Angioimmunoblastic T-cell lymphoma is the most common T-cell lymphoma in two distinct French information data sets

Reliable information regarding the current prevalence of peripheral T-cell lymphoma (PTCL) entities is missing. Herein we report on the frequency of PTCL entities in France between 2010 and 2013. Using Lymphopath, a national lymphoma network established by the French National Cancer Agency, which covers about 70 % of all lymphomas currently diagnosed in France, we found that PTCL comprised 6.5 % of non-cutaneous lymphomas with angioimmunoblastic T-cell lymphoma (AITL) being the most frequent (739 cases; 36 %), followed by peripheral Tcell lymphoma not otherwise specified (PTCL-NOS) (550 cases, 27%). These data were verified in an independent data set from a transnational research consortium active in three European countries. In comparison to epidemiologic data reported previously, we show that AITL is by far the most common PTCL subtype. In light of the results of recent molecular findings highlighting the heterogeneity of T-cell lymphomas and the advent of targeted therapies, these data have important implications for both basic and clinical research.

Peripheral T-cell lymphomas (PTCLs) represent diverse and complex diseases, estimated to represent an overall 10-15% of all lymphomas worldwide, with the highest incidence rates occurring in Asia.¹² The relative prevalence of PTCL entities delineated according to the criteria of the REAL (1994) /WHO (2001) classification systems, was evaluated in multiple institutions in the late 1990s and early 2000s, based on retrospective cohorts of patients. In those series, PTCL-NOS (a "by default" diagnosis for cases not fulfilling the criteria for other more specific entities) was the most frequent entity, followed by anaplastic large cell lymphoma (ALCL) and AITL, while extranodal entities, in general, accounted for a small proportion of the cases.^{13,4}

This worldwide epidemiology was most recently addressed by the International PTCL study, which reviewed more than 1,300 patients diagnosed with PTCL between 1990 and 2002 in North America, Europe and Asia.⁵ In this cohort, PTCL-NOS was the most common diagnosis (25,9%), followed by AITL (18,5%), representing 30.4% and 21.7% of non-cutaneous PTCL, respectively (Figure 1).⁵ This study also confirmed geographic variations in the distribution of PTCL entities, notably demonstrating that the highest frequencies of AITL and enteropathy-associated T-cell lymphoma (EATL) are in Europe.

Here, we provide recent data obtained from a large prospective survey in France. The prevalent analysis of PTCL was derived from data collected through *Lymphopath*, a national network of 33 expert reference centers for hematopathology which was established by the French National Cancer Agency in 2010. Non-expert pathologists are encouraged to refer every newly diagnosed lymphoma for review to a *Lymphopath* center. Diagnoses provided by experts, following slide review and additional ancillary techniques performed in the reference center, are entered into a central database. Of the 31,401 non-cutaneous lymphomas reviewed in Lymphopath over four years (2010-2013), there were 2,046 cases of PTCLs (6.5%) which comprised: 739 AITL (36.1%), 550 PTCL-NOS (26.9%), 176 ALK+ ALCL (8.6%), 162 ALK- ALCL (7.9%), 107 extranodal NK/T-cell lymphomas, nasal-type (5.2%), 77 EATL (3.8%), 52 adult Tcell leukemia/lymphoma (2.5%) and 20 hepatosplenic T-cell lymphomas (1.0%). The remaining 163 cases (8%) included



Figure 1. Relative frequency of non-cutaneous PTCL entities according to the International peripheral T-cell lymphoma (PTCL) project (worldwide (n=1314) and European (n=450) statistics)⁵ and in the Lymphopath registry (France) (n=2046). ALCL: anaplastic large cell lymphoma; ALK: anaplastic lymphoma kinase; ATLL: adult T-cell leukemia/lymphoma; EATL: enteropathy-associated T-cell lymphoma; HSTL: hepatosplenic T-cell lymphoma; NK/T: extranodal NK/T-cell lymphoma; PTCL-NOS: peripheral T-cell lymphoma, not otherwise specified. For the International T-cell lymphoma project, the percentage of the different non-cutaneous PTCL entities have been adjusted, after exclusion of cases not confirmed to be T-cell lymphomas (for the international statistics) and exclusion of categories corresponding to cutaneous lymphomas and "other disorders".

