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The peripheral T-cell lymphoma: can we pivot from the chemotherapy-predicated paradigm?

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Abstract

Peripheral T-cell lymphomas (PTCL) are uncommon and aggressive diseases that are difficult to study. Combination chemotherapy such as cyclophosphamide, doxorubicin, vincristine, and prednisone has been the mainstay of treatment for almost 30 years, but outcomes remain poor. The development of new targeted therapies is changing the landscape of how we treat patients with these difficult diseases. For instance, the addition of brentuximab vedotin to combination chemotherapy enhanced the outcomes in patients with CD30-positive anaplastic large cell lymphomas, but there is still a need for better therapies in the other numerous subtypes. Here we discuss the data for the existing treatment paradigm of PTCL as well as the merits of shifting toward a chemotherapy-free approach.

Keywords: Peripheral T-cell lymphomas, PTCL, chemotherapy-free, epigenetics, targeted therapy

INTRODUCTION

Peripheral T-cell lymphomas (PTCL) are a group of diseases that are both very heterogeneous and rare, making it a significant challenge to advance care for patients affected by these diseases. The most recent World Health Organization (WHO) classification recognizes 29 distinct types, with the cell-of-origin of these non-Hodgkin lymphomas (NHL) arising from post-thymic T- and natural killer (NK) cells^[1]. Due to



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such a wide range in subtypes, clinical manifestations, and response to therapies, clinical trials are often limited by small numbers and lack of representation for the whole spectrum of disease. Adding to the complexity, prevalence also varies geographically, suggesting both genetic and environmental factors. In North America, the most frequent subtypes are PTCL not otherwise specified (NOS), angioimmunoblastic T-cell lymphoma (AITL), and ALK-positive anaplastic large cell lymphoma (ALCL), at 34%, 16%, and 16%, respectively^[2]. In Europe, the most diagnosed subtypes are PTCL NOS, AITL, and ALK-negative ALCL, making up 34%, 29%, and 9% of the cases, respectively^[3]. In Asia, however, the most common subtypes are adult T-cell leukemia/lymphoma (ATLL), PTCL NOS, and NK/T-cell lymphoma (NKTCL) at 25%, 22%, and 22% of PTCL cases, respectively^[3]. With the exception of ALK-positive ALCL, the PTCL carry a poor prognosis with the 5-year overall survival (OS) ranging from 14% in ATLL, 20% for enteropathy-type, 32% for AITL and PTCL NOS, to 59% in ALK-negative ALCL and 70% in ALK-positive ALCL^[3].

The poor outcomes are broadly related to intrinsic chemo-resistance, with at least 15%-20% of cases that are primary refractory to frontline treatment and short progression-free survival (PFS) in patients who do initially respond. Based on the very disappointing results of standard chemotherapy in the majority of PTCL subtypes, excluding ALK-positive ALCL, it should not come as a surprise that one additional novel agent combined with the chemotherapy backbone has often yielded disappointing results^[4-6]. In addition, retrospective data have demonstrated that patients who received novel drugs had improved outcomes when compared to those who were never treated with targeted agents^[7,8], further suggesting that combinations of novel agents could yield the future treatment for patients with PTCL. In this review, based on the intrinsic characteristics of these diseases, we present the merits of shifting to a chemotherapy-free paradigm.

THE MOLECULAR DRIVERS OF PTCL

In T-cell lymphomagenesis, the molecular pathways are poorly understood given the wide array of clinical and biologic presentations associated with so many subtypes, but despite these challenges, it has been possible to identify some recurring themes. We describe below the more prominent genetic and epigenetic aberrations described in PTCL NOS and AITL.

Mutation in epigenetic factors

Epigenetics refer to clonal changes in the pattern of gene expression without alterations in the primary genetic sequence. Over the last 10 years, it has become evident that PTCL are characterized by gross epigenetic dysregulation. The first suggestion that epigenetic pathways were relevant in the pathogenesis of PTCL comes from the clinic, where PTCL exhibit a consistent sensitivity to the histone deacetylase inhibitors (HDAC). PTCL are in fact the only malignant disease for which we have 4 HDAC inhibitors approved as monotherapy around the world. Only about thirty percent of patients can expect a response to HDAC inhibitors, but the duration of response that patients may experience is indeed remarkable. For example, cutaneous T-cell lymphomas (CTCL), for which HDAC inhibitors were first approved, are diseases currently being redefined and reclassified based on their genetic, epigenetic, and transcriptional changes landscape^[9,10].

The epigenetic machinery is responsible for maintaining the adaptive transcriptional memory of normal T-cells, and this improves the efficiency of T-helper subsets to transcribe inducible genes when needed^[11,12]. Dysregulation of epigenetic modulation likely contributes to the progression of normal T-cells in the development of PTCL, especially with recurrent reports of mutations in genes involved in methylation, acetylation, and other epigenetic functions^[13-16]. These mutations may lead to hypomethylation across the genome to silence tumor suppressor genes, especially in specific subtypes including AITL and PTCL with T-follicular helper (TFH) phenotype, which may be targeted with novel treatment paradigms.

Mutations in tet methylcytosine dioxygenase 2 (*TET2*), which encodes a methylcytosine dioxygenase that is important in normal cell function, are commonly reported in myeloid diseases but are also seen in lymphoid malignancies: 2% of B-cell lymphomas and almost 12% of PTCL including 33% of AITL^[17]. *TET2* mutations with and without DNA methyltransferase 3 alpha (*DNMT3A*) mutations can be found in non-tumor DNA in some PTCL patients, signifying that clonal hematopoiesis may contribute to pathogenesis^[18-20]. *TET2* mutations are common and may occur in almost half of AITL and over one-third of PTCL-NOS cases^[21]. A second study showed three-quarters of patients with AITL have mutations in *TET2*, with two-thirds of these cases with more than one mutation in the gene^[22]. Preclinical studies have demonstrated a causal relationship between mutations in *TET2* and the development of both lymphoid and myeloid cancers^[17]. Murine models with mutated *TET2* demonstrated TFH cell proliferation and progression to PTCL with TFH phenotype after a prolonged latency, possibly due to dysregulation of both cytokine expression and T-cell differentiation^[23,24]. Mutations in *TET2* have been found in patients without malignancies, related to an entity named clonal hematopoiesis of indeterminate potential (CHIP), which highlights that aberrancies in *TET2* are not enough for transformation but can predispose the cells to transform into diverse malignancies upon acquisition of secondary genetic hits.

DNA methyltransferase 3 alpha (*DNMT3A*) encodes a DNA methyltransferase and is found to be mutated in 10%-40% of AITL cases^[22]. Alterations in *DNMT3A* are thought to be the initial event in malignant transformation and often occur with mutations in *TET2*^[22,25]. Murine models demonstrate loss-of-function mutations in both *TET2* and *DNMT3A* lead to lymphomagenesis^[26].

