#### **CLINICAL TRIALS AND OBSERVATIONS**

# Concurrent pembrolizumab with AVD for untreated classic Hodgkin lymphoma

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#### KEY POINTS

- Concurrent APVD was safe and effective in untreated HL without clinically significant treatment delays.
- CR rates by PET were lower than expected at all time points despite only one relapse, and no patient who cleared ctDNA has relapsed to date.

Concurrent administration of pembrolizumab with chemotherapy in untreated classic Hodgkin lymphoma (CHL) has not been studied previously. To investigate this combination, we conducted a single-arm study of concurrent pembrolizumab with AVD (doxorubicin, vinblastine, and dacarbazine; APVD) for untreated CHL. We enrolled 30 patients and met the primary safety end point with no observed significant treatment delays in the first 2 cycles. Twelve patients experienced grade 3 or 4 nonhematologic adverse events (AEs), most commonly febrile neutropenia and infection/sepsis. Grade 3 or 4 immunerelated AEs, including alanine aminotransferase elevation and aspartate aminotransferase elevation were observed in 3 patients. One patient experienced an episode of grade 2 colitis and arthritis. Six patients missed at least 1 dose of pembrolizumab because of AEs, primarily grade 2 or higher transaminitis. Among 29 response-evaluable patients, the best overall response rate was 100% and the complete response rate was 90%. With a median follow-up of 2.1 years, the 2-year progression-free survival (PFS) and overall

survival were 97% and 100%, respectively. To date, no patient who has withheld or discontinued pembrolizumab because of toxicity has progressed. Clearance of circulating tumor DNA (ctDNA) was associated with superior PFS when measured after cycle 2 and at the end of treatment (EOT). None of the 4 patients with persistent uptake by fluorodeoxyglucose positron emission tomography (PET) at EOT yet negative ctDNA have relapsed to date. Concurrent APVD shows promising safety and efficacy but may yield spurious PET findings in some patients. This trial was registered at www.clinicaltrials.gov as #NCT03331341.

#### Introduction

ABVD (doxorubicin, bleomycin, vinblastine, and dacarbazine) as well as combinations with brentuximab vedotin (BV; AAVD), have been the standard frontline treatment regimen for classic Hodgkin lymphoma (CHL) in North America, with the duration of treatment and use of consolidative radiotherapy dependent on baseline risk factors and stage.<sup>1-5</sup> Despite high cure rates, many patients will still relapse and require more intensive second-line therapies<sup>6-8</sup> followed by high-dose chemotherapy and autologous stem cell transplant to achieve a long-term remission.

Three drugs have been approved for CHL in the United States since 2011 ( $BV^{9}$ , nivolumab,<sup>10</sup> and pembrolizumab<sup>11</sup>), based on significant activity in the relapsed/refractory setting.

Pembrolizumab was found to have superior progression-free survival (PFS) vs BV in relapsed/refractory patients who were ineligible for autologous transplant.<sup>12</sup> Subsequent studies using these agents in the frontline setting have been promising, and recent long-term follow-up of the ECHELON-1 study suggests a potential overall survival (OS) benefit with BV-AVD vs ABVD in patients who are at advanced stages.<sup>5</sup> Programmed cell death protein 1 (PD1) inhibitor–based combinations so far have focused on performing a lead-in of PD1 inhibitor monotherapy followed by chemotherapy alone<sup>13,14</sup> or concurrent PD1 inhibitor + chemotherapy.<sup>14,15</sup> A phase 3 randomized study comparing concurrent nivolumab + AVD vs BV + AVD is ongoing, and results are not yet available.<sup>16</sup>

Interim positron emission tomography after 2 cycles of ABVD (PET2) has been identified as a prognostic marker  $^{\rm 17}$  and

subsequently validated using the Deauville 5-point scoring system, 18-20 leading to the development of several PETadapted trials.<sup>2,21,22</sup> Although initial results of these studies suggested that treatment intensification may improve outcomes compared with historical controls, the lack of a control arm has limited the interpretation of this data. Prospective data from uniformly treated patients with interim PET (PET2) results but who did not undergo treatment intensification or deescalation have been made available through long-term results from the ECHELON-1 study.<sup>23</sup> These findings suggest that improved surrogate markers are needed beyond PET2 for both risk stratification and early identification of treatment efficacy. Circulating tumor DNA (ctDNA) can be sequenced in plasma samples from patients with CHL,<sup>24,25</sup> and response assessments may be enhanced with newer ultrasensitive minimal residual disease (MRD) techniques.<sup>26</sup>

We hypothesized that concurrent pembrolizumab with AVD (APVD) chemotherapy would be safe, feasible, and effective therapy for untreated patients with CHL and would be delivered more efficiently than similar sequential approaches. In exploratory analyses, we aim to examine the role of ctDNA in response assessment and the examination of MRD compared with traditional radiographic approaches.

#### Methods

#### Trial conduct

This was a single-center, open-label, investigator-initiated clinical trial of APVD in untreated CHL. The protocol was approved by the institutional review board at our institution. The trial was conducted in accordance with the good clinical practice guidelines of the International Council for Harmonisation of Technical Requirements for Pharmaceuticals for Human Use and the principles of the Declaration of Helsinki.<sup>27</sup> All the patients provided written, informed consent. Drug-only support was provided by Merck, with research funding provided by philanthropic sources, including the Richard Hotes Foundation. Our institutional data safety monitoring committee reviewed safety data on a regular basis. The first draft of the manuscript was written by the first author. All the authors reviewed the data and contributed to the preparation of the final version of the manuscript. The authors vouch for the completeness and accuracy of the data and for the fidelity of the trial to the protocol and statistical analysis plan.

#### Patients

Eligible patients were at least 18 years of age having any stage of CHL with no prior systemic treatment for lymphoma, measurable disease of at least 1 cm in extranodal sites and 1.5 cm in nodal sites, an Eastern Cooperative Oncology Group performance status of 0 or 1, an ejection fraction of at least 50%, and adequate hematologic and organ function. Notable exclusion criteria include current or prior autoimmune disease (except vitiligo), current or prior history of pneumonitis/interstitial lung disease requiring corticosteroids, or current use of supplemental oxygen. Details of trial methods, including complete eligibility criteria, are provided in the protocol in the supplemental Appendix, which is available on the *Blood* website.

