

Treatment of Richter's syndrome

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Richter's syndrome (RS) is an aggressive histologic transformation of chronic lymphocytic leukemia (CLL), most commonly to diffuse large B-cell lymphoma (DLBCL). Outcomes are generally poor, with complete remission (CR) rates of only about 20% and less than 20% long-term survival with chemoimmunotherapy (CIT). RS is biologically heterogeneous, and in 80% of patients with CLL who develop DLBCL, the disease is clonally related to the CLL. Clonally unrelated cases are genetically and immunologically distinct from clonally related DLBCL-RS, have more favorable responses to CIT, and are best treated as de novo DLBCL. Relatively favorable outcomes with CIT are also seen in patients who have never previously received treatment for CLL and who lack *TP53* mutation or deletion. For the remaining patients, treatment on a clinical trial is optimal. Fortunately, numerous agents are now in clinical development that show encouraging results. Here we review clinical data for some of the most promising approaches. DLBCL-RS tumor cells frequently express programmed cell death 1 protein (PD-1), and several studies have demonstrated activity for PD-1 inhibitors, especially in combination with ibrutinib. The BCL2 inhibitor venetoclax in combination with R-EPOCH CIT achieved CR in 50% of patients, and a study of venetoclax-R-CHOP is ongoing. The noncovalent Bruton's tyrosine kinase inhibitor pirtobrutinib has achieved responses in approximately two-thirds of heavily pretreated patients and, given its favorable toxicity profile, appears ideally suited to combining with other active agents. Finally, we review available data for bispecific antibodies, antibody-drug conjugates, and chimeric antigen receptor T-cell therapy, which, after revolutionizing the treatment of DLBCL, are now being evaluated in RS.

LEARNING OBJECTIVES

- Understand the clinical features, diagnosis, and prognostic features of Richter's syndrome
- Understand the novel therapeutic approaches and how to select the optimal approach

CLINICAL CASE

The patient, a fit 73-year-old woman with Richter's syndrome (RS), was treated for chronic lymphocytic leukemia (CLL) in 2011 with 6 cycles of fludarabine, cyclophosphamide, and rituximab (FCR) chemoimmunotherapy (CIT). She achieved complete remission (CR) but progressed with CLL in 2017. At progression, genomic evaluation revealed unmutated *IGHV* with *IGHV4-39-IGHJ5* (subset no. 8) utilization. A chromosomal analysis revealed a diploid karyotype, with trisomy 12 identified in 34% of interphases by fluorescence in situ hybridization (FISH). Next generation sequencing showed a *NOTCH1* mutation. In 2018 she was enrolled in a clinical trial with 2 years of fixed-duration ibrutinib and venetoclax, achieving CR with undetectable minimal residual disease to a level lower than 10^{-4} in the bone marrow. In 2021, just over a year after completion of ibrutinib and venetoclax, she progressed with rapid nodal enlargement, splenomegaly, and constitutional

symptoms. Positron emission tomography/computed tomography (PET/CT) showed widespread nodal disease, with a maximum standardized uptake value (SUV) of 10.9. Biopsy confirmed diffuse large B-cell lymphoma (DLBCL). Genotyping of the biopsy specimen showed clonally related disease, with identical *IGHV4-39-IGHJ5* utilization; no analyzable metaphases for karyotyping; trisomy 12 by FISH; *NOTCH1* mutation; no *TP53* mutation. Analysis of untransformed CLL cells at the time of progression showed trisomy 12 with a complex karyotype (46,XX,add(1(p36.1),-7,add(7)(q22),-10,+12,-14,+2mar).

Diagnosis

RS affects 2% to 15% of CLL patients, with an incidence of 0.5% to 1% per year.^{1,2} RS is defined as the development of a histologically aggressive lymphoma in a patient with a previous or concurrent diagnosis of CLL/small lymphocytic lymphoma, most commonly (~90% of cases) DLBCL, which

in 80% of cases is clonally related to the underlying CLL. Transformation to classical Hodgkin lymphoma (CHL; ~10% of cases) or rare lymphoma subtypes occurs less frequently.³ The bulk of this review focuses on the DLBCL subtype (DLBCL-RS).