Finally, isocitrate dehydrogenase-2 (*IDH2*) encodes a mitochondrial enzyme that is involved in 2-oxoglutarate generation. Mutations in *IDH2* at residue 172, where arginine is replaced by other amino acids, have been found in up to half of AITL cases^[27]. Mutant forms have altered enzymatic activity leading to the overproduction of 2-hydroxyglutarate, an oncometabolite that antagonizes the TET family of enzymes. Specifically, *IDH2* mutations lead to altered enzyme activity^[28] that catalyzes the conversion of α -keto glutarate (α KG) to 2-hydroxyglutarate (2HG). Elevated levels of intracellular 2HG interfere with the conversion of the DNA base 5-methylcytosine (5mC) into 5-hydroxymethylcytosine (5hmC), which is important in DNA demethylation. Interestingly, AITL is the only subtype of the T-cell lymphomas with mutations in *IDH2* and remarkably all the mutations occur at position R172, which is associated with abnormally high levels of 2HG. The mutation in *IDH2* (p.Arg172) leads to hypermethylation of gene promoters in a T-cell line^[29], consistent with a mechanism that is mediated by *IDH2* and *TET2* mutations, suggesting the cause of AITL pathogenesis. As seen with alterations in *DNMT3A*, mutations in *IDH2* appear to occur frequently with aberrancies in *TET2*^[22].

Mutations in RhoA

RHOA is part of the Rho family of GTPases and is important in signaling pathways, such as cellular responses and cell motility^[30-32]. The role of RHOA in tumorigenesis is controversial, but cell-type-specific functions of RHOA have recently been better defined by gene targeting techniques. Previously, there was little understanding of RHOA in cancer pathogenesis until *RHOA* mutations were repeatedly reported in AITL and PTCL NOS TFH subtypes^[14,33,34]. The *RHOA* mutations are unique to these subtypes of PTCL, where glycine at the 17th position is replaced by valine (G17V). *RHOA* G17V did not bind GTP like its normal counterpart, but rather, it inhibited the GTP binding to wild-type RHOA. The exact biochemical role of RHOA in T-cell signaling, however, is far from resolved. As described above, *TET2* is the most frequently mutated gene in AITL, and these tumors are commonly found to occur with *RHOA* G17V^[14,33]. Multiple transgenic murine models with *RHOA* G17V have demonstrated increased TFH proliferation as well as an overly reactive response to T-cell receptor (TCR) stimuli^[35,36]. Coupling *RHOA* G17V expression with *TET2* mutations resulted in the development of PTCL with TFH phenotype, i.e., AITL^[36,37]. Adding

evidence to the clonal hematopoiesis story, mice with defects in both *TET2* and *RHOA* developed myeloid tumors before seven months but tended to produce T-cell lymphomas after this time^[35]. One possibility, suggested by the experimental evidence described above, is that loss-of-function mutation in epigenetic factors, e.g., *TET2*, is an early event in the process of transformation that requires a second and final hit in *RHOA* G17V to induce cell differentiation toward TFH and the development of AITL.

VAV1 encodes a hematopoietic-specific Rho family-specific guanine exchange factor and is found to be mutated in up to 10% of AITL and PTCL-NOS patients^[38]. These mutations generally include recurrent frame deletion that is generated by alternative splicing and multiple *VAV1* gene fusions. Mutations in *RHOA* and *VAV1* are mutually exclusive, suggesting that they could be affecting the same genetic pathway^[39].

Mutations in T-cell receptor pathway genes

Defects in TCR signaling can result in aberrant activation, differentiation, and proliferation of T-cells. AITL or PTCL-NOS with a TFH phenotype specimens have recurrent mutations in T-cell receptor signaling including phospho-lipase C gamma 1 (*PLCG1*), CD28, phosphoinositide-3-kinase (*PI3K*) regulatory and catalytic subunits as well as other genes^[40]. Activating TCR alterations seemed to be adequate to drive malignant transformation since these cases rarely had more than one TCR gene mutation. TCR dysregulation was also associated with activation of NF- κ B, leading to cellular proliferation. CD28 mutations were reported in both AITL and PTCL NOS subtypes and was found to bind its ligand more tightly than wild-type CD28, which presumably is responsible for its activating ability^[41]. Interestingly, CD28-mutated AITL patients have worse survival compared to wild-type cases^[41]. The most common mutation in *PLCG1* p.Ser345Phe interferes with the enzymatic catalytic domain to increase NF- κ B/NFAT activity and, interestingly, is also found in CTCL, suggesting a shared mechanism of disease^[42].

Other insights into the genetic drivers of PTCL NOS

Consistent with the heterogeneous presentations of the PTCL, there is a variety of mutations reported to describe the diversity. In addition to the molecular drivers of disease described above, chromosomal translocations may offer insight into oncogenic pathways and potential therapeutic targets. The translocation t(5;9)(q33;q22) between interleukin-2-inducible T-cell kinase (ITK) and spleen tyrosine kinase (SYK) has been reported in 17% of PTCL NOS and 18% of PTCL NOS TFH patients^[43,44]. By immunohistochemistry, the majority (94%) of PTCL samples overexpress SYK^[45]. A mechanistic murine model demonstrated that overexpression of an ITK-SYK fusion gene is sufficient to activate TCR signaling and drive the production of T-cell lymphomas^[44,46]. Other novel kinase fusion genes have been described and warrant further investigation^[38]. Recent analysis of a large PTCL NOS cohort has identified additional altered genes, such as *YTHDF2* and *PD1*, and even more interestingly, a novel subtype characterized by *TP53* and/or *CDKN2A* mutations and deletions has been described and associated, not surprisingly, with a worse prognosis^[47].

In summary, the implementation of high-throughput genomic techniques in the field of PTCL has brought significant improvement in our understanding of the molecular complexity and genetic drivers of the disease. The identification of recurrently deregulated pathways has paved the way to the development of more rationally planned drug development and to the design of early phase clinical trials aiming to target these pathways that will hopefully yield better clinical outcomes for patients with PTCL.

STANDARDS OF CARE

Cyclophosphamide, doxorubicin, vincristine, and prednisone

Combination chemotherapy with cyclophosphamide, doxorubicin, vincristine, and prednisone (CHOP) is currently considered the standard of care based on a large, randomized phase 3 study led by Fisher *et al.* that explored the value of different intensified chemotherapy regimens compared to CHOP in patients with intermediate- and high-grade NHL that included both B-cell and T-cell lymphomas^[48]. End points of the study were overall response rate (ORR), time to treatment failure (TTF), disease-free survival (DFS), overall survival (OS), and incidence of severe or life-threatening toxicity. This landmark trial showed no difference in ORR, PFS, and DFS, but patients treated with CHOP developed fewer adverse events. At the time of the study, there was no standardized differentiation between aggressive B-cell and T-cell lymphomas, and CHOP chemotherapy has become the default standard of therapy for both subtypes for the past 30 years. PTCL patients treated with doxorubicin-based chemotherapy have a 3-year PFS of 32% and overall survival (OS) of 55%^[49]. Since then, there has been better characterization of lymphomas, e.g., diffuse large B-cell lymphoma (DLBCL) and addition of anti-CD20 monoclonal antibody rituximab with improvement in survival, but much less progress has been made for the T-cell lymphomas compared to the B-cell counterpart.