#### Treatment and assessments

A cycle of therapy was given every 28 days, with doses of AVD (doxorubicin [IV, 25 mg/m<sup>2</sup>], vinblastine [IV, 6 mg/m<sup>2</sup>], and dacarbazine [IV, 375 mg/m<sup>2</sup>]) on days 1 and 15, per the standard dosing.<sup>21</sup> Steroid premedication was allowed per institutional standard of care. Pembrolizumab (IV, 200 mg) was given every 21 days starting on cycle 1 day 1 (eq, cycle 1 day 1, cycle 1 day 22, cycle 2 day 15, etc). Treatment was allowed regardless of absolute neutrophil count with or without granulocyte colony-stimulating factor (GCSF) support, which was optional based on previous publication and institutional practice.<sup>28</sup> We recommended primary GCSF prophylaxis for patients who were older (ie, aged  $\geq$ 60 years), with comorbidities, or with prior episodes of febrile neutropenia, at the discretion of the investigator. All patients were required to receive a minimum of 2 cycles of study therapy, with up to 6 total cycles allowed, at the discretion of the investigator, based on stage and baseline risk factors. Preplanned consolidative radiotherapy was allowed. Growth factor support was allowed and optional. Treatment responses were collected per Lugano criteria<sup>20</sup> after 2 cycles of therapy, and if >2 cycles were administered, at the completion of all systemic therapy. Adverse events (AEs) were noted per Common Terminology Criteria for Adverse Events version 5.0 at the end-of-treatment (EOT) visit. Additional follow-up beyond the EOT visit was not specified in the protocol and was up to the institutional standard of care.

#### End points and statistical considerations

The primary objective of this study was to estimate the safety and tolerability of 2 cycles of the study treatment. At the time of study conception, there were concerns regarding the toxicity observed in the phase 1 study of BV + ABVD or AVD.<sup>29</sup> We did not want the addition of pembrolizumab to the chemotherapy backbone to lead to delays in chemotherapy because of immune-related toxicity. Using the phase 1 BV + chemotherapy study as a benchmark, we determined that we would deem the regimen safe if >85% of the patients are able to complete 2 cycles of treatment without a dose delay due to toxicity of >3 weeks. This was used as a stopping rule and served as the primary end point. Criteria for interruption or discontinuation of pembrolizumab because of immune-related AEs (IRAEs) was outlined in the protocol and available in the supplemental Material. Dose modification or interruption of AVD were performed per the institutional standard of care.

Our secondary objective was to estimate the number of patients in a complete response (CR; Deauville 1-3) via PET2/computed tomography (CT). Exploratory objectives include markers of efficacy, including overall response rate (ORR), CR rate, PFS, and OS, for the intention-to-treat population. We also examined the predictive capacity of PET2 with long-term outcomes in a prespecified analysis. Additional exploratory analyses, including survival analyses, were performed using ctDNA collected at baseline as well as after cycle 1, cycle 2, and the EOT. Using a separate institutional retrospective protocol, we were able to perform a post hoc tumor metabolic tumor volume (TMTV) analysis for all patients at baseline and for each response assessment.

#### ctDNA profiling and MRD detection

Phased variant enrichment and detection sequencing (PhasEDseq) for tracking MRD and cancer personalized profiling by deep sequencing for ctDNA profiling were performed at Stanford University as previously described,<sup>26,30</sup> with exceptions noted here. Whole blood was collected in either K2 EDTA tubes and processed within 6 hours or Streck BCT tubes and processed within 72 hours. Tubes were centrifuged twice at 1600g for 10 minutes at room temperature. After centrifugation, plasma was stored at -80°C in 1.8 mL aliquots until cell-free DNA (cfDNA) was isolated. Plasma-depleted whole blood was stored at -80°C until DNA was isolated from leukocytes. cfDNA was extracted from 1 to 11 mL of plasma (median, 5 mL) using the QIAamp Circulating Nucleic Acid Kit (Qiagen), following the manufacturer's instructions. After isolation, cfDNA was quantified using the Qubit dsDNA High Sensitivity Kit (Thermo Fisher Scientific) and the High Sensitivity NGS Fragment Analyzer Kit (Agilent). Genomic DNA (gDNA) from matched plasmadepleted whole blood was extracted using the Qiagen Dneasy Blood and Tissue Kit, quantified using the Qubit dsDNA High Sensitivity Kit, and fragmented to a target size of 170 base pairs using the Covaris S2 sonicator. Fragmented gDNA was purified using the QIAquick PCR Purification Kit (Qiagen). For cfDNA, a median of 32 ng (range, 6-50 ng) was inputted for library preparation. For gDNA from leukocytes, 75 ng of fragmented gDNA was inputted for library preparation. After sequencing on Illumina HiSeq4000/X-ten or NovaSeq6000 machines, data preprocessing, alignment, identification of single nucleotide variants and phased variants and detection of MRD were performed as previously described.<sup>26,30</sup> Quantitative levels of ctDNA were measured in haploid genome equivalents per mL and determined as the product of total cfDNA concentration and the mean allele fraction of somatic mutations expressed on a logarithmic scale (log10 haploid genome equivalents per mL).<sup>30,31</sup> ctDNA change from baseline was calculated as log10 (on-treatment ctDNA level/pretreatment ctDNA level) for visualization in waterfall plots. Wilcoxon ranksum test and Spearman correlation were used to correlate ctDNA levels with stage and TMTV, respectively. Plasma was prospectively banked at serial time points, with ctDNA MRD analysis performed later in batches, but not in real time.