Clinical features

Suggestive clinical features of DLBCL-RS are high-grade fevers, rapidly enlarging lymph nodes, unexplained weight loss, hypercalcemia, markedly elevated lactate dehydrogenase (LDH), and the development of extranodal disease. These features are nonspecific, and biopsy is always required for diagnostic confirmation and to obtain tissue for genomic evaluation, which is prognostically relevant and may direct therapeutic decisions.¹

PET/CT and biopsy

We perform PET/CT in any patient with clinical suspicion for RS. In 1 study done prior to the targeted-agent era, the negative predictive value of a PET/CT maximum SUV (SUV_{max}) lower than 5 was 92%, with a positive predictive value of 38%.⁴ The positive predictive value of an SUV_{max} of 5 was lower in patients progressing on Bruton's tyrosine kinase inhibitors (BTKis) or PI3K-delta inhibitors, but an SUV lower than 5 retains excellent negative predictive value.⁵

We generally obtain a tissue diagnosis in patients with an SUV greater than or equal to 5 on PET/CT. However, there is considerable nuance to this decision, and clinical judgment can be exercised regarding features that increase or decrease the likelihood of RS being present. As an example, a patient with widespread lymph node progression clinically may be less likely to have RS than one with dominant and rapid progression at 1 or few sites. Similarly, significant differences in the SUV_{max} from 1 tumor site to another may suggest transformation at the site with a high SUV, while a patient with uniform SUVs of just above 5 throughout all lymph node groups may be more likely to have CLL progression. Overall, if there is suspicion, a biopsy should be considered. Excisional lymph node biopsy is ideal.¹ If the lesion is inaccessible for surgical biopsy, then an image-guided core needle biopsy (not fine needle aspiration) should be performed. Where possible, the lesion with the highest SUV should be biopsied.

Risk factors, molecular pathogenesis, and prognosis

High-risk genomic characteristics of CLL increase the risk of transformation to RS—notably, unmutated *IGHV* status, *IGHV* stereotyped subset number 8 (*IGHV4-39-IGHJ5*), activating *NOTCH1* mutations, *TP53* deletion and/or mutation, and *del11q*.^{1,3} Near tetraploidy has been associated with a high risk of RS in patients receiving ibrutinib.⁶

The clonal relationship between the CLL cells and the RS cells is determined by sequencing the *IGH* gene in the transformed cells and comparing it to a concurrent or historical *IGH* sequence from the patient's CLL cells. Discordant light-chain expression between the CLL and the DLBCL-RS cells is, unfortunately, not a reliable surrogate for the lack of a clonal relationship, with cases reported in which light-chain discordance between CLL and RS cells exists despite identical *IGH* rearrangements.⁷ Given the distinct genomic characteristics and clinical outcomes of clonally related vs clonally unrelated cases, these are best thought of as distinct entities.

Clonally related DLBCL-RS is genomically distinct from clonally unrelated DLBCL-RS. The most common genomic alterations

in clonally related DLBCL-RS are *TP53* mutation (60%-80%),⁸ *CDKN2A* deletion (30%), *MYC* overexpression (40%), and activating *NOTCH1* mutation (~30%).⁹ *NOTCH1* mutations cluster among patients with trisomy 12 and are largely mutually exclusive with *TP53* mutations and *CDKN2A* deletions.⁹ Notably, clonally unrelated DLBCL-RS has a lower rate of *TP53* disruption (~20%), akin to de novo DLBCL. Additionally, stereotyped immunoglobulin genes (particularly *IGHV4-39/IGHD6-13/IGHJ5*) are found in 50% of clonally related DLBCL-RS but almost never in clonally unrelated DLBCL-RS.⁸