Cyclophosphamide, doxorubicin, vincristine, etoposide, and prednisone

Adding more chemotherapy to the CHOP backbone has been studied as a means to improve outcomes. In a secondary analysis of the German High-Grade NHL Study Group trials, the addition of etoposide to standard CHOP chemotherapy has been shown to improve the 3-year event-free survival (EFS) to 75% from 51% with CHOP alone ($P = 0.003$) in patients younger than 60 with normal lactate dehydrogenase though there was not a significant difference in OS^[50]. This finding is mostly because of treatment effect on patients with ALK-positive ALCL, where treatment with Cyclophosphamide, doxorubicin, vincristine, etoposide, and prednisone (CHOEP) was associated with a 3-year EFS of 91% compared to 57% when treated with CHOP ($P = 0.012$); the effect was no longer significant in other subtypes of PTCL^[50]. An additional retrospective study of Danish and Swedish patients with ALK-positive ALCL treated with CHOEP was associated with better PFS (hazard ratio [HR] 0.48, $P = 0.034$) and OS (HR 0.34, $P = 0.008$) compared to treatment with CHOP^[51]. In a meta-analysis performed in Asian patients, however, the addition of etoposide did not improve response rates when compared to CHOP^[52].

Infusional etoposide, doxorubicin, and vincristine with cyclophosphamide and prednisone

Given evidence of multidrug resistance by expression in relapsed and refractory lymphoma, prolonged exposure to doxorubicin and vincristine over a 4-day infusion may overcome resistance mechanism compared to the 1-day infusion used in CHOP^[53]. In the phase II study of dose-adjusted Infusional etoposide, doxorubicin, and vincristine with cyclophosphamide and prednisone (EPOCH), 41 previously untreated patients with PTCL (excluding ATLL) were treated, leading to an ORR of 78% (with 61% achieving a complete response [CR]). The most common grade 3 or higher adverse events included neutropenia (75%), anemia (41%), and thrombocytopenia (22%). After a median follow-up of 2 years, the median PFS and OS were not reached. The 2-year PFS and OS were 53% and 73%, respectively. The decision to choose CHOP, CHOEP, or EPOCH is usually dependent on the provider and/or institution.

Brentuximab vedotin plus CHP

Adding brentuximab vedotin (an antibody drug conjugate linking a monoclonal antibody against cluster of differentiation 30 (CD30) and cytotoxic agent, monomethyl auristatin E [MMAE], Bv) to cyclophosphamide, doxorubicin, and prednisone (Bv-CHP) compared to CHOP in the phase 3 frontline ECHELON-2 trial met its primary endpoint and was the first regimen to get FDA approval in frontline CD30-positive (defined as 10% of more expression by immunohistochemistry) PTCL^[6]. The median

progression-free survival (PFS) was 48.2 months and 20.8 months after a median follow-up of 36.2 months in the Bv-CHP and CHOP arms, respectively. The median OS was not reached for either group, but the hazard ratio was 0.66 (95%CI: 0.46-0.95), $P = 0.0244$. The ORR and CR rate for the Bv-CHOP and CHOP arms were 83% and 68% vs. 72% and 56%, respectively. Both groups had similar rates of adverse events (AE), grade 3 or higher AE, and serious AE. This is now the standard of care in patients with CD30-positive PTCL, though the trial was enriched for ALCL and not sufficiently powered for other subtypes. The 5-year update demonstrated that after a median follow-up of 47.6 months, the median PFS was 62.3 months vs. 22.8 months and median OS was not reached vs. 54 months in the Bv-CHP and CHOP arms, respectively^[54]. A multicenter study is now recruiting to study Bv-CHP in patients with untreated PTCL with less than 10% CD30 expression (NCT04569032).

Does adding a novel agent to CHOP-like regimens improve outcomes?

Save the addition of brentuximab vedotin in CD30-positive PTCL, the treatment for PTCL has not shared the same advancement of B-cell lymphomas with the addition of the CD20 monoclonal antibody rituximab despite many recent attempts to add additional targeted agents FDA-approved in the relapsed or refractory (R/R) setting to the CHOP-backbone in the past 5 years^[55].

Pralatrexate plus CHOP-like Combinations

Pralatrexate was the first drug approved for patients with relapsed or refractory PTCL. It is a folate analog that binds the reduced folate carrier (RFC), leading to the inhibition of dihydrofolate reductase (DHFR) intracellularly^[56]. PROPEL was an international single-arm Phase 2 study conducted in patients with R/R PTCL that led to the US FDA approval in 2009^[57]. Following cyclophosphamide, etoposide, vincristine, and prednisone (COEP) on day 1 of each cycle, pralatrexate was added on days 15, 22, and 29 in a phase 2 study to treat 33 patients with untreated PTCL^[58]. The ORR was 70% (CR 50%). After a median follow-up of 21.5 months, the estimated 2-year PFS and OS were 64% and 80%, respectively. Common grade 3 or 4 AE included anemia (27%), febrile neutropenia (18%), and mucositis (18%). This combination did not demonstrate improvement in outcomes seen with CHOP and was not deemed a regimen that warranted additional exploration by the authors.

The sequential addition of pralatrexate followed by COEP did not improve outcomes compared to CHOP, so the folate analog metabolic inhibitor was added to CHOP (Fol-CHOP) in a phase I study (NCT02594267)^[5]. The study was completed in October 2020, but the results are not readily available at the time of writing this manuscript.

Romidepsin plus CHOP (Ro-CHOP)

It has been a decade since romidepsin, a histone deacetylase (HDAC) inhibitor, was approved in some countries based on the phase II study in patients with R/R PTCL by Coiffier *et al.*^[59-61]. The phase III Ro-CHOP study (NCT01796002) comparing CHOP with CHOP plus romidepsin in patients with untreated PTCL, excluding ALK-positive ALCL, was a highly anticipated trial. Unfortunately, the clinical trial revealed that the addition of romidepsin did not improve ORR, PFS, or OS compared to CHOP alone^[4]. After a median follow up of almost 28 months, the study failed to meet its primary endpoint with a median PFS of 12.0 months compared to 10.2 months in the Ro-CHOP and CHOP groups, respectively. The hazard ratio of 0.81 was not significant ($P = 0.096$)^[4]. The secondary endpoints of overall response rate (ORR) and OS were also not statistically different, but the addition of HDAC inhibitor, however, was associated with more treatment-emergent adverse events related to cytopenias and gastrointestinal side effects. This led to the withdrawal of romidepsin's PTCL indication in August 2021, though appeals are pending.

Romidepsin was also added to CHOEP but the trial was stopped because there was no improvement in the primary endpoint of PFS^[62].

Azacytidine and CHOP

Given the enrichment of mutations related to the epigenetic machinery in AITL and PTCL NOS TFH, an oral hypomethylating agent (HMA), azacytidine was added to the CHOP backbone in a phase II study that prioritized PTCL TFH subtypes (NCT03542266). At a median follow up of 21 months, the CR was 75% overall ($n = 20$) and 88% in patients with TFH subtypes ($n = 17$)^[63]. The 2-year OS was 68% for the entire cohort and 76% for TFH subtypes, with an estimated median PFS of 36 months^[64]. In addition, the *TET2* mutations were associated with CR ($P = 0.007$), favorable PFS ($P = 0.004$), and OS ($P = 0.015$) in contrast to *DNMT3A* mutations, which were associated with a worse PFS ($P = 0.016$)^[63]. The biomarker approach may be a promising way to decide which patient gets the right treatment. Longer follow-up is needed to determine the impact on survival.

