Mogamulizumab: a new tool for management of cutaneous T-cell lymphoma

This article was published in the following Dove Medical Press journal:
OncoTargets and Therapy

Abstract: Cutaneous T-cell lymphoma (CTCL) poses unique treatment challenges, given its range of presentations and numerous systemic therapy options. These options often lack comparative evidence or are characterized by low response rates and short remission duration in relapsed/refractory disease. The approval of mogamulizumab, a humanized, glycoengineered IgG1κ monoclonal antibody targeting the chemokine receptor type 4 (CCR4) chemokine receptor, brings a novel tool into the spectrum of treatment options for advanced CTCL and adult T-cell leukemia/lymphoma (ATLL). CCR4 is expressed in almost all cases of ATLL, and in a majority of CTCLs, particularly when blood involvement is present. In a Phase III randomized trial, mogamulizumab was associated with 28% overall response rate among patients with relapsed CTCL, median progression-free survival of 7.7 months, and median duration of remission of 14.1 months. Responses are more frequent among patients with Sézary syndrome and within the blood compartment. Common adverse effects include rash and infusion reactions, which are usually low grade. Sentinel reports indicate that exposure to mogamulizumab may result in severe or refractory graft vs host disease after allogeneic bone marrow transplantation, highlighting the need for vigilance and expert management. Further research may establish incremental efficacy of combining mogamulizumab with cytotoxic or immunomodulatory agents in CTCL, ATLL, and possibly other lymphomas and even solid tumors.

Keywords: cutaneous T-cell lymphoma, mogamulizumab, CCR4, adult T-cell leukemia, lymphoma, Sezary syndrome, mycosis fungoides

Introduction

The conventional approach to treatment of T-cell lymphomas is often to use regimens active in B-cell lymphomas, but this approach has suffered from lack of an effective T-cell-directed monoclonal antibody to use in place of rituximab. The recent Food and Drug Administration (FDA) approval of mogamulizumab in adult patients with relapsed or refractory mycosis fungoides (MF) or Sézary syndrome (SS) after at least one prior systemic therapy provides a new clinical tool that may fill in this historical gap. Cutaneous T-cell lymphoma (CTCL) is a rare non-Hodgkin lymphoma of skin-homing T-cells with a wide range of presentation and prognosis. In the USA, it has an overall incidence of 7.5 per million with the two most prevalent forms, MF and SS, accounting for two thirds of cases.1,2 Less common types of CTCL include primary cutaneous CD30+ lymphoproliferative disorders (lymphomatoid papulosis and primary cutaneous anaplastic large cell lymphoma [PCALCL]), primary cutaneous gamma/delta T-cell lymphoma, CD8-positive aggressive epidermotropic cytotoxic T-cell lymphoma, acral CD8-positive T-cell lymphoma, and CD4+ small/medium T-cell lymphoproliferative disorder.3 Furthermore, skin involvement may occur in other T-cell lymphomas, including subcutaneous panniculitis-like T-cell lymphoma,
extranodal natural killer (NK)/T-cell lymphoma, nasal type, or peripheral T-cell lymphoma (PTCL), not otherwise specified. Conventional thought had been that MF and SS represent different phases of disease progression, although molecular analysis suggests that the two forms of CTCL may in fact be distinct diseases. MF typically presents with plaques and patches, and despite low associated mortality, the lesions can cause significant pruritus affecting the quality of life. They may also cause pain and disfigurement. Making the diagnosis of early-stage MF can be difficult as it resembles many benign dermatologic conditions. A recent large international study found a median time to diagnosis from first symptom of 36 months. Advanced MF involves lymph nodes, blood, and organs outside the skin. In contrast, SS presents with diffuse erythroderma and blood involvement, with or without lymphadenopathy, and is characterized by a more aggressive course with 5-year overall survival (OS) of only about 26%. In advanced MF or SS, treatment is multidisciplinary and overall palliative. Relapses usually occur upon cessation of therapy, necessitating chronic management. Outside of allogeneic bone marrow transplantation, neither form of CTCL is curable.