Finally, DLBCL-RS appears immunologically distinct from both CLL and from de novo DLBCL. Notably, the malignant B cells in DLBCL-RS express programmed cell death 1 protein (PD-1) in up to 80% of cases, while PD-1 expression is rare in de novo DLBCL, which may have therapeutic relevance.¹⁰

The risk of RS most likely relates to underlying disease biology, rather than treatment received. Among numerous randomized trials in the front-line and relapsed/refractory settings reviewed here,³ none showed a significant difference in RS incidence between treatment arms, except a lower rate for FCR vs FC in the CLL8 study.¹¹ Notably, there was no difference in RS risk between treatment arms in the E1912 study of ibrutinib-rituximab vs FCR or the CLL14 study of chlorambucil-obinutuzumab vs venetoclax-obinutuzumab.^{12,13} The risk for RS, however, increases in studies in relapsed/refractory CLL compared to studies conducted in front-line patients, likely due to higher proportions of patients in relapsed/refractory studies with high-risk disease biology and clonal evolution during therapy.

Prognostic features

Patients untreated for CLL at the time of DLBCL-RS have relatively favorable progression-free survival (PFS; median 46.3 vs 7.8 months) (Figure 1A).¹⁴ Beyond this, the most important prognostic factor in DLBCL-RS is the clonal relationship of DLBCL-RS to the underlying CLL. In 1 study, patients with clonally unrelated disease had a median PFS of 62.5 months vs 14.2 months for clonally related disease (Figure 1B).⁸ Beyond clonal relationship to the underlying CLL (which is not available in most case series reported) and number of prior therapies for CLL, there is some discordance between different case series regarding the prognostic significance of certain markers. In the 2011 case series by Rossi et al, patients with *TP53* disruption had a median PFS of 9.4 months vs 47.1 months without *TP53* disruption (Figure 1B). In the Mayo Clinic series, however, *TP53* disruption was not independently associated with inferior outcomes, with the most important negative prognostic markers being elevated LDH and prior treatment for CLL. In another study, a complex karyotype in the CLL cells and an increasing number of prior therapies for CLL were associated with inferior PFS/OS, while LDH and *TP53* deletion by FISH were not.

Variants

The CHL subtype of RS (CHL-RS) is rare (~1/10th as frequent as DLBCL-RS) but is the second most common histologic transformation. A retrospective analysis of 94 patients demonstrated relatively favorable outcomes, especially for those patients who received chemotherapy with doxorubicin hydrochloride, bleomycin sulfate, vinblastine sulfate, and dacarbazine (ABVD; median OS, 13.2 years).¹⁵ Similar to data for DLBCL-RS, patients with no prior therapy for CLL had superior outcomes to previously treated