# Assessment of TMTV and correlation with ctDNA levels

TMTV was analyzed using PET/CTs performed at baseline, after cycle 2, and at the EOT. Native support for the calculation of TMTV was performed using MIM Encore (version 7.1.3; Cleveland, OH), which uses a threshold-based, semi-automated technique to identify any tumor with a standardized uptake value above the liver background of maximum standardized uptake value.<sup>32</sup> TMTV analyses were performed retrospectively.

#### Results

#### Patients

Between 1 February 2019 and 18 June 2021, 30 patients were enrolled on the study (Figure 1). Six additional patients consented but subsequently withdrew because of patient decision (n = 4) and lack of insurance clearance (n = 2). Enrollment ended when the accrual goal of 30 patients was met.

Patient characteristics are presented in Table 1. The median age was 33 years (range, 18-69 years), and most patients were



Figure 1. CONSORT diagram.

females (18; 60%). Patients at all stages enrolled on the study, but most were in the advanced stage (18; 60%). At diagnosis, 13 patients (43%) had B symptoms, 6 (20%) had mediastinal bulk (>10 cm), 11 (37%) had an elevated erythrocyte sedimentation rate, and 7 (23%) had spleen involvement. Among the 12 patients at early stages, 6 (50%) were unfavorable per the National Comprehensive Cancer Network criteria. Among patients at advanced stages, 5 (28%) had an international prognostic score ranging from 4 to 7. Most patients received 6 cycles of treatment (22; 73%), with others receiving 2 cycles (5; 17%) and 4 cycles (3; 10%). Nine patients (30%) used growth factors for at least 1 cycle of treatment, including 4 (13%) for primary prophylaxis and 5 (17%) for secondary prophylaxis of febrile neutropenia.

#### Treatment exposure and safety

Thirty patients received at least 1 dose of study therapy and were evaluable for safety. A list of treatment-emergent AEs, observed in >10% of patients, as well as all grade 3 and 4 AEs are listed in Table 2. No deaths were observed in any study patient, either during treatment or during long-term follow-up. Serious AEs were observed in 6 patients (20%), including 5 patients with febrile neutropenia and 1 patient with a grade 4 large bowel obstruction/abscess in the setting of diverticulitis with no evidence of colitis. The most common grade 3 or 4 nonhematologic AEs observed include febrile neutropenia (5; 17%), hyponatremia (3; 10%), syncope (3; 10%), and sepsis (2; 7%). One secondary malignancy was observed, a diffuse large B-cell lymphoma (DLBCL) in a 69-year-old patient that presented 11 months after completion of the study therapy. In this patient, at diagnosis, the CHL was tested as Epstein-Barr encoding region in situ hybridization-positive, whereas the secondary DLBCL was Epstein-Barr virus-negative. For this reason, we felt this was more likely to represent a distinct malignancy and unlikely to represent a relapse of a composite lymphoma, though comparative clonality studies are not available.

#### Table 1. Patient characteristics

Characteristic	n = 30
Male sex, n (%)	12 (40)
Age, median (range), y	33 (18-69
Stage, n (%)         	1 (3) 11 (37) 7 (23) 11 (37)
B symptoms, n (%)	13 (43)
Mediastinal bulk, >10 cm, n (%)	6 (20)
Elevated erythrocyte sedimentation rate, >50, n (%)	11 (37)
Extranodal involvement, n (%)	11 (37)
Spleen involvement, n (%)	7 (23)
Early-stage unfavorable, National Comprehensive Cancer Network (n = 12), n (%)	6 (50)
International Prognostic Score (advanced stage, n = 18), n (%)	
0-1	6 (33)
2 and 3	7 (39)
4, 5, 6, and 7	5 (28)

There were no treatment delays >21 days observed during the first 2 cycles of therapy, meeting the study's primary end point. Three patients had treatment delays because of AEs, including 2 patients with grade 3 febrile neutropenia (2 and 5 days, respectively). One patient with a grade 4 large bowel obstruction/abscess in the setting of diverticulitis had treatment delay and doxorubicin omitted from cycle 3 onward because of newly onset grade 1 heart failure and discontinued all study treatment from cycle 4 day 22 onward because of recurrent postoperative complications. Six patients (20%) missed at least 1 dose of pembrolizumab because of AEs, primarily grade 2 or higher transaminitis (5; 17%). Three patients (10%) experienced grade 3 or higher aspartate aminotransferase (AST) or alanine aminotransferase (ALT) elevation, which required permanent discontinuation of pembrolizumab. One patient with grade 3 ALT elevation who was required to discontinue pembrolizumab was transitioned to AAVD therapy at investigator discretion but did not have evidence of progressive disease. One other patient who omitted at least 1 dose of pembrolizumab did so because of a newly onset grade 1 pleural effusion (subsequently found to be due to reduced ejection fraction), which was attributed to doxorubicin because there was no elevation of troponins or evidence of myocarditis observed. After several cycles with pembrolizumab and doxorubicin being held because of toxicity, all treatment was discontinued before cycle 5, and the patient has remained in remission for >1 year after completing all therapy.

We have outlined all grades of AEs deemed to be IRAEs in Table 3. In addition to the previously mentioned elevations of AST and ALT, we also observed grade 2 hypothyroidism (2; 7%),

rash (any grade, 13 [43%] and grade 2+, 3 [10%]), grade 2 arthritis (1; 3%), and grade 2 colitis (1; 3%). All elevations of AST or ALT are transient and reversible. Five patients with grade 2 or higher AST or ALT elevation were managed with holding pembrolizumab and corticosteroids as required by the protocol, including 3 patients with grade 3 or 4 transaminitis that required permanent discontinuation of pembrolizumab. All patients who required corticosteroids during the study treatment were able to discontinue them before the completion of chemotherapy. The episode of grade 2 colitis occurred 1 month after the completion of study treatment, in which all expected doses of pembrolizumab were received. The patient was able to discontinue all steroids ~3 months later. No patient who required any dose modification, discontinuation, or treatment delay has experienced progression to date. No pneumonitis or other grade  $\geq$ 3 IRAEs were observed. All IRAEs resolved by the EOT or during long-term follow-up, except those in patients who continued thyroid hormone supplementation.