	Mourad Blood 2008 ⁷	Lachenal Medicine 2007 [®]	Tokunaga Blood 2012°	Federico JCO 2012 ¹⁰	TENOMIC
Time period for case collection	1987-2008	1990-2004	1990-2008	1990-2002	1999-2009
	GELA	Lyon	Japan	International T-cell	France, Belgium,
				project	Switzerland
N.	157	77	207	243	246
Median age	62	64	67	65	66
(year) [range]	[20-89]	[30-91]	[34-91]	[20-86]	[27-92]
Male to female ratio	1.5:1	1.26:1	1.8:1	1.3:1	1.5:1
B symptoms	72%	77%	60%	69%	64%
	(112/155)	(59/77)	(122/202)	(168/202)	(139/217)
Stage III - IV	81% (126/156)	92% (71/77)	90% (186/207)	89% (214/241)	98% (235/240)
Bone marrow involvement	47% (71/151)	60% (NA)	29% (59/202)	28% (67/228)	42% (91/218)
Splenomegaly	55% (86/155)	51% (39/77)	NA	35% (NA)	26% (56/215)
Hepatomegaly	31% (38/154)	26% (20/77)	9% (18/192)	26% (NA)	15% (35/230)
Anemia	65% [†] (101/155)	51% [‡] (39/76)	61% [‡] (126/207)	33% ^{**} (81/205)	61% [‡] (135/220)
Coombs positivity	33% (30/92)	58% (38/66)	46% (29/63)	NA	58% (88/153)
Hypergammaglobulinemia	50% (73/146)*	51% (37/73)¥	54% (89/166) [*]	30% (74/166)*	65% (102/156)*
LDH >1N	66%	71%	75%	60%	70%
	(98/149)	(55/77)	(154/205)	(146/223)	(146/208)
IPI score ≥ 2	89% (117/132)	87% (NA)	89% (175/197)	79% (192/222)	95% (217/229)
PIT score ≥2	74% (106/144)	NA	77% (153/198)	63% (154/233)	78% (167/213)
Median follow-up (months)	69	35,5	42	NA	61,5
3-years survival	NA	49%	54%	NA	41%
5-years survival	33%	NA	41%	32%	34%

Table 1. Clinical characteristics of 246 AITL patients with follow-up data from the Tenomic collection with comparison with other published series.

[†]Hemoglobin<12 gr/dL, [†]Hemoglobin<11.5 gr/dL, ^{††}undefined Hemoglobin level for anemia, ^{*}>12 gr/dL, ^{*}>16 gr/dL, NA: not available.

74 cases of NK/T-cell leukemias (15 aggressive NK-cell leukemias, 14 T-prolymphocytic leukemias and 45 large granular lymphocyte leukemias), and 89 other, or unclassifiable cases. Thus, in *Lymphopath*, AITL represents the most frequent non-cutaneous PTCL entity (36.1%) followed by PTCL-NOS (26.9%), which is distinctly less recurrent.

In comparison to the European data of the International PTCL study (Figure 1), the current findings differ principally with respect to the relative prevalence of AITL and PTCL-NOS.5 The reason for this discrepancy is unclear. The international study comprised a total of 450 European cases collected from eight submitting centers (two centers from Spain and Italy, and one center each from France, Germany, UK, and Norway), whereas the current analysis encompasses a much higher number of cases, estimated to be comprised of more than 70% of all new lymphoma diagnoses in France. Yet, despite the coverage not being exhaustive, it thus far represents the most recent and largest prospectively collected series validated by expert hematopathologists, with unrestricted access to ancillary diagnostic tools.