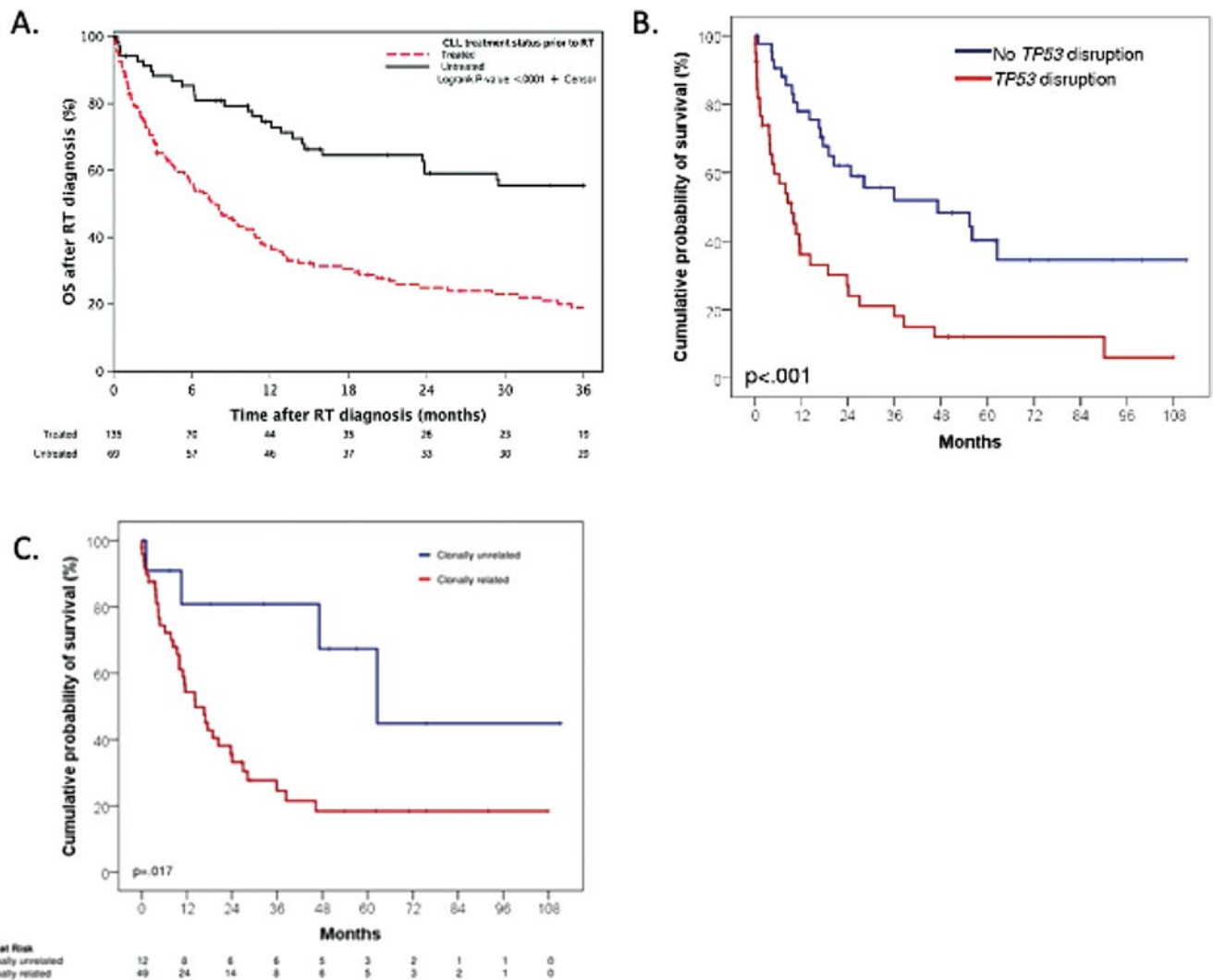


Figure 1. (A) PFS in Mayo Clinic patients with DLBCL-RS by prior CLL treatment status. Used with the permission of Wang et al.¹⁴ (B) PFS in patients in a case series according to the presence or absence of T53 mutation/deletion. (C) PFS of patients in a case series by clonal relationship to the underlying CLL. (B) and (C) used with the permission of Rossie et al.⁸ RT, Richter's transformation.

patients, and those never treated with purine analogue chemotherapy had superior outcomes to those previously treated with purine analogues. Few patients underwent allogeneic stem cell transplant (alloSCT) in first remission, but within the limitations of the small numbers of patients analyzed, alloSCT did not appear to have an impact on survival outcomes.

CIT in the treatment of DLBCL-RS

Initial treatment with CIT, analogous to the treatment of de novo DLBCL, is generally given, but outcomes for patients with DLBCL-RS are poor. CIT studies have generally resulted in CR rates of 20% or lower, with a median overall survival (OS) of 6 to 12 months (Table 1).

