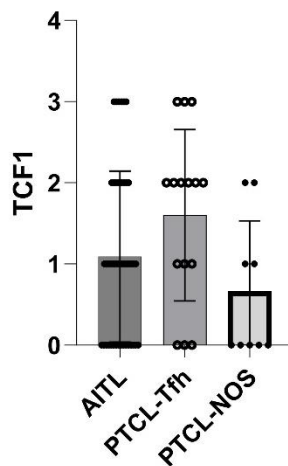


A

TCF1



B



C

		Total				Non-AITL				PTCL-NOS			
		TBX21	GATA3	Unclassified	P	TBX21	GATA3	Unclassified	P	TBX21	GATA3	Unclassified	P
		(n=37)	(n=17)	(n=12)		(n=10)	(n=10)	(n=4)		(n=3)	(n=4)	(n=2)	
TCF1	Negative	8 (21.6)	7 (41.2)	8 (66.7)	0.014	1 (10.0)	5 (50.0)	2 (50.0)	0.122	1 (33.3)	3 (75.0)	1 (50.0)	0.539
	Positive	29 (78.4)	10 (58.8)	4 (33.3)		9 (90.0)	5 (50.0)	2 (50.0)		2 (66.7)	1 (25.0)	1 (50.0)	

Supplementary Fig. S1. Expression pattern of TCF1 in nodal peripheral T-cell lymphoma. Representative immunohistochemical image of TCF1 (A). Expression patterns for TCF1 in 3 PTCL entities (B). Correlation between TCF1 expression and TBX21 and GATA3 subtype classification in 3 PTCL entities (C). GATA3, GATA binding protein 3; PTCL, peripheral T-cell lymphoma; TBX21, T-box transcription factor 21; TCF1, T-cell factor-1; AITL, angioimmunoblastic T-cell lymphoma; PTCL-NOS, peripheral T-cell lymphoma, not otherwise specified; PTCL-Tfh, peripheral T-cell lymphoma of follicular helper T-cell phenotype.