#### Efficacy

Twenty-nine patients had at least 1 response assessment and were evaluable for efficacy. The best ORR was 100%, including 90% in CR. A PET2/CT showed an ORR of 100% and CR rate of 66%. In the 23 patients with early-stage unfavorable or advanced-stage disease, there was a PET2 CR rate of 57%. Three patients at early-stage disease (all favorable per the German Hodgkin Study Group criteria) with a PET2 CR proceeded with preplanned consolidative radiotherapy. At the EOT, 82% of patients were in complete remission. Five patients at the EOT had persistent or new fluorodeoxyglucose (FDG) uptake, 1 of whom ultimately developed biopsy-proven progressive disease. Two patients remained in partial response (Deauville 4) throughout the study treatment, without disease progression to date. One of these patients was followed up with surveillance CT and has remained progression-free 28 months later. Another patient had persistent FDG uptake upon follow-up PET/CT 3 months later, underwent a mediastinoscopy, showed the presence of benign thymic tissue, and has remained progression-free 28 months after treatment completion. Two other patients achieved a PET2 CR but developed FDG uptake upon EOT PET/CT. One of these patients had an additional PET/CT 4 months later that remained FDG-avid but with stable lymph node size. The investigator opted to transition to CT-based surveillance, which has not demonstrated progressive lymphoma to date, 36 months after the completion of therapy. Another patient's new FDG uptake resolved by the subsequent PET/CT, and the patient has remained progressionfree 33 months after completion of therapy. These 4 patients were followed up with serial imaging and/or biopsies, and, to date, none have developed recurrent HL 28, 28, 33, and 36 months after completion of study treatment. With a median follow-up of 2.1 years, only 1 patient has suffered a relapse, resulting in a 2-year PFS and OS of 97% (95% confidence interval, 90%-100%) and 100% (95% confidence interval, 100%-100%), respectively (Figure 2).

## MRD analysis with PhasED-Seq and correlation with TMTV

Banked plasma samples for all 29 response-evaluable patients were analyzed for ctDNA at baseline. Twenty-six patients had sufficient ctDNA detected at baseline and were eligible for MRD monitoring. One patient did not have detectable ctDNA

#### Table 2. Treatment-emergent AEs observed in more than 10% of patients and all grade 3 or 4 AEs

AE	Grade 1, n	%	Grade 2, n	%	Grade 3, n	%	Grade 4, n	%	All grade, n	%
Neutropenia	0	0	2	7	1	3	25	83	28	93
Anemia	18	60	8	27	2	7	0	0	28	93
Constipation	21	70	2	7	0	0	0	0	23	77
Nausea	16	53	4	13	0	0	0	0	20	67
ALT elevation	15	50	2	7	2	7	1	3	20	67
Fatigue	12	40	1	3	0	0	0	0	13	43
Rash	10	33	2	7	1	3	0	0	13	43
Peripheral sensory neuropathy	9	30	1	3	0	0	0	0	10	33
AST elevation	8	27	1	3	0	0	1	3	10	33
Dyspepsia	6	20	3	10	0	0	0	0	9	30
Oral pain	8	27	1	3	0	0	0	0	9	30
Headache	7	23	1	3	0	0	0	0	8	27
Insomnia	7	23	1	3	0	0	0	0	8	27
Diarrhea	8	27	0	0	0	0	0	0	8	27
Pain	8	27	0	0	0	0	0	0	8	27
Night sweats	8	27	0	0	0	0	0	0	8	27
Myalgia	7	23	1	3	0	0	0	0	8	27
Vomiting	5	17	2	7	1	3	0	0	8	27
Dyspnea	6	20	1	3	0	0	0	0	7	23
Alopecia	5	17	1	3	0	0	0	0	6	20
Hypokalemia	6	20	0	0	0	0	0	0	6	20
Upper respiratory tract infection	0	0	5	17	1	3	0	0	6	20
Thrombocytopenia	4	13	2	7	0	0	0	0	6	20
Bone pain	4	13	1	3	0	0	0	0	5	17
Back pain	5	17	0	0	0	0	0	0	5	17
Mucositis oral	5	17	0	0	0	0	0	0	5	17
Febrile neutropenia	0	0	0	0	5	17	0	0	5	17
Cough	3	10	1	3	0	0	0	0	4	13
Anxiety	3	10	1	3	0	0	0	0	4	13
Chills	4	13	0	0	0	0	0	0	4	13
Dry mouth	4	13	0	0	0	0	0	0	4	13
Arthralgia	4	13	0	0	0	0	0	0	4	13
Hyponatremia	0	0	0	0	3	10	0	0	3	10

AE	Grade 1, n	%	Grade 2, n	%	Grade 3, n	%	Grade 4, n	%	All grade, n	%
Syncope	0	0	0	0	3	10	0	0	3	10
Sepsis	0	0	0	0	1	3	1	3	2	7
Colonic obstruction	0	0	0	0	0	0	1	3	1	3
Colon abscess	0	0	0	0	0	0	1	3	1	3
Conjunctivitis	0	0	0	0	1	3	0	0	1	3
Total	30	100	23	77	13	43	25	83	30	100

#### Table 2 (continued)

at baseline, and 2 others had insufficient mutation reporters at baseline to allow for sensitive MRD detection. Of note, all 3 patients with insufficient baseline ctDNA at baseline had early-stage disease. Of the 26 patients evaluable for MRD assessment, all banked plasma samples could successfully be profiled at cycle 3 day 1 (n = 26) and the EOT (for those with >2 total cycles, n = 24).

Baseline quantitative ctDNA levels and tumor volume were compared with those at baseline stage and total metabolic tumor volume (Figure 3). ctDNA levels were lower in patients at early stage compared with patients at advanced stage (P = .001). In addition, TMTV was strongly correlated with ctDNA at baseline ( $R_{\rm S} = 0.72$ ; P < .0001).