A phase II cooperative group study is testing the addition of duvelisib or azacytidine to CHOP with or without etoposide in patients with CD30-negative PTCL (NCT04803201).

Alemtuzumab plus CHOP

The phase 3 Act-2 trial added alemtuzumab (a monoclonal antibody targeting CD52) to CHOP in older patients with untreated PTCL and also failed to demonstrate improvement compared to CHOP alone^[64]. There was no statistically significant benefit in ORR, EFS, PFS, or OS, but there were higher rates of toxicity, including grade 3 or higher infections (40% vs. 21%)^[64]. The planned sample sizes for elderly and younger cohorts were not reached due to poor accrual.

Lenalidomide plus CHOP

Lenalidomide, an oral immune modulator that acts by inhibiting cereblon, was added to 8 cycles of CHOP chemotherapy as frontline therapy to treat older patients with AITL in a phase 2 trial^[65]. Patients also received 4 injections of intrathecal methotrexate during the study. The ORR was 56% (CR 41%), which was not better than CHOP alone in historic controls (e.g., ECHELON-2 study)^[6,65]. After a median follow-up of 45 months, the 2-year PFS and OS were 42% and 59%, respectively. The most common AE was neutropenia, with 71% of patients developing grade 4 neutropenia. In addition, 23% developed grade 3 or 4 infections and 5% succumbed to grade 5 sepsis. The authors did not believe the results warranted the addition of lenalidomide to CHOP in this subset of patients. Lenalidomide is being studied with CHOP-like therapies in different populations, both with CHOP (NCT04922567) and CHOEP in the frontline setting (NCT02561273) with results pending.

Nivolumab and EPOCH

A multicenter pilot study investigating nivolumab with EPOCH showed that 78% of 18 patients enrolled developed immune related adverse events (irAEs), and 44% required discontinuation of nivolumab due to irAEs^[66]. The trial protocol permitted patients to receive one cycle of chemotherapy prior to enrollment, and interestingly, patients who received one cycle of EPOCH prior to the addition of nivolumab to subsequent cycles did not experience dose-limiting irAEs. These data suggest pre-treatment with chemotherapy may result in less toxicity. Of note, the ORR was 83% at the end of induction and 2 patients proceeded with a consolidative autologous stem cell transplant^[66].

Ongoing “CHOP-like plus” Trials

There are still many studies looking to add targeted agents to combination chemotherapy and results are pending. In China, a phase 3 study comparing decitabine plus CHOP vs. CHOP is ongoing (NCT03553537).

The study began in 2015, was completed in 2020, and is pending analysis and publication of the results. Two phase II frontline studies investigate the safety and efficacy of adding chidamide, an HDAC inhibitor, or apatinib, a multitargeted kinase inhibitor, to CHOP in China (NCT03268889 and NCT03631862, respectively). [Table 1](#) summarizes completed and ongoing trials studying CHOP-like regimens with and without targeted therapies.

THE MANY NEW DRUGS COMING ALONG IN PTCL

Approved agents

About 70% of patients with PTCL treated with chemotherapy eventually will relapse or have refractory disease, and these patients have particularly poor outcomes. The British Columbia Cancer Agency Lymphoid Cancer found that patients with PTCL who relapse or are refractory to first-line therapy had a short median PFS of 3.1 months and median OS of 5.5 months^[67]. Over the past decade, the FDA approved pralatrexate in the United States in 2009, romidepsin for PTCL in the United States in 2011, mogamulizumab for ATLL in Japan in 2012, belinostat in the United States in 2014, brentuximab vedotin in CD30-positive ALCL and mycosis fungoides in the United States in 2014, chidamide in China in 2014 based on registration trials^[57,59,68-71]. Treatment with novel agents in the relapsed and refractory setting is associated with improved survival compared to additional, less efficacious chemotherapy^[7,8]. In August 2021, romidepsin was withdrawn from the indication, despite the efficacy of novel combinations. Over the past 5 years, pralatrexate has been approved in Japan and China, romidepsin, forodesine, and mogamulizumab were approved for the treatment of R/R PTCL in Japan, and numerous other targeted agents have been studied worldwide.

Pralatrexate

As previously mentioned, pralatrexate was the first novel agent approved in the treatment of R/R PTCL in 2009 based on the results from the phase 2 registration PROPEL study^[57]. The ORR in these patients was 29%, and treatment was associated with a median duration of response (DOR) of 12.4 months^[57]. In the past 5 years, pralatrexate has been studied around the world and has obtained additional FDA approvals in Asia. In two phase II registration studies in Japan and China, pralatrexate was studied in R/R PTCL, resulting in ORRs of 45%-52% (CR 9%-10%) in patients from both cohorts^[72,73]. The most common AEs included mucositis (68%-88%), thrombocytopenia (37%-68%), and liver function test abnormalities (41%-64%). Almost half of the patients (48%-49%) developed serious AEs, including febrile neutropenia and pneumonia. A study in Taiwan using pralatrexate in R/R PTCL is also ongoing (NCT03150602). Pralatrexate remains an important option in the treatment of these patients with R/R PTCL.

Romidepsin

Romidepsin was studied in patients with R/R T-cell lymphomas with ORR ranging from 25% to 34%, leading to its approval in R/R CTCL in 2009 and R/R PTCL in 2011^[59,74,75]. More recently, the HDAC inhibitor has been studied in 40 Japanese patients with R/R PTCL, leading to FDA approval in Japan^[76]. There were no unexpected AEs, and the most common grade 3 or higher AEs included lymphopenia (74%), neutropenia (54%), leukopenia (46%), and thrombocytopenia (38%). The ORR was 43% with 23% of patients achieving a CR, which is better than previous cohorts. In patients with the rarer subtype, extranodal NK/T-cell lymphoma (ENKTCL), however, treatment with romidepsin is associated with reactivation of EBV infection, likely due to changes in histone acetylation of genes that normally play a role in viral repression^[77]. Phosphodiesterase inhibitors have been shown to prevent romidepsin-induced EBV reactivation in preclinical studies that may lead to future studies^[78]. As mentioned above, however, romidepsin was unexpectedly withdrawn from the PTCL indication after the highly anticipated phase 3 R-CHOP trial failed to meet its primary endpoint, though combination studies are ongoing.