As outlined above, attempts to improve outcomes through intensification of chemotherapy have been unsuccessful. As a result, there is no standard of care CIT regimen for DLBCL-RS, and there is an unmet need for effective treatments in this disease. Currently, treatment choice is based on age, comorbidities,

prior therapies, and the experience of the treating center. Given poor outcomes with standard therapy, all patients should be treated in clinical trials where possible. Combinations with novel targeted therapies, checkpoint inhibitors, cellular therapy, and trials of several nonchemotherapy regimens are ongoing, as described below.

Hematopoietic progenitor cell transplantation in DLBCL-RS

Evaluation of the impact of alloSCT and autologous stem cell transplant (autoSCT) is limited by the lack of prospective randomized studies, introducing selection bias into the comparison of survival between transplanted and nontransplanted patients. With these caveats, there appears to be curative potential of alloSCT, especially for patients who achieve a CR prior to transplant. In a European Society for Blood and Marrow Transplantation analysis of 25 patients (9/25 with progressive disease prior to alloSCT), there was a plateau on the long-term relapse-free survival (RFS) curve.

Table 1. Outcomes in DLBCL-RS with standard CIT

Study and years of patient recruitment	Regimen	n	Median age (years)	Results		
				ORR	CRR	Median OS
Anthracycline-containing regimens						
Langerbeins et al ¹⁶ (2003–2008)	R-CHOP	15	69 (N/A)	67%	7%	21 months
Dabaja et al ¹⁷ (published 2000)	HyperCVXD	29	61 (36–75)	41%	38%	10 months
Tsimberidou et al ¹⁸ (1999–2001)	Rituximab and GM-CSF with alternating hyperCVAD and MTX/cytarabine	30	59 (27–79)	43%	18%	8.5 months
Rogers et al ¹⁹ (2006–2014)	R-EPOCH	46	67 (38–83)	39%	N/A	5.9 months
Platinum-containing regimens						
Tsimberidou et al ²⁰ (2004–2006)	OFAR1	20	59 (34–77)	50%	20%	8 months
Tsimberidou et al ²¹ (2007–2010)	OFAR2	35	63 (40–81)	43%	8.6%	6.6 months
Fludarabine-containing regimens						
Giles et al ²² (1992–1996)	PFA or CFA	12	59 (49–74)	45%	N/A	17 months
Tsimberidou et al ²³ (1997–2001)	FACPGM	15	62 (42–74)	5%	5%	2.2 months

CFA, cyclophosphamide-fludarabine-arabinosyl cytosine; FACPGM, fludarabine–cytarabine–cyclophosphamide–cisplatin–GM-CSF; GM-CSF, granulocyte-macrophage colony-stimulating factor; HyperCVAD, fractionated cyclophosphamide–vincristine–liposomal daunorubicin–dexamethasone; MTX, methotrexate; N/A, not available; OFAR, oxaliplatin–fludarabine–cytarabine–rituximab; PFA, cisplatin, fludarabine, cytarabine.

However, overall results remained poor, with 3-year RFS of only 27% post alloSCT and with 47% relapse (10 with DLBCL-RS, 2 with CLL) and 26% nonrelapse mortality at 3 years. Chemosensitive disease and the use of reduced-intensity conditioning were associated with superior RFS (largely due to lower nonrelapse mortality), stressing the importance of more effective therapy prior to alloSCT. Thirty-four patients in the same cohort who had chemosensitive disease underwent autoSCT. No plateau was seen on the RFS curve after autoSCT, with 3-year RFS of 45% (11 relapses with DLBCL-RS and 6 with CLL).²⁴

Novel approaches to DLBCL-RS treatment

Here, we cover reported clinical trial data on novel approaches in DLBCL-RS (Table 2). Unfortunately, the numbers of patients treated with most novel approaches are small, but several show encouraging results. In addition, the visual abstract also indicates some approaches with promising preclinical data and/or active clinical investigation, without publicly reported data available.