We compared the log reduction in ctDNA levels with the baseline levels at both cycle 3 day 1 and the EOT (Figure 4). Undetectable ctDNA levels were common at both time points. At cycle 3 day 1, 21 patients (81%) were tested as ctDNA-negative, including 7 of 8 (88%) with early-stage favorable disease and 14 of 18 (78%) with early unfavorable/advanced-stage disease. At the EOT, ctDNA clearance increased to 24 of 26 (92%) in all patients, with 8 of 8 (100%) patients who were early-stage favorable and 16 of 18 patients who were early-stage unfavorable or advanced stage showing no detectable disease. The only known patient who relapsed had detectable ctDNA at all time points, whereas 4 patients with positive EOT PETs had undetectable ctDNA. None of these 4 patients have

had recurrent disease at 28, 28, 33, and 36 months since the completion of treatment. One additional patient had detectable ctDNA at the EOT. However, this patient had discontinued treatment during cycle 4 after developing a cardiomyopathy, so the EOT sample was drawn nearly 2 months earlier than expected. This patient remains without evidence of recurrence more than a year after discontinuing therapy. We performed a survival analysis comparing patients who were tested as ctDNA-positive vs those who were tested as ctDNA-negative at both cycle 3 day 1 and the EOT. ctDNA clearance was associated with improved PFS at both cycle 3 day 1 (P = .025) and the EOT (P = .0012).

#### Discussion

The therapeutic landscape for newly diagnosed CHL continues to evolve, with additional data on the use of various novel agents in combination with chemotherapy. In long-term follow-up, BV-AVD has shown continued PFS and now an OS benefit when compared with ABVD.<sup>5</sup> However, the risks of peripheral sensory neuropathy, febrile neutropenia, and the mandatory use of growth factors support the need for further improvements. In the relapsed setting, pembrolizumab has been shown to be superior to BV in a head-to-head comparison, leading to its exploration in the front-line setting by our study.

To our knowledge, these initial results of APVD in untreated CHL are the first to demonstrate the tolerability and activity of

AE	Any grade, n	%	Grade 1, n	%	Grade 2, n	%	Grade 3, n	%	Grade 4, n	%
Elevated ALT	20	67	15	50	2	7	2	7	1	3%
Elevated AST	10	33	8	27	1	3	—	_	1	3%
Rash	12	43	10	33	2	7	1	3	_	—
Hypothyroidism	2	7	_	_	2	7	_	_	_	_
Arthritis	1	3	_	_	1	3	_	_	_	_
Colitis	1	3	—	_	1	3	—	_	—	_
Total	26	87	24	80	7	23	3	10	1	3

#### Table 3. IRAEs of any grade



Figure 2. Kaplan-Meier estimates. PFS (A) and OS (B) for the response-evaluable study population. CI, confidence interval.

concurrent PD1 inhibitor anthracycline-based chemotherapy in patients at advanced stages without a PD1 inhibitor lead-in, building on the work of other trials using a sequential approach that demonstrated single-agent frontline anti-PD1 activity.<sup>13-15</sup> We were able to show that the APVD regimen showed promising safety and efficacy with similar overall outcomes over a much shorter time course than those of sequential PD1 inhibitor regimens. This shortened timeline not only resulted in a diminished total treatment duration for patients but also allowed for a more rapid achievement of disease control. This approach also avoids the possibility of experiencing an IRAE during the PD1 monotherapy portion that could delay or prevent the administration of combination chemotherapy.

The APVD regimen was well tolerated, and the IRAEs observed were manageable and reversible in all cases (excluding hypothyroidism). The concurrent approach saw increased rates of elevated AST and ALT compared with previously published monotherapy and sequential chemotherapy combinations, including cases of grade 3 or higher. Our findings are comparable with the results of using pembrolizumab with liposomal doxorubicin, gemcitabine, and vinorelbine and may indicate the potential for low-grade hepatotoxicity with concurrent chemotherapy and anti-PD1 strategies.<sup>8</sup> Although our protocol did call for up to 8 total doses of pembrolizumab (vs 3 total in the sequential approach<sup>13</sup>), most toxicity occurred early and likely was not a reflection of an increased cumulative dose of pembrolizumab. Fortunately, all patients remained asymptomatic, had no elevations of total bilirubin, and did not suffer lower remission rates.

We also observed a febrile neutropenia rate of 17%, compared with 10% observed in approaches with a PD1 inhibitor lead-in (Checkmate-205<sup>15</sup> and sequential pembrolizumab + AVD<sup>13</sup>) and 8% seen in ABVD.<sup>33</sup> Our patients experienced higher rates of at least 1 episode of neutropenia (93% any grade and 83% grade 4) than reported in these other frontline combination regimens. Despite this, we observed only 3 patients (10%) with grade 3 or 4 infections, in line with ABVD findings,<sup>34</sup> and below the 18% grade 3 or higher infection rate observed with AAVD that led to the mandatory use of GCSF. More importantly, we did not observe any pneumonitis in our study, which is said to affect up to 7% of patients treated with bleomycin. Careful attention is needed in larger PD1 and chemotherapy combination studies to determine whether checkpoint inhibitors continue to contribute to higher rates of febrile neutropenia



Figure 3. Total metabolic tumor volume and ctDNA analyses. (A) Comparison of baseline ctDNA levels in patients at early-stage I or II compared with in those at stage III or IV. (B) Scatterplot comparing baseline ctDNA levels to TMTV. hGE, haploid genome equivalents.



Figure 4. Exploratory analyses of ctDNA and PET with clinical outcomes. Waterfall plot with x-axis plotted from treatment initiation representing log change in ctDNA levels at cycle 3 day 1 (A) and EOT (B). Green bars represent patients with no evidence of relapse and the red bar represents the only patient with evidence of recurrence. Below each bar is a square that indicates the PET result at the same time point. Blue squares represent a complete metabolic response (Deauville 1-3) and orange squares indicate FDG uptake (Deauville 4, 5, or X). Of note, 2 patients with detectable ctDNA at baseline that only received 2 total cycles of treatment on study were represented on both waterfall plots. One patient who came off study after cycle 3 day 1 because of toxicity and received alternate frontline treatment was represented only on the cycle 3 day 1 waterfall plot. Survival analyses comparing PFS in patients with detectable or undetectable ctDNA at cycle 3 day 1 (C) and the EOT (D). ND, not detected.

and thus determine whether GCSF use should be mandated and also to assess the risk of rare IRAEs more intricately.