The majority of patients with CTCL are diagnosed as early-stage MF (stage IA–II according to the tumor-node-metastasis-blood (TNBM) classification), and observation or local therapy alone are reasonable first steps. Local therapy includes topical steroids, topical chemotherapy (mechloroethamine), immunomodulators (imiquimod), radiation, or phototherapy, usually at the direction of a dermatologist. The 5-year disease-specific survival in early-stage MF approaches 90% compared with 30%–50% for advanced disease (stage IIB–IVB). In fact, in a large single-center series of patients with CTCL, only 11.6% progressed to a higher stage, and for T1 disease, median OS and disease-specific survival were not reached. In SS, patients with <1,000 Sézary cells/μL have median OS of 7.6 years, yet with ≥10,000 cells/μL, median survival plummets to 2.1 years. MF/SS patients often proceed through many lines of systemic treatment through their lifetime. Standard therapies include extracorporeal photopheresis, retinoids (bexarotene), histone deacetylase (HDAC) inhibitors (vorinostat and romidepsin), interferon α, methotrexate, pralatrexate, alemtuzumab, brentuximab vedotin, and now, mogamulizumab (Table 1). All have relatively low response rates ranging from 14% to 60% (mostly 20%–30%) and median duration of response rarely exceeding 1 year.

Given that monoclonal antibodies have revolutionized treatment for B-cell lymphoma, with survival outcomes often markedly differing between the eras before and after introduction of rituximab, the elusive goal in T-cell lymphoma has been to develop an equally safe and effective targeted antibody. Alemtuzumab (a monoclonal antibody against CD52) has shown efficacy, but with unacceptably high risk of infectious complications due to profound T- and B-cell depletion. While a naked anti-CD30 monoclonal antibody SGN-30 had only modest activity, brentuximab vedotin, an anti-CD30 antibody-drug conjugate, has demonstrated high efficacy in CD30+ CTCL and has been approved by FDA for PCALCL and for CD30-expressing MF. However, there is an ongoing need to identify other monoclonal antibodies that could selectively target T-cell-specific antigens with clinical efficacy and acceptable toxicity and that could be potentially combined with cytotoxic agents. Recently, mogamulizumab (Poteligeo®, Kyowa Kirin, Tokyo, Japan), a fully humanized, defucosylated IgG1 antibody against the C–C chemokine receptor type 4 (CCR4), has been granted FDA approval in CTCL based on improved outcomes demonstrated in the Phase III MAVORIC study (Figure 1).

### CCR4 expression and genomic profiling in T-cell lymphomas

CCR4 is a transmembrane chemokine receptor, which plays an important role in T-cell’s ability for homing and migration to the skin. While it has a particularly high expression on malignant cells, CCR4 is normally expressed on T<sub>reg</sub> cells and serves as the dominant chemokine receptor on Th2 and cutaneous lymphocyte antigen-expressing skin-homing T-cells. CCR4 is a receptor for the C–C chemokine ligand 17 (CCL17, originally termed thymus- and activation-regulated chemokine) and for CCL22, also known as macrophage-derived chemokine. It is one of the 18 known human chemokine receptors, whose overall job is to coordinate cell migration.

CCR4 expression by lymphoma cells varies by T-cell subtype and between studies, which use different qualitative and quantitative approaches to determine positive status, as well as various methods of assessment: immunohistochemistry of the skin biopsies (with cutoffs of “any,” 5%, or 10% positive cells), flow cytometry analysis of the blood, or reverse transcription polymerase chain reaction (Table 2). In adult T-cell leukemia/lymphoma (ATLL), CCR4 expression is nearly universal, in contrast to CTCL and PTCL, where expression varies greatly but appears to correlate overall with advanced or relapsed/refractory disease, particularly with blood involvement. Some T-cell lymphomas are Th1-polarized with high CXCR3 expression and no or minimal CCR4 expression;
Emerging evidence suggests also that CCR4 expression may be regulated by class I HDAC, specifically HDAC2.\(^{39}\) In an elegant study, Kitade et al assessed CCR4 expression before and after vorinostat therapy and found expression that may be regulated by class I HDAC, specifically HDAC2.