Small-molecule targeted agents

BTKis have dramatically improved outcomes for patients with CLL. However, results in DLBCL-RS have been more modest. Acalabrutinib monotherapy achieved responses, mostly partial responses, in 40% of patients, with a median duration of response (DOR) of 6 months.²⁵ Pirtobrutinib, a noncovalent BTKi

with a prolonged half-life, achieved at least partial responses in 6 of 9 very heavily pretreated patients with DLBCL-RS, all of whom had previously received a covalent BTKi, with a highly favorable toxicity profile.²⁶ Data from a much larger cohort treated with pirtobrutinib are eagerly awaited.

Three of 7 patients with DLBCL-RS responded to single-agent venetoclax in the phase 1 study.²⁷ This led to a multicenter phase 2 study of venetoclax in addition to R-EPOCH (rituximab, etoposide, prednisone, vincristine, cyclophosphamide, doxorubicin), which demonstrated an encouraging 50% CR rate.²⁸ Standard dosing of dose-adjusted R-EPOCH was utilized. Cycle 1 was given without venetoclax. Prior to cycle 2, rapid venetoclax ramp-up over 5 days, with inpatient tumor lysis syndrome monitoring, was performed, with no tumor lysis syndrome seen. Subsequently, venetoclax at 400 mg/d for 10 days was given concurrently with cycle 2 through 6 of dose-adjusted R-EPOCH. Of note, *TP53* alterations did not negatively affect the CR rate. Hematologic toxicity was substantial, and consequently, the median number of cycles of R-EPOCH given was 4. There were 2 therapy-related deaths, both occurring during cycle 1 (prior to the initiation of venetoclax). An additional cohort is being evaluated utilizing R-CHOP rather than R-EPOCH in the hope of mitigating hematologic toxicity and improving deliverability without compromising outcomes (NCT03054896). A retrospective analysis of 10 patients treated off protocol showed an encouraging 50% CR rate with venetoclax–R-CHOP.²⁹

Table 2. Novel approaches to DLBCL-RS treatment with published results

Treatment	Number of patients	Median number prior Rx (CLL + RT)	ORR/CRR (%)	Median PFS/DOR(mo)	Median OS (mo)
Small-molecule targeted agents					
Venetoclax monotherapy ³⁸	7	NR	43/0	NR/NR	NR
Acalabrutinib monotherapy ²⁵	25	1 for RT	40/8	3.2/6.2	NR
DTRM-555 (novel BTKi DTRMWXHS-12- everolimus- pomalidomide) ³⁹	24	5	45/9	NR/NR	NR
Pirtobrutinib ²⁶	9	6 (including 100% treated with covalent BTKi)	67/NR	NR/NR	NR
CIT + targeted agents					
R-EPOCH-venetoclax ²⁸	26	1 for CLL, 0 for RT	62/50	10.1/NR	19.6
Checkpoint inhibitors					
Pembrolizumab ³⁰	9	5	44/0	NR/NR	10.1
Pembrolizumab ³¹	23 (2 with CHL)	3 for RT, NR for CLL	5/0 (excluding 2 responders with CHL)	1.6/NR	3.8
Ibrutinib-nivolumab ⁴⁰	24	3	43/35	NR/10.	13.8
Ibrutinib- nivolumab ⁴¹	20	2	65/10	5.0/6.9	10.3
Venetoclax-obinutuzumab-atezolizumab ³³	7	NR	100/71	Not reached/not reached	NR
Bispecific antibodies					
Blinatumomab monotherapy (<i>Leukemia</i> , in press)	9	4 for CLL +2 for DLBCL-RS	22/11	1.9/NR	10.3
Blinatumomab after R-CHOP ³⁴	31	2 for CLL	54/39	NR/NR	NR
Antibody-drug conjugates					
Zilovetamab vedotin ³⁶	6	NR	67/17	NR/NR	NR
CAR T					
CD19 CAR T ⁴²	6 (DLBCL only)	5	67/67	NR/NR	NR
Axicabtagene ciloleuce ⁴³	8	4	100/63	NR/NR	NR
Lisocabtagene maraleuce ^l (European Breyanzi label)	4	NR	50/25	NR/2	NR

NR, not reported; RT, Richter's transformation.