The role of PET-adapted therapy in the era of novel agent chemotherapy combinations remains undefined. Long-term follow-up from the ECHELON-1 study found that patients who were tested as PET2<sup>+</sup> and continued receiving AAVD and ABVD had a 5-year PFS of 60.6% and 45.9%, respectively.<sup>23</sup> These results exceed those shown in previously published retrospective reports,<sup>17,19</sup> and in the case of AAVD, they are very similar to the outcomes observed among patients who escalated to BEACOPP (bleomycin, etoposide, doxorubicin, cyclophosphamide, vincristine, procarbazine, prednisone) in prior PETadapted studies.<sup>21,22</sup> PET2-negativity was also an imperfect prognostic marker in both AAVD- and ABVD-treated patients, with a 5-year PFS of 84.9% and 78.9%, respectively, in these groups. APVD yielded a lower PET2 CR rate (66%) as well as EOT CR rate (82%) compared with historical regimens, yet only 1 out of 30 patients enrolled ultimately relapsed. Importantly, no patient who tested PET2<sup>+</sup> has relapsed to date, and only 1 out of 5 patients with FDG uptake on EOT imaging ultimately

relapsed. It is not clear why APVD was associated with increased rates of presumably false-positive results, but we have observed high rates of false-positive FDG-PET in other settings of pembrolizumab-chemotherapy combinations for lymphoma (PR-CHOP [pembrolizumab, rituximab, cyclophosphamide, doxorubicin, vincristine, prednisone]).<sup>35</sup> These false-positive findings were not reported in the sequential pembrolizumab + AVD approach<sup>13</sup> but were observed in the Checkmate-205<sup>15</sup> and NIVAHL<sup>14</sup> studies, suggesting that perhaps the concurrent use of these drugs may increase the likelihood of spurious FDG uptake. These results imply that most of the patients who test PET2<sup>+</sup> and are treated with APVD will achieve a long-term remission, countering the use of PET2adapted escalation approaches with this or similar regimens. These findings broadly call into question the utility of an interim PET in patients without signs of clinical progression treated with similar approaches.

To improve response assessment beyond FDG-PET, we explored ctDNA as a potential tool for measuring both tumor burden before therapy and residual disease during and after

therapy. Similar to findings in DLBCL, 26, 30, 31, 36, 37 we found baseline ctDNA levels in HL to reflect tumor burden, with ctDNA negativity after treatment being associated with favorable treatment outcomes. Previous efforts to demonstrate the utility of MRD in CHL have used immunoglobulin sequencing,<sup>3</sup> droplet polymerase chain reaction,  $^{39}$  or capture-based strategies tracking point mutations,  $^{24,25,40,41}$  generally associated with sensitivities of ~1:10 000 or less when using typical volumes of plasma. However, as with radiographic responses, deep molecular responses in CHL seem to be achieved substantially faster than in DLBCL, with most prior techniques demonstrating limited clinical sensitivity for the early detection of residual disease. Accordingly, highly sensitive assays such as PhasED-Seq<sup>26</sup> that allow for ultrasensitive MRD detection at levels  $\sim 10^{-6}$  are likely to be critical for measuring MRD after therapy of CHL. Remarkably, >80% of patients with HL in this study tested negative for ctDNA MRD after just 2 cycles of APVD therapy, using PhasED-seq. In contrast, less than half of the patients with DLBCL appear to clear ctDNA MRD after 2 cycles of therapy, using this same assay.<sup>42</sup>

Here, no patients who cleared ctDNA had recurrent CHL, and those with persistent FDG uptake received additional scans and biopsies. If PD1 inhibitor-based chemotherapy combinations are associated with increased false-positive PET scans,<sup>14,15</sup> then they may represent a tool to help avoid the anxiety and morbidity of additional scans and/or biopsies or potentially obviate the need for follow-up FDG imaging. Given the poor performance of interim PET in our and similar studies, a better understanding of ctDNA dynamics in patients treated for CHL may inform the design of future response-adapted clinical trials for treatment de-escalation. The negative predictive value of ctDNA MRD appears high in this study as well as in a pooled analysis,<sup>43</sup> which could inform future abbreviation strategies. In contrast, although the positive predictive value of ctDNA for treatment failure seems somewhat lower, particularly on interim assessments, the positive predictive value of interim ctDNA appears to be at least as high as that of interim PET in the context of APVD therapy. Nevertheless, future strategies for leveraging the positive predictive value ctDNA MRD for treatment escalation are likely to be most efficient when considering patients who test ctDNA MRD<sup>+</sup> at the end-of-induction landmark.

The generalizability of our results to date is limited given the small sample size, but our APVD data provide early insight into the safety and efficacy of this strategy. This was a single-arm study that enrolled a heterogeneous patient population, including limited and advanced stage patients who received different treatment durations based on their stage and baseline risk factors. Our study was not enriched for male patients (who are at higher-risk) or those having an international prognostic score ranging from 4 to 7. An additional cohort has since been added after completing this study to treat exclusively patients who are advanced stage to better define efficacy and further examine the dynamics of response assessment with PET and ctDNA. Another limitation is the every 3-week pembrolizumab dosing, which adds 4 additional visits over the course of 6 cycles and was required by the Merck Investigator Studies Program at the time of study conception despite the availability of every 6-week dosing outside of a clinical trial. The program has since allowed future studies to incorporate every 6-week dosing, which would allow for future studies to use a dosing schedule that would avoid additional infusion visits outside of the chemotherapy administration days.

In conclusion, concurrent APVD represents a time-efficient regimen with promising safety and efficacy as frontline therapy for CHL. However, a positive posttreatment FDG-PET after this approach was poorly predictive of persistent ctDNA and relapse. ctDNA may represent a more sensitive and specific treatment-agnostic response assessment tool that merits further study in larger data sets.