Table 1. CHOP plus trials in Untreated PTCL

Phase	Population	Study drugs	Primary endpoint	Result	Reference
3	Untreated CD30-positive PTCL	Bv-CHP vs. CHOP (ECHELON-2)	PFS, met	Median PFS Bv-CHP 48.2 months vs. CHOP 20.8 months, HR 0.71 $P = 0.0110$	[6]
2	Untreated PTCL with < 10% CD30 expression	Bv-CHP	ORR	Recruiting	NCT04569032
2	Untreated patients with PTCL-NOS, AITL, ALCL	Sequential COEP followed by pralatrexate	CR, met	CR 51% (goal CR > 40%), but 2-year PFS and OS are not better than historic controls	[58]
1	Untreated PTCL	Pralatrexate + CHOP	MTD	Study completed 8 Oct 2020, data unavailable	[5]
3	Untreated PTCL, excluding ALK-positive ALCL	Ro-CHOP vs. CHOP	PFS, not met	Median PFS Ro-CHOP 12.0 months vs. CHOP 10.2 months, HR 0.81 $P = 0.0962$	[4]
2	Untreated PTCL, excluding ALK-positive ALCL	Romidepsin plus CHOEP	18-month PFS, not met	18-month PFS 48% (goal 70%)	[62]
2	Untreated PTCL	Chidamide + CHOP vs. CHOP	CR	Results pending	NCT03268889
2	Untreated PTCL	Chidamide + CHOEP and ASCT	2-year PFS	Results pending	NCT02987244
2	Untreated PTCL	Chidamide + CHOPE vs. CHOPE	CR	Results pending	NCT03617432
2	Untreated PTCL, TFH prioritized	Azacytidine and CHOP	CR	CR overall 75%, PTCL-TFH 88% (goal N/A)	[63]
2	Untreated PTCL with < 10% CD30	Azacytidine or Duvelisib plus CHO(E)P	CR	Recruiting	NCT04803201
3	Untreated PTCL	Decitabine + CHOP vs. CHOP	3-year PFS	Results pending	NCT03553537
1/2	Untreated PTCL	Nivolumab + EPOCH	Safety, tolerability, ORR	Grade 3/4 IRAE 39%, ORR 83%	[66]
3	Untreated PTCL	Alemtuzumab + CHOP vs. CHOP	3-year EFS, not met	3-year EFS A-CHOP 32% vs. CHOP 25%, $P = 0.159$	[64]
2	Untreated older AITL patients	Lenalidomide and CHOP (REVAIL)	CMR, not met	CMR 41% (goal 55%)	[65]
1/2	Untreated PTCL	Lenalidomide and CHOP	MTD, safety, ORR	Recruiting	NCT04423926
2	Untreated PTCL	Lenalidomide + CHOP vs. CHOP	ORR	Recruiting	NCT04922567
2	Untreated PTCL	Biomarker targeted treatments plus CHOP vs. CHOP	CR	Recruiting	NCT04480099
3	Untreated PTCL	Apatinib + CHOP vs. CHOP	2-year PFS and ORR	Results pending	NCT03631862
2	Untreated PTCL	Tenalisib + CHOP	CR	Not yet recruiting	NCT05239910

AITL: Angioimmunoblastic T-cell lymphoma; ALCL: anaplastic large cell lymphoma; Bv-CHP: brentuximab vedotin, cyclophosphamide, doxorubicin, prednisone; CHOEP/CHOPE: cyclophosphamide, doxorubicin, vincristine, etoposide, prednisone; CHOP: cyclophosphamide, doxorubicin, vincristine, prednisone; CMR: complete metabolic response; CR: complete response; HR: hazard ratio; IRAE: immune related adverse event; ORR: overall response rate; MTD: maximum tolerated dose; N/A: not applicable; PFS: progression free survival; PTCL: peripheral T-cell lymphoma; PTCL NOS: peripheral T-cell lymphoma not otherwise specified; Ro-CHOP: romidepsin, cyclophosphamide, doxorubicin, vincristine, prednisone; TFH: T-follicular helper.

Belinostat

Belinostat is a hydroxamic acid-derived pan-HDAC inhibitor that was approved in 2014 for the treatment of R/R PTCL because of the phase 2 registration BELIEF trial^[69]. Patients enrolled in the study tolerated treatment with the most common grade 3 or 4 adverse events consisting of anemia (11%), thrombocytopenia (7%), neutropenia (6%), and dyspnea (6%)^[69]. Similar to other HDAC inhibitors, the ORR was 26% with a median DOR of 13.6 months. In a subset analysis, it was notable that in patients with AITL, the ORR was 46%, suggesting a higher sensitivity to this class of drug^[79].

Forodesine

Purine nucleoside phosphorylase (PNP) seems important in T-cell biology, and forodesine is a novel oral PNP inhibitor that has been approved in Japan for the treatment of R/R PTCL^[80]. Of 41 evaluable patients, the ORR was 25% (CR 10%). Given the mechanism of action, the AEs were expected with grade 3/4 lymphopenia (96%), leukopenia (42%), neutropenia (35%), and thrombocytopenia (25%). Five patients (3 with AITL, 2 with PTCL NOS) developed secondary B-cell lymphomas. The drug has not been approved elsewhere.

Mogamulizumab

Mogamulizumab is an antibody that targets C-C chemokine receptor 4 (CCR4) and was approved in Japan for R/R PTCL with CCR4 expression (mostly ATLL) almost a decade ago. In order to study other ATLL patient populations, a phase II international study was conducted in 22 centers to compare mogamulizumab and investigator's choice (IC): pralatrexate, gemcitabine plus oxaliplatin (GemOx), or dexamethasone, cisplatin, and cytarabine (DHAP)^[81]. The ORR was 11% (CR 2%) in the mogamulizumab arm compared to 0% in the IC arm. Though treatment was well tolerated, mogamulizumab did not improve outcomes for this difficult-to-treat group of patients. In addition, caution must be taken if mogamulizumab is given to patients prior to allogeneic stem cell transplant because there is an increased risk of graft-versus-host-disease (GVHD)-related mortality, which is normally tempered by CCR4-expressing Tregs that would be inhibited by the drug^[82].

Chidamide

In China, chidamide, another HDAC inhibitor, was approved for the treatment of R/R PTCL patients in 2014 based on the trial that produced an ORR of 29% with a median DOR of 10 months^[71]. There are many clinical trials studying chidamide-based combinations in the treatment of Chinese patients with PTCL.

Investigational novel agents

Given the need for more efficacious treatments in this difficult group of diseases, there are many new targeted agents being studied as single agents. Below we discuss data from investigational novel agents as single agents and summarize ongoing studies in [Table 2](#).

Azacytidine

The serendipitous discovery of the effectiveness of azacytidine in the treatment of AITL began with the treatment of 2 patients with both AITL and chronic myelomonocytic leukemia who achieved CR^[83,84]. Due to recurrent mutations in the epigenetic machinery in AITL and PTCL TFH, azacytidine is a logical choice of therapy and is associated with an ORR of 9/12 (75%) with 6/12 achieving CR^[85]. All patients had a *TET2* mutation, and some had additional mutations in *DNMT3A*, *IDH2*, and *RHOA*. The median PFS and OS were 15 months and 21 months, respectively. Since then, there have been many clinical studies testing azacytidine in combinations to treat patients with untreated and R/R PTCL.