Although a selection bias inherent to the design of the international study (retrospective collection, limited number of submitting centers, university medical centers only) cannot be excluded, the differences observed may in fact reflect geographical differences within Europe. An overrepresentation of EATL in Nordic countries is well recognized, but this largely rare disease has only minimal impact on global statistics. Interestingly, according to the data collected by the national Swedish Lymphoma Registry over a 10-year period (2000-2009), the 755 PTCL (non-cutaneous and non-leukemic) registered in Sweden comprised 34% PTCL-NOS, 29% ALCL, only 14% AITL, and 9% EATL, a distribution also markedly different from both the published European statistics and from our findings, thus further suggesting geographic variations.⁶

Another potentially conflicting difference is that the international study cohort comprises cases diagnosed between 1990 and 2002 and reviewed with reference to the 2001 WHO classification, while the *Lymphopath* dataset is derived from a more recent observation period (2010-2013). In the interim, it was discovered that AITL is associated to CD10 expression and derived from T follicular helper (TFH) cells. These perceptions were incorporated into the description of AITL in the 2008 WHO classification,² and since then CD10 and novel markers associated to TFH differentiation have been validated for diagnostic use and increasingly incorporated into routine practice.⁷ Thus, consideration must be given to whether the higher prevalence of AITL observed recently might be linked in some way to the availability of novel ancillary tools to support the diagnosis.

Confronted with the lack of a similar systematic database antedating Lymphopath in France, we used another nonoverlapping set of PTCL patients to compare the distribution of PTCL entities over an earlier time period. This set of patients was collected through Tenomic, a transnational research consortium on T-cell lymphomas involving several LYSA (Lymphoma Study Association) centers in France, Belgium and Switzerland. The *Tenomic* database (approved by the ethical committee "CPP Ile-de-France IX 08-009") is a retrospective and prospective collection of PTCL samples with available frozen tissue, and corresponding clinical annotations (Online Supplementary Methods), including a subset of patients enrolled in GELA/LYSA studies. All Tenomic cases are reviewed by at least two expert hematopathologists belonging to the consortium, and classified according to the 2008 WHO criteria. The Tenomic database comprised 623 non-anaplastic PTCL cases diagnosed between 1999 and 2009, reviewed after 2008. Remarkably, within the limits of this retrospective collection, there was an even higher ratio of AITL (n=293) to PTCL-NOS (n=171) in the Tenomic dataset (1.7:1) than in that of Lymphopath (1.35:1). According to the detailed records available for 236 AITL cases (Online Supplementary Methods and Online Supplementary Table S1), the characteristic morphological features of AITL (polymorphic cellular infiltrate (100%), arborizing vessels (98%) blast cells (94%), clear cells (69%) and sinus sign (72%)), were present in most cases. In addition, an expansion of follicular dendritic cells (FDCs), regarded as a hallmark of AITL,⁷ was present in 93% of the cases, and EBV-positive blasts were evidenced in 77%. The percentages of cases expressing CD10 (83%) and the TFHassociated molecules PD1 (78%), CXCL13 (76%) and BCL6 (62%) in the neoplastic cells were comparative to those previously reported.⁷ In addition, among the 246 patients who had available follow-up data, the frequencies of B symptoms (67%), advanced stage disease (98%), IPI score>2 (94%), hypergammaglobulinemia (64%) and positive Coombs' test (41%) (Table 1), were overall similar to those recorded in other series, 8-11 providing evidence that the pathological AITL diagnoses were concordant with the clinical and biological features of that entity. The five-year overall survival (OS) for the entire group, whose treatment characteristics are detailed in the Online Supplementary Table S2, was 34% (IC 95% [27%-40%]) (Online Supplementary Figure S1), which is also in agreement with other series.⁸⁻¹¹

Interestingly, 43 of the 293 *Tenomic* AITL cases (14.6%) had initially been diagnosed as PTCL-NOS, and reclassified as AITL cases by expert review. The pathological and clinical features (shown in *Online Supplementary Tables S3 and S4*) of these reclassified cases did not differ from those of the entire cohort, suggesting that the apparent "increase" in AITL frequency may at least partly reflect the underrecognition or underdiagnosis of this entity in previous years.

This report further confirms the poor long-term outcome for AITL patients, and in view of the high prevalence of this disease there are essential, but as yet unfulfilled, needs for new therapeutic options. Importantly, the response to novel agents in AITL patients may be distinctively different from that seen in other PTCL patients.¹² In that respect, the recent findings of recurrent mutations in genes coding for enzymes controlling DNA methylation (*TET2, IDH2* and *DNMT3*),^{15,14} and in *RHOA* (coding for a small GTPase),¹⁵ offer rationale for the use of demethylating agents and/or specific inhibitors in AITL patients.¹⁶ As a whole, the data presented herein are highly relevant at a time when there is a shift towards the development of individualized therapies in PTCL, and should be taken into consideration for the design of future clinical studies.