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## Authorship

Contribution: R.C.L. and A.K.G. designed the research and wrote the manuscript; R.C.L., C.S.U., C.P., E.H.W., S.D.S., M. Shadman, B.T., V.M.R., H.C., M. Shelby, S.O., S.K., K.V., Y.T., P.M., and E.M. performed the research; S.A., A.A.A., A.G., and D.L.C. contributed vital new reagents or analytical tools; R.C.L., H.D., J. Vandermeer, A.G., and H.R. collected data; R.C.L., S.A., A.A.A., A.G., D.L.C., J. Voutsinas, and A.K.G. analyzed and interpreted the data; and J. Voutsinas, S.A., and A.A.A. performed statistical analysis.

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#### REFERENCES

- Gordon LI, Hong F, Fisher RI, et al. Randomized phase III trial of ABVD versus Stanford V with or without radiation therapy in locally extensive and advanced-stage Hodgkin lymphoma: an intergroup study coordinated by the Eastern Cooperative Oncology Group (E2496). J Clin Oncol. 2013;31(6):684-691.
- Andre MPE, Girinsky T, Federico M, et al. Early positron emission tomography response-adapted treatment in stage I and II Hodgkin lymphoma: final results of the randomized EORTC/LYSA/FIL H10 trial. J Clin Oncol. 2017;35(16):1786-1794.
- Radford J, Illidge T, Counsell N, et al. Results of a trial of PET-directed therapy for earlystage Hodgkin's lymphoma. N Engl J Med. 2015;372(17):1598-1607.
- Engert A, Plutschow A, Eich HT, et al. Reduced treatment intensity in patients with early-stage Hodgkin's lymphoma. N Engl J Med. 2010;363(7):640-652.
- Ansell SM, Radford J, Connors JM, et al. Overall survival with brentuximab vedotin in stage III or IV Hodgkin's lymphoma. N Engl J Med. 2022;387(4):310-320.
- Lynch RC, Cassaday RD, Smith SD, et al. Dose-dense brentuximab vedotin plus ifosfamide, carboplatin, and etoposide for second-line treatment of relapsed or refractory classical Hodgkin lymphoma: a single centre, phase 1/2 study. Lancet Haematol. 2021;8(8):e562-e571.
- LaCasce AS, Bociek RG, Sawas A, et al. Three-year outcomes with brentuximab vedotin plus bendamustine as first salvage therapy in relapsed or refractory Hodgkin lymphoma. Br J Haematol. 2020;189(3): e86-e90.
- Moskowitz AJ, Shah G, Schöder H, et al. Phase II trial of pembrolizumab plus gemcitabine, vinorelbine, and liposomal doxorubicin as second-line therapy for relapsed or refractory classical Hodgkin lymphoma. J Clin Oncol. 2021;39(28):3109-3117.
- Gopal AK, Chen R, Smith SE, et al. Durable remissions in a pivotal phase 2 study of brentuximab vedotin in relapsed or refractory Hodgkin lymphoma. *Blood*. 2015;125(8): 1236-1243.
- Ansell SM, Lesokhin AM, Borrello I, et al. PD-1 blockade with nivolumab in relapsed or

refractory Hodgkin's lymphoma. N Engl J Med. 2015;372(4):311-319.

- Armand P, Shipp MA, Ribrag V, et al. Programmed death-1 blockade with pembrolizumab in patients with classical Hodgkin lymphoma after brentuximab vedotin failure. J Clin Oncol. 2016;34(31):3733-3739.
- 12. Kuruvilla J, Ramchandren R, Santoro A, et al. Pembrolizumab versus brentuximab vedotin in relapsed or refractory classical Hodgkin lymphoma (KEYNOTE-204): an interim analysis of a multicentre, randomised, openlabel, phase 3 study. *Lancet Oncol.* 2021; 22(4):512-524.
- Allen PB, Savas H, Evens AM, et al. Pembrolizumab followed by AVD in untreated early unfavorable and advancedstage classical Hodgkin lymphoma. *Blood*. 2021;137(10):1318-1326.
- 14. Bröckelmann PJ, Bühnen I, Meissner J, et al. Nivolumab and doxorubicin, vinblastine, and dacarbazine in early-stage unfavorable Hodgkin lymphoma: final analysis of the randomized German Hodgkin Study Group phase II NIVAHL trial. J Clin Oncol. 2023;41(6):1193-1199.
- 15. Ramchandren R, Domingo-Domènech E, Rueda A, et al. Nivolumab for newly diagnosed advanced-stage classic Hodgkin lymphoma: safety and efficacy in the phase II CheckMate 205 study. J Clin Oncol. 2019; 37(23):1997-2007.
- 16. Herrera AF, Li H, Castellino SM, et al. SWOG S1826: a phase III, randomized study of nivolumab plus AVD or brentuximab vedotin plus AVD in patients with newly diagnosed advanced stage classical Hodgkin lymphoma [abstract]. *Blood*. 2020;136(suppl 1):23-24.
- Gallamini A, Hutchings M, Rigacci L, et al. Early interim 2-[18F]fluoro-2-deoxy-Dglucose positron emission tomography is prognostically superior to International Prognostic Score in advanced-stage Hodgkin's lymphoma: a report from a joint Italian-Danish study. J Clin Oncol. 2007; 25(24):3746-3752.
- Meignan M, Gallamini A, Meignan M, Gallamini A, Haioun C. Report on the first international workshop on interim-PET-scan in lymphoma. *Leuk Lymphoma*. 2009;50(8): 1257-1260.
- **19.** Gallamini A, Barrington SF, Biggi A, et al. The predictive role of interim positron emission

#### **Footnotes**

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Proposals for collaborations using patient-level data are available on request from the corresponding author, Ajay K. Gopal (agopal@uw.edu).

The online version of this article contains a data supplement.

There is a Blood Commentary on this article in this issue.

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tomography for Hodgkin lymphoma treatment outcome is confirmed using the interpretation criteria of the Deauville five-point scale. *Haematologica*. 2014;99(6):1107-1113.