Phosphatidylinositol 3-kinase (PI3K) inhibitors

Duvelisib, a dual PI3K- δ,γ inhibitor, has been studied in patients with R/R PTCL (NCT03372057) with treatment limited by low CD4 counts^[86]. Various doses were studied, but the patients enrolled in the expansion phase received oral duvelisib 75 mg twice daily for 2 cycles, followed by 25 mg twice daily until disease progression or toxicity^[87]. The ORR in 78 evaluable patients in the expansion phase was 50% (CR 32%) with a median DOR of 233 days. The most common grade 3 or 4 adverse events included neutropenia (38.5%), ALT and AST elevation (24.4%, 21.8%, respectively)^[87]. Of note, 3 deaths from pneumonitis, Epstein-Barr associated lymphoproliferative disorder, and sepsis occurred on trial, possibly due to treatment. Due to the risk of infections, trial investigators recommend antiviral prophylaxis to be

Table 2. Novel Therapies in Ongoing Clinical Trials

Phase	Patient population	Drug	Mechanism of action	NCT number
2	Untreated PTCL, ineligible for chemotherapy	Brentuximab vedotin, intravenous	Antibody-drug conjugate targeting CD30	NCT01716806
2b	R/R PTCL	HBI-8000, oral	HDAC inhibitor	NCT02953652
2	R/R PTCL	Valemetostat tosylate (DS-3201b), oral	Enhancer of zeste homolog 1/2 dual inhibitor	NCT04703192
2	R/R PTCL	TQ-B3525, oral	PI3K α/δ inhibitor	NCT04615468
2	R/R PTCL	YY-20394, oral	PI3K δ inhibitor	NCT04705090
2	R/R PTCL	AZD4205, oral	Janus kinase inhibitor	NCT04105010
2	R/R PTCL	AZD4573, intravenous	CDK9 inhibitor	NCT05140382
2	R/R PTCL	Tipifarnib, oral	Farnesyl protein transferase inhibitor	NCT02464228
2	R/R PTCL	F520, intravenous	Anti-PDL1 antibody	NCT04457830
2	R/R malignancies	STI-3031	Anti-PDL1 antibody	NCT03999658
1/2	R/R PTCL	AK104, intravenous	PD1/CTLA4 bispecific antibody	NCT04444141
1/2	R/R PTCL	Tolinapant (ASTX660), oral	Dual antagonist of cellular and X-linked inhibitors of apoptosis proteins	NCT04362007
1/2	R/R solid tumors or lymphomas with WT TP53	ALRN-6924	Dual MDMX and MDM2 inhibitor	NCT02264613
2	R/R PTCL	Fenretinide, intravenous emulsion	Synthetic retinoid derivative	NCT02495415
1	R/R PTCL	nanoFenretinide	Synthetic retinoid derivative	NCT04234048
1	R/R PTCL	CPI-818, oral	Interleukin-2-Inducible T-Cell Kinase Inhibitor	NCT03952078
1	R/R lymphoid malignancies	BP1002	L-Bcl2 antisense oligonucleotide	NCT04072458
1	R/R solid tumors and lymphomas	SEA-TGT	Inhibitory immune receptor	NCT04254107
1	R/R PTCL	Carfilzomib	Proteasome inhibitor	NCT01336920
1	R/R PTCL TFH	MEDI-570	Anti-ICOS monoclonal antibody	NCT02520791

BCL2: B-cell lymphoma 2; CDK9: cyclin-dependent kinase 9; CTLA4: cytotoxic T-lymphocyte-associated protein 4; HDAC: histone deacetylase; ICOS: inducible T cell costimulatory; MDM2: mouse double minute 2; MDMX: murine double minute X; PD1: programmed cell death protein 1; PDL1: programmed death ligand 1; PI3K: phosphoinositide 3-kinase; PTCL: peripheral T-cell lymphoma; R/R: relapsed or refractory; TFH: T-follicular helper.

administered with duvelisib. In another phase 1 study, Tenalisib, a dual PI3K δ/γ Inhibitor, was studied in 58 patients with R/R PTCL and CTCL (NCT02567656)^[88]. In all patients, the most common AE was fatigue (45%), AST elevation (36%), ALT elevation, and diarrhea (33%). The ORR in the PTCL subset was 46.7% (CR 20%). To increase efficacy, tenalisib is being studied in combination with romidepsin with preliminary data discussed below (NCT03770000).

Immune checkpoint inhibitors

Pembrolizumab is an anti-Programmed cell death protein 1 (PD1) monoclonal antibody and was studied in a phase II trial that resulted in an ORR of 33% in patients with R/R PTCL ($n = 15$)^[89]. Nivolumab, another drug in the same class, was studied in a phase II trial and resulted in an ORR of 33% in patients with R/R PTCL^[90]. There were reports of hyperprogression during treatment in 10 out of the 12 patients. Both trials were terminated early due to futility and the latter also cited safety concerns. Nivolumab was also associated with hyperprogression in patients with ATLL, likely due to effects on tumor-infiltrating Tregs after immune checkpoint inhibition^[91,92]. Due to concern for hyperprogression in ATLL, a phase 2 multicenter study in R/R PTCL, excluding ATLL, was performed and showed that anti-PD1 antibody geptanolimab was safe without report of hyperprogression^[93]. In this study by Shi *et al.*, patients with PDL1 expression of 50% or greater seemed to predict response to treatment with ORR of 53% compared PDL1 expression of < 50% with ORR of 25%^[93].

In R/R ENKTCL where there is high Programmed death ligand 1 (PDL1) expression (54% of cases), however, pembrolizumab is associated with an ORR of 6/14 (43%) to 7/7 (100%) with 2/7 (29%) to 5/7 (71%) achieving a CR^[94-96]. Given the expression of PDL1 in ENKTCL, avelumab, an anti-PDL1 monoclonal antibody, has been used in a phase 2 trial with R/R ENKTCL^[97]. Out of 21 patients, the ORR was 38% (CR 24%) with a median PFS of 2.7 months after a median follow-up of 15.7 months. In this patient population, the trial was terminated because of lower response rate than expected. Avelumab is also being studied alone in a phase 2 trial in R/R PTCL (NCT03046953) as well as in a phase 1 trial in combination with interleukin-15 (NCT03905135).

Immunotherapy agents have been studied in combination to treat PTCL without cases of hyperprogression, warranting further study as later discussed below.

Cerdulatinib

Spleen tyrosine kinase (SYK) and Janus kinase (JAK) are enzymes important in cellular processes and have been found to be important in T-cell lymphomagenesis^[98]. Cerdulatinib is a SYK, JAK1, and JAK3 inhibitor that is being studied in R/R lymphomas, including PTCL (NCT01994382). 65 patients with R/R PTCL were treated with cerdulatinib in a phase 2 study, resulting in an ORR of 36.2% in 58 evaluable patients^[98]. Of note, the AITL/TFH subtype had a higher ORR of 51.9% (CR 37%) with a median DOR of 12.9 months after a median follow-up of 16.4 months. The most common treatment-emergent grade 3 or higher adverse events were elevation in amylase (23%), anemia (20%), elevation in lipase (18.5%) as well as neutropenia (12%) and sepsis (9.2%)^[58]. As we understand disease biology, we can better tailor treatments to optimize efficacy while limiting side effects.

Alisertib

Alisertib is an aurora kinase inhibitor that was randomized against investigator's choice of pralatrexate, romidepsin, and gemcitabine in the phase III LUMIERE trial in patients with R/R PTCL^[99]. Surprisingly, the ORR and CR rates achieved with alisertib compared to comparators were 33% and 18% vs. 45% and 27%, respectively. The median PFS and OS did not differ significantly between the two groups.