Antibody-based therapy

Immune checkpoint blockade has shown some promising results. A study of the PD-1 inhibitor pembrolizumab, as monotherapy, showed that 4 of 9 patients achieved PR,³⁰ but a larger, multicenter follow-up study was disappointing.³¹ Somewhat superior results were seen with ibrutinib-nivolumab: 35% CR was seen in data from MD Anderson, with more favorable responses in ibrutinib-naïve patients.³² A follow-up multicenter study showed a 65% overall response rate (ORR; 10% complete response rate [CRR]) but with a median DOR of only 6.9 months.³² More

recently, atezolizumab-venetoclax-obinutuzumab achieved CR in 5 of 7 patients.³³

Responses have been seen with blinatumomab either alone (Thompson et al, *Leukemia* 2022, accepted) or as consolidation after R-CHOP.³⁴ Numerous bispecific antibodies are being studied in DLBCL, targeting CD19, CD20, CD22, CD37, and ROR1, reviewed elsewhere.³⁵

Finally, the ROR1-targeting antibody-drug conjugate zilovetamab vedotin achieved an ORR of 50% in patients heavily pretreated for CLL and/or RS.³⁶

Chimeric antigen receptor T-cell and chimeric antigen receptor natural killer cell therapy

In a retrospective evaluation of patients treated with axicabtagene ciloleucel for DLBCL-RS, 9 patients received the drug, 7 in combination with a BTKi.³⁶ The ORR was 100% in 8 evaluable patients, with 5 of 8 CRs. Similarly, in preliminary results of an ongoing clinical trial in Israel, 8 patients with DLBCL-RS received anti-CD19 chimeric antigen receptor (CAR) T cells, and 5 patients achieved CR. In the TRANSCEND-NHL001 trial, 4 patients with DLBCL-RS were treated with lisocabtagene ciloleucel (liso-cel), and 2 of 4 patients responded, with 1 of 4 achieving CR (European Breyanzi label). A single patient with RS in a phase 1 study of CAR natural killer (NK) therapy achieved CR.³⁷ Several studies with novel CAR NK products are ongoing.

Trials of interest

The ongoing PLATFORM trial (NCT03310619) is studying a combination of liso-cel plus targeted or immunotherapies in different combination cohorts. One cohort combined the checkpoint inhibitor durvalumab. One patient with relapsed/refractory DLBCL-RS achieved a CR for 2 years with this combination, but she unfortunately died of therapy-related myelodysplastic syndrome. On the ibrutinib as well as nivolumab combination cohorts in this trial, some other DLBCL-RS patients are experiencing durable remissions (data not published). Taken together, these results support further exploration of anti-CD19 CAR T for DLBCL-RS, potentially in combination with a BTKi as well as a checkpoint inhibitor. An investigator-initiated study of liso-cel-ibrutinib-nivolumab is planned at City of Hope National Medical Center and MD Anderson Cancer Center.

Bispecific antibodies targeting CD20 have achieved impressive rates of durable CRs in de novo DLBCL.^{44,45} Epcoritamab, a CD3x20 bispecific antibody, is being evaluated in DLBCL-RS (NCT04623541), and a multicenter investigator-initiated study is planned with glofitamab in RS (Davids M, personal communication).

Pirtobrutinib is active and very well tolerated in DLBCL-RS. Given its favorable toxicity profile, it appears an ideal agent to explore in combination with other active agents. A phase 2 study of pirtobrutinib-venetoclax-obinutuzumab in CLL and DLBCL-RS is planned at MD Anderson.