- 20. Cheson BD, Fisher RI, Barrington SF, et al. Recommendations for initial evaluation, staging, and response assessment of Hodgkin and non-Hodgkin lymphoma: the Lugano classification. *J Clin Oncol.* 2014; 32(27):3059-3068.
- 21. Johnson P, Federico M, Kirkwood A, et al. Adapted treatment guided by interim PET-CT scan in advanced Hodgkin's lymphoma. *N Engl J Med.* 2016;374(25): 2419-2429.
- 22. Press OW, Li H, Schöder H, et al. US intergroup trial of response-adapted therapy for stage III to IV Hodgkin lymphoma using early interim fluorodeoxyglucose-positron emission tomography imaging: Southwest Oncology Group S0816. J Clin Oncol. 2016; 34(17):2020-2027.
- 23. Straus DJ, długosz-Danecka M, Connors JM, et al. Brentuximab vedotin with chemotherapy for stage III or IV classical Hodgkin lymphoma (ECHELON-1): 5-year update of an international, open-label, randomised, phase 3 trial. *Lancet Haematol*. 2021;8(6):e410-e421.
- 24. Spina V, Bruscaggin A, Cuccaro A, et al. Circulating tumor DNA reveals genetics, clonal evolution, and residual disease in classical Hodgkin lymphoma. *Blood.* 2018; 131(22):2413-2425.
- 25. Sobesky S, Mammadova L, Cirillo M, et al. Indepth cell-free DNA sequencing reveals genomic landscape of Hodgkin's lymphoma and facilitates ultrasensitive residual disease detection. *Med (N Y)*. 2021;2(10): 1171-1193.e11.
- 26. Kurtz DM, Soo J, Co Ting Keh L, et al. Enhanced detection of minimal residual disease by targeted sequencing of phased variants in circulating tumor DNA. Nat Biotechnol. 2021;39(12):1537-1547.
- Association WM. World Medical Association Declaration of Helsinki: ethical principles for medical research involving human subjects. JAMA. 2013;310(20):2191-2194.
- 28. Evens AM, Cilley J, Ortiz T, et al. G-CSF is not necessary to maintain over 99%

dose-intensity with ABVD in the treatment of Hodgkin lymphoma: low toxicity and excellent outcomes in a 10-year analysis. *Br J Haematol*. 2007;137(6):545-552.

- 29. Younes A, Connors JM, Park SI, et al. Brentuximab vedotin combined with ABVD or AVD for patients with newly diagnosed Hodgkin's lymphoma: a phase 1, open-label, dose-escalation study. *Lancet Oncol.* 2013; 14(13):1348-1356.
- Kurtz DM, Scherer F, Jin MC, et al. Circulating tumor DNA measurements as early outcome predictors in diffuse large B-cell lymphoma. J Clin Oncol. 2018;36(28): 2845-2853.
- Alig S, Macaulay CW, Kurtz DM, et al. Short diagnosis-to-treatment interval is associated with higher circulating tumor DNA levels in diffuse large B-cell lymphoma. J Clin Oncol. 2021;39(23):2605-2616.
- 32. Morland D, Triumbari EKA, Maiolo E, et al. Healthy organs uptake on baseline (18)F-FDG PET/CT as an alternative to total metabolic tumor volume to predict event-free survival in classical Hodgkin's lymphoma. Front Med (Lausanne). 2022;9:913866.
- **33.** Connors JM, Ansell SM, Fanale M, Park SI, Younes A. Five-year follow-up of brentuximab

vedotin combined with ABVD or AVD for advanced-stage classical Hodgkin lymphoma. *Blood.* 2017;130(11):1375-1377.

- 34. Connors JM, Jurczak W, Straus DJ, et al. Brentuximab vedotin with chemotherapy for stage III or IV Hodgkin's lymphoma. N Engl J Med. 2018;378(4):331-344.
- 35. Smith SD, Till BG, Shadman MS, et al. Pembrolizumab with R-CHOP in previously untreated diffuse large B-cell lymphoma: potential for biomarker driven therapy. Br J Haematol. 2020;189(6):1119-1126.
- **36.** Roschewski M, Dunleavy K, Pittaluga S, et al. Circulating tumour DNA and CT monitoring in patients with untreated diffuse large B-cell lymphoma: a correlative biomarker study. *Lancet Oncol.* 2015;16(5): 541-549.
- Meriranta L, Alkodsi A, Pasanen A, et al. Molecular features encoded in the ctDNA reveal heterogeneity and predict outcome in high-risk aggressive B-cell lymphoma. *Blood.* 2022;139(12):1863-1877.
- 38. Oki Y, Neelapu SS, Fanale M, et al. Detection of classical Hodgkin lymphoma specific sequence in peripheral blood using a nextgeneration sequencing approach. Br J Haematol. 2015;169(5):689-693.

- 39. Camus V, Stamatoullas A, Mareschal S, et al. Detection and prognostic value of recurrent exportin 1 mutations in tumor and cell-free circulating DNA of patients with classical Hodgkin lymphoma. *Haematologica*. 2016; 101(9):1094-1101.
- Desch AK, Hartung K, Botzen A, et al. Genotyping circulating tumor DNA of pediatric Hodgkin lymphoma. *Leukemia*. 2020;34(1):151-166.
- 41. Camus V, Viennot M, Lequesne J, et al. Targeted genotyping of circulating tumor DNA for classical Hodgkin lymphoma monitoring: a prospective study. *Haematologica*. 2021;106(1): 154-162.
- 42. Roschewski M, Kurtz DM, Westin J, et al. MRD-negativity as a potential surrogate endpoint after frontline DLBCL therapy: pooled analysis of trials & implications for clinical trial design [abstract]. *Blood.* 2022; 140(suppl 1):785-786.
- 43. Alig SK, Shahrokh Esfahani M, Li MY, et al. Distinct molecular subtypes of classic Hodgkin lymphoma identified by comprehensive noninvasive profiling [abstract]. *Blood*. 2022;140(suppl 1): 1295-1296.