Valemetostat

Given known epigenetic dysfunction found in PTCL, valemetostat, a selective inhibitor of enhancer of zeste homolog 1 and 2 (EZH1/EZH2), has been studied in R/R PTCL patients^[100,101]. EZH1 and EZH2 catalyze methylation of histone 3 to downregulate gene expression, and dysregulation of the delicate balance of tumor suppression and cell processes may lead to lymphomagenesis^[101]. Valemetostat is given orally once a day until disease progression or toxicity, leading to an ORR of 48%^[101]. Longer follow-up is needed for these ongoing studies.

NOVEL DRUG COMBINATIONS: COULD THEY FORM THE BASIS OF NEW TREATMENT PLATFORMS?

The track record of adding targeted therapies to CHOP-based chemotherapy has so far mostly been disappointing, and the field is looking to study novel combinations in the treatment of both untreated and R/R PTCL. We discuss the combinations where data are available and list other ongoing trials with novel combination in [Table 3](#).

Azacytidine and Romidepsin

Preclinical data by Marchi *et al.* show that the combination of HDAC inhibitors and HMAs are synergistic in modulating gene expression leading to the rationale development of this regimen in patients with PTCL^[102,103]. Twenty-three patients (10 untreated, 13 R/R) were evaluable in the phase 2 trial with oral

Table 3. Novel combinations in ongoing trials

Phase	Patient population	Drugs	Mechanisms of action	NCT number
2b	R/R PTCL	Azacytidine and romidepsin vs. investigator's choice	HMA and HDAC inhibitor vs. approved agents	NCT04747236
2	R/R PTCL	Sintilimab and chidamide	PD1 antibody and HDAC inhibitor	NCT04512534
2	R/R PTCL	Sintilimab, chidamide and azacytidine	PD1 antibody, HDAC inhibitor, HMA	NCT04052659
1	R/R NHL	Carfilzomib and belinostat	Proteasome inhibitor and HDAC inhibitor	NCT02142530
2	R/R PTCL	Nivolumab and cabrializumab	PD1 antibody and anti-CSF-1R antibody	NCT03927105
2	R/R PTCL	Ixazomib and romidepsin	Proteasome inhibitor and HDAC inhibitor	NCT03547700
1/2	R/R PTCL	Romidepsin and carfilzomib	HDAC inhibitor and proteasome inhibitor	NCT03141203
2	R/R PTCL	Brentuximab vedotin and pembrolizumab	Antibody-drug conjugate targeting CD30 and PD1 inhibitor	NCT04795869
2	R/R PTCL	Chidamide and lenalidomide	HDAC inhibitor and immune modulator	NCT04329130
2	Untreated PTCL	Azacytidine and chidamide	HMA and HDAC inhibitor	NCT04480125
2	Untreated PTCL	Romidepsin and lenalidomide	HDAC inhibitor and immune modulator	NCT02232516
1	R/R PTCL	Bortezomib and azacytidine	Proteasome inhibitor and HMA	NCT01129180
1/2	R/R PTCL	Pembrolizumab and pralatrexate	PD1 inhibitor and antifolate	NCT03598998
1/2	R/R PTCL	Durvalumab and lenalidomide	PDL1 inhibitor and immune modulator	NCT03011814
2	R/R PTCL	Venetoclax and romidepsin	BCL2 inhibitor and HDAC inhibitor	NCT03534180
1/2	R/R PTCL	Pembrolizumab and romidepsin	PD1 inhibitor and HDAC inhibitor	NCT03278782
1/2	R/R PTCL	Pralatrexate and romidepsin	Antifolate and HDAC inhibitor	NCT01947140
1	R/R PTCL	Brentuximab vedotin and lenalidomide	Antibody-drug conjugate targeting CD30 and immune modulator	NCT03302728
1	R/R PTCL	Romidepsin, azacytidine, and lenalidomide	HDAC inhibitor, HMA, immune modulator	NCT04447027
2	R/R EBV-positive lymphomas	Nanatinostat and valganciclovir	HDAC inhibitor and antiviral	NCT05011058
1	R/R CD30-positive lymphomas	AFM13-NK and AFM13	Modified immune cells and bispecific antiCD30 and CD16A antibody	NCT04074746
1	R/R PTCL	Ruxolitinib and duvelisib	JAK inhibitor and PI3K inhibitor	NCT05010005

BCL2: B-cell lymphoma 2; HDAC: histone deacetylase; HMA: hypomethylating agent; JAK: Janus kinase; NHL: non-Hodgkin lymphoma; PD1: programmed cell death protein 1; PI3K: phosphoinositide 3-kinase; PTCL: peripheral T-cell Lymphoma; R/R: relapsed or refractory.

azacytidine and romidepsin^[103]. Though all subtypes were enrolled, patients with AITL or PTCL TFH ($n = 15$) had the best response compared to other subtypes ($n = 8$) with ORR 80% (CR 60%) vs. 25% (12.5%), respectively. The median PFS and OS for AITL or PTCL TFH compared to other subtypes was 8.9 months vs. 2.3 months and not reached vs. 9.4 months, respectively. The most common grade 3 or higher AEs included thrombocytopenia (48%), neutropenia (40%), and lymphopenia (32%). Most patients had mutations in epigenetic modifiers and a correlation between mutations and response was limited due to sample size. An ongoing multicenter phase 2 randomized trial comparing oral azacytidine and romidepsin vs. investigator's choice in R/R PTCL will confirm these findings (NCT04747236).

Pralatrexate and Romidepsin

In a murine model of T-cell lymphoma, the administration of pralatrexate in conjunction with romidepsin demonstrated synergy in treatment efficacy than either agent alone^[104]. To further explore the safety and efficacy of this combination, patients with R/R lymphomas enrolled in a phase 1 study of pralatrexate and romidepsin^[105]. The most common grade 3 and 4 AEs included anemia (29%), thrombocytopenia (28%), febrile neutropenia (14%), and oral mucositis (14%). The ORR for PTCL compared to non-PTCL were 71% (CR 40%) vs. 33% (CR 0%), respectively. Enrollment of patients with R/R PTCL in the phase 2 trial is ongoing (NCT01947140).

Tenalisib and Romidepsin

Tenalisib, a selective PI3K and SIK3 inhibitor, was studied in combination with romidepsin in 33 patients with R/R T-cell lymphomas^[106]. In the phase 1/2 trial, there were no dose-limiting toxicities, but the most frequent grade 3 or 4 treatment-emergent adverse events included thrombocytopenia (21%), ALT elevation (18%), and neutropenia (15%). The ORR was 63% in 27 evaluable patients (CR 26%), and 3 out of 6 PTCL patients who achieved a CR were bridged to transplant^[106].

Immune Checkpoint Inhibitors Combined with Epigenetic Modifiers

To improve on the results of the doublets: romidepsin plus azacytidine and romidepsin plus pralatrexate, two innovative trials are combining immune checkpoint inhibitors with epigenetic modifiers^[107]. In the phase 1 EMBOLDEN trial, pembrolizumab, an anti-PD1 monoclonal antibody, is combined with decitabine and/or pralatrexate and has resulted in an ORR of 2/13 (CR 1/13). Durvalumab in different combinations with pralatrexate, azacytidine, and romidepsin in the phase I/IIa DURABILITY trial has promising results with an ORR of 3/5 (CR 3/5). These trials are ongoing and actively enrolling patients (NCT03240211, NCT03161223).