How we treat

We treat patients with CHL-RS similarly to patients with de novo CHL, usually with ABVD or similar as initial therapy. Anecdotally, we have seen excellent responses to PD-1 inhibition in patients with relapsed CHL-RS.

Among patients with DLBCL-RS, optimal risk stratification is essential. This requires consideration of patient fitness, extent of prior therapy, determination of clonal relationship to the underlying CLL, and *TP53* mutation/deletion status. We recognize that *IGHV* sequencing of tumor tissue to determine the clonal relationship of the DLBCL-RS to the underlying CLL is not universally available.

The above evaluation allows us to identify the small groups of patients who have relatively favorable outcomes with CIT: (1) those with clonally unrelated disease (who should be treated as de novo DLBCL) and (2) those who have untreated CLL and may lack *TP53* alteration. These patients can receive R-CHOP CIT alone, without alloSCT. For all other patients, the goal is to

achieve remission and then proceed with alloSCT for eligible patients, which is potentially curative for RS and CLL.

We stress that, where possible, all patients, especially those with poor-risk disease, should be enrolled in clinical trials. Outside clinical trials, patients predicted to have poor response to CIT (patients with *TP53* alterations and patients with clonally related disease who have previously received treatment for CLL) could be considered for an alternative approach, utilizing off-label therapy. The 2 approaches we generally utilize for such patients are R-CHOP or R-EPOCH combined with venetoclax or ibrutinib at 420 mg/d plus nivolumab at 3 mg/kg intravenously every 2 weeks. In general, we favor R-CHOP-venetoclax in patients who are CIT-naïve, given the approximate 50% CR rate with CIT plus venetoclax and the lesser degree of toxicity compared to R-EPOCH-venetoclax seen in our experience and in the de novo DLBCL literature. For patients previously treated with CIT for CLL or DLBCL-RS, we favor ibrutinib-nivolumab, especially in patients who are BTKi-naïve, given that response rates to this regimen are substantially higher in BTKi-naïve (64%) vs BTKi pretreated patients (23%).⁴⁰ When utilizing PD-1/programmed cell death 1 ligand 1 inhibitors as a bridge to alloSCT, it is important to remember the potential for increased severe acute graft-versus-host disease risk post alloSCT. This risk may be ameliorated by the use of posttransplant cyclophosphamide.⁴⁶

CLINICAL CASE (Continued)

The patient was enrolled in a clinical trial with R-CHOP-venetoclax (NCT03984448). Cycle 1 (R-CHOP alone, standard doses) was poorly tolerated, with grade 4 ileus and febrile neutropenia. Subsequent cycles, with the addition of venetoclax per protocol, were given without vincristine and with a dose reduction to 400mg/m² of cyclophosphamide and 25mg/m² of doxorubicin and were well tolerated. She achieved CR on PET/CT after cycle 3 and undetectable minimal residual disease to a level lower than 10⁻⁶ in the bone marrow by next generation sequencing following cycle 6, at which point she proceeded with alloSCT from a matched unrelated donor. She remains well thus far and in CR 3 months post alloSCT.

Conflict of interest disclosure

Philip A. Thompson: research funding: Adaptive Biotechnologies, Genentech, AbbVie, Pharmacyclics, Amgen, Lilly; advisory board: Adaptive Biotechnologies, Janssen, Pharmacyclics, AstraZeneca, Beigene, AbbVie, Genentech, Lilly; lecturer: Janssen Australia. Tanya Siddiqi: speaker: Astra Zeneca, Bristol Myers Squibb, BeiGene; advisory board: Astra Zeneca, BeiGene, Bristol Myers Squibb, Celgene, AbbVie, Pharmacyclics, Gilead; honoraria: Dava Oncology, ResearchToPractice.

Off-label drug use

Ibrutinib and nivolumab for Richter's syndrome; venetoclax combined with chemoimmunotherapy for Richter's syndrome.

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