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References

- Anderson JR, Armitage JO, Weisenburger DD. Epidemiology of the non-Hodgkin's lymphomas: distributions of the major subtypes differ by geographic locations. Non-Hodgkin's Lymphoma Classification Project. Ann Oncol. 1998;9(7):717-20.
- Swerdlow S, Campo E, Harris N, et al. WHO Classification of Tumours of Haematopoietic and Lymphoid Tissues. Lyon: IARC Press, 2008.
- Melnyk A, Rodriguez A, Pugh W, et al. Evaluation of the Revised European-American Lymphoma classification confirms the clinical relevance of immunophenotype in 560 cases of aggressive non-Hodgkin's lymphoma. Blood. 1997;89: 4514-4520.
- Lopez-Guillermo A, Cid J, Salar A, et al. Peripheral T-cell lymphomas: initial features, natural history, and prognostic factors in a series of 174 patients diagnosed according to the R.E.A.L. Classification. Ann Oncol. 1998;9(8):849-855.
- Vose J, Armitage J, Weisenburger D. International TCLP. International peripheral T-cell and natural killer/T-cell lymphoma study: pathology findings and clinical outcomes. J Clin Oncol. 2008;26(25):4124-4130.
- Ellin F, Landstrom J, Jerkeman M, et al. Real-world data on prognostic factors and treatment in peripheral T-cell lymphomas: a study from the Swedish Lymphoma Registry. Blood. 2014;124(10):1570-1577.
- Attygalle AD, Cabecadas J, Gaulard P, et al. Peripheral T-cell and NKcell lymphomas and their mimics; taking a step forward - report on the lymphoma workshop of the XVIth meeting of the European Association for Haematopathology and the Society for Hematopathology. Histopathology. 2014;64(2):171-199.
- Mourad N, Mounier N, Briere J, et al. Clinical, biologic, and pathologic features in 157 patients with angioimmunoblastic T-cell lymphoma treated within the Groupe d'Etude des Lymphomes de l'Adulte (GELA) trials. Blood. 2008;111(9):4463-4470.
- Lachenal F, Berger F, Ghesquieres H, et al. Angioimmunoblastic T-cell lymphoma: clinical and laboratory features at diagnosis in 77 patients. Medicine (Baltimore). 2007;86(5):282-292.
- Federico M, Rudiger T, Bellei M, et al. Clinicopathologic characteristics of angioimmunoblastic T-cell lymphoma: analysis of the international peripheral T-cell lymphoma project. J Clin Oncol. 2013; 31(2):240-246.
- Tokunaga T, Shimada K, Yamamoto K, et al. Retrospective analysis of prognostic factors for angioimmunoblastic T-cell lymphoma: a multicenter cooperative study in Japan. Blood. 2012;119(12):2837-2843.
- Moskowitz AJ, Lunning MA, Horwitz SM. How I treat the peripheral T-cell lymphomas. Blood. 2014;123(17):2636-2644.
- Lemonnier F, Couronne L, Parrens M, et al. Recurrent TET2 mutations in peripheral T-cell lymphomas correlate with TFH-like features and adverse clinical parameters. Blood. 2012;120(7):1466-1469.
- Cairns RA, Iqbal J, Lemonnier F, et al. IDH2 mutations are frequent in angioimmunoblastic T-cell lymphoma. Blood. 2012;119(8):1901-1903.
- Sakata-Yanagimoto M, Enami T, Yoshida K, et al. Somatic RHOA mutation in angioimmunoblastic T cell lymphoma. Nat Genet. 2014;46(2):171-175.
- Cheminant M, Bruneau J, Kosmider O, et al. Efficacy of 5-Azacytidine in a TET2 mutated angioimmunoblastic T cell lymphoma. Br J Haematol. 2015;168(6):913-916.

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