CHIMERIC ANTIGEN RECEPTOR (CAR) T CELL THERAPY

CAR T cell therapy has been approved for B-cell malignancies and is currently in development for T-cell malignancies. Due to the genetic engineering of T-cells to target neoplastic T-cells, several considerations have arisen, such as T-cell aplasia, fratricide, and the contamination of modified T-cells with neoplastic T-cells that can lead to resistance^[108]. Targets have included CD4, CD5, CD7, CD30, and TCR beta chain-1 (TRBC1) have progressed to clinical trials as summarized in [Table 4](#). Results in patients are limited. LB1901 is a CAR T cell therapy targeting CD4, for example, has resulted in low CD4 T-cell counts, leading to a clinical hold^[109]. A phase I trial using autologous CD5 CAR T-cells to bridge 9 patients with R/R T-cell malignancies to allogeneic transplant resulted in ORR of 44% without T-cell aplasia^[110]. CD30-targeted CAR T cell therapies have been studied in ALCL where of 2 patients treated, 1 had a CR and 1 died of progression^[111]. This is a promising modality for treating patients with R/R disease.

CONCLUSION

The future of PTCL care could look like a precision oncology-based approach where the biology of the disease is targeted for optimal efficacy and limited side effects. Combination chemotherapy has been the default standard of care, but the minority of patients are cured and there are lasting side effects that may impact quality of life. Rather than categorizing patients with PTCL, the future may focus on subtypes to make more appropriate application of clinical trials rather than extrapolating to all patients, even when rarer subtypes are not represented. Patients with ATLL have a range of outcomes based on the subtype of disease, but more treatments need to be developed for the aggressive acute and lymphomatous types. HDAC inhibitors are approved for R/R PTCL but can be associated with hyperprogression in ATLL, highlighting that caution is needed when generalizing clinical trials. Historically, patients with ALCL are associated with better outcomes and CD30-targeted therapy have improved outcome, but more options are needed in those who relapse. Given the enrichment of epigenetic dysregulation as pathogenesis, AITL and TFH are associated with better outcomes when treated with hypomethylating agents and HDAC inhibitors, and there are ongoing trials around the world that are building on the success of these doublets. Patients with ENKTCL have aggressive disease that have found success with immunotherapy in the R/R setting, which can be a different story for patients with nodal PTCL. Nivolumab may be associated with hyperprogression as a single agent, and yet, it has also been found to be efficacious when combined with other drugs, both chemotherapy and targeted agents. CAR T cell therapy offers another modality of treating patients with R/R disease but is still early in development.

Table 4. CAR T cell therapies in ongoing trials

Phase	Patient population	CAR T	Target antigen	Costimulatory domain	NCT number
1	R/R PTCL or CTCL with CD4 expression	LB1901 (Legend Biotech, on clinical hold)	CD4	4-1BB	NCT04712864
1	R/R CD5 positive lymphomas	CT125A (Huazhong)	CD5	Unknown	NCT04767308
1/2	R/R CD5 positive PTCL	MT-101 (Myeloid)	CD5	Unknown	NCT05138458
1	R/R T-cell leukemia or lymphoma with CD7 expression	CD7.CAR/CD28zeta (Baylor)	CD7	CD28	NCT03690011
1	R/R CD7 positive hematologic malignancies	PG-CART-07-001 (PersonGen)	CD7	Unknown	NCT04480788
1	R/R T-cell hematologic malignancies with CD7 expression	SHYSXY-202105-CD7-CART (Shanghai)	CD7	Unknown	NCT05290155
1/2	R/R CD7 positive leukemia and lymphoma	anti-CD7 CAR-pNK cells (PersonGen)	CD7	Unknown	NCT02742727
1	R.R CD30 positive lymphomas	Fully human anti-CD30 CAR (National Cancer Institute)	CD30	CD28 ^[113]	NCT03049449
1/2	R/R CD30 positive lymphomas	CD30.CAR T cells (Baylor)	CD30	CD28 ^[114]	NCT02690545, NCT02917083
2	R/R CD30 positive lymphomas	ATLCAR.CD30 T cells (North Carolina)	CD30	CD28	NCT04083495
1	R/R PTCL with CD30 expression	CD30.CAR-T (Tessa)	CD30	Unknown	NCT04526834
1	R/R CD30 positive lymphocyte malignancies	WHUH-CART-CD30-01 (Wuhan)	CD30	Unknown	NCT04008394
1	R/R CD30 positive	ARM002-CN-2021-01 (Zhejiang)	CD30	Unknown	NCT05208853
1	CD70 positive malignant hematologic diseases	CD70 CAR T-cells (Zhejiang)	CD70	Unknown	NCT04662294
1/2	R/R PTCL with TRBC1 expression	AUTO4, RQR8/aTRBC1 (Autolus)	TRBC1	N/A	NCT03590574
1	R/R TRBC1 positive T-cell hematologic malignancies	SHYSXY-202101-CART (Shanghai)	TRBC1	N/A	NCT04828174

CTCL: Cutaneous T-cell lymphoma; PTCL: peripheral T-cell lymphoma; R/R: relapsed or refractory; TRBC1: T-cell receptor beta constant 1.

The PTCL are rare and heterogeneous, making it difficult to study but many advances, such as technology, diagnostics, and treatments, have been made since 1993, when CHOP was determined to be the standard. Technology permits interconnectedness in a global scientific community where communication across the world can be instant and international, multicenter clinical trials are more feasible than ever before. In addition, the diagnose of lymphomas has become more sophisticated. The Non-Hodgkin's Lymphoma Pathologic Classification Project from 1982 was used to classify patients with low, intermediate, and high-grade lymphomas^[112]. In the trial by Fisher and colleagues, patients with aggressive lymphomas were included and there was no differentiation between B-cell and T-cell^[48]. In this trial, subtypes from this working classification, subtypes encompassed both intermediate- and high-grade groups: follicular lymphoma that was predominantly large cell, diffuse small-cleaved cell, diffuse mixed small and large cell, diffuse large cell, immunoblastic, and small noncleaved^[89,112]. If these lymphomas were diagnosed today, the revised 2016 WHO classification could categorize these intermediate and aggressive lymphomas in 50 different ways^[1]. Finally, the development of novel agents has transformed how to diagnosis and treat lymphomas. In the past, there were only combinations of chemotherapy, and maybe the specific subtype would not change how patients could be treated. Now there are novel agents that specifically target disease pathways and specifying the subtype of lymphoma may change management.

International, multicenter trials are easier to coordinate than ever before, and global consortiums are important in systematically designing trials that answer important questions in the treatment of patients with PTCL. Given the difficulty of enrolling patients in clinical trials, which is also coupled with the fact that

at any given moment, there are numerous competing trials, collaborations to enroll patients on trials specific for their disease subtype should be prioritized. Drug development for orphan diseases has drawbacks and challenges as often the number of patients limits impact, but FDA pathways to prioritize will continue to help support these efforts.

DECLARATIONS

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Authors' contributions

Made substantial contributions to conception and design of the study and performed data analysis and interpretation: Ma H, Marchi E

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Ethical approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

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