## Abbreviations

- CTCL: cutaneous T-cell lymphoma
- PFS: progression-free survival
- DOR: duration of remission
- NR: not reported
- ORR: overall response rate
- MAVORIC: a randomized, double-blind, placebo-controlled trial of vorinostat in patients with relapsed disease
- ALCANZA: a randomized, double-blind, placebo-controlled trial of vorinostat in patients with relapsed disease
- ATLL: adult T-cell leukemia/lymphoma
- CCR4: chemokine (C-C motif) receptor 4
- HDAC: histone deacetylase
- SS: Sézary syndrome
- PTCL: peripheral T-cell lymphoma
- ALK: anaplastic lymphoma kinase
- TP53: tumor protein p53

## Table 1: Comparison of mogamulizumab with other systemic treatment options for CTCL

<table>
<thead>
<tr>
<th>Drugs</th>
<th>Study/reference</th>
<th>Disease</th>
<th>N</th>
<th>Phase</th>
<th>Median prior lines</th>
<th>ORR (%)</th>
<th>PFS (months)</th>
<th>DOR (months)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mogamulizumab</td>
<td>Ogura et al (2014)(^{29})</td>
<td>PTCL, CTCL</td>
<td>37</td>
<td>2</td>
<td>2</td>
<td>38(^{b})</td>
<td>3.0</td>
<td>NR</td>
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<tr>
<td>Mogamulizumab</td>
<td>Duvic et al (2015)(^{35})</td>
<td>CTCL</td>
<td>42</td>
<td>1/2</td>
<td>3</td>
<td>37</td>
<td>11.4</td>
<td>10.4</td>
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<td>Mogamulizumab</td>
<td>Kim et al (2018)(^{20})</td>
<td>CTCL</td>
<td>186</td>
<td>3</td>
<td>3</td>
<td>28</td>
<td>7.7</td>
<td>14.1</td>
</tr>
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<td>Alemtuzumab</td>
<td>Lundin et al (2003)(^{26})</td>
<td>CTCL</td>
<td>22</td>
<td>2</td>
<td>3</td>
<td>55</td>
<td>NR</td>
<td>12.0</td>
</tr>
<tr>
<td>Belinostat</td>
<td>Foss et al (2015)(^{29})</td>
<td>PTCL, CTCL</td>
<td>53</td>
<td>2</td>
<td>4</td>
<td>14(^{c})</td>
<td>1.4(^{a})</td>
<td>2.7(^{a})</td>
</tr>
<tr>
<td>Bendamustine</td>
<td>Damaj et al (2013)(^{20})</td>
<td>PTCL, CTCL</td>
<td>60</td>
<td>2</td>
<td>1</td>
<td>50</td>
<td>3.6</td>
<td>3.5</td>
</tr>
<tr>
<td>Bexarotene</td>
<td>Duvic et al (2001)(^{31})</td>
<td>CTCL</td>
<td>94</td>
<td>2</td>
<td>5</td>
<td>45, 55(^{c})</td>
<td>NR</td>
<td>9.8, 12.6(^{c})</td>
</tr>
<tr>
<td>Bexarotene (59%) or</td>
<td>Prince et al (2017)(^{23})</td>
<td>CD30+</td>
<td>64</td>
<td>3</td>
<td>4</td>
<td>13</td>
<td>3.5</td>
<td>18.3</td>
</tr>
<tr>
<td>methotrexate (41%)(^{a})</td>
<td></td>
<td>CTCL</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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</tr>
<tr>
<td>Brentuximab vedotin</td>
<td>Prince et al (2017)(^{23})</td>
<td>CD30+</td>
<td>66</td>
<td>3</td>
<td>4</td>
<td>56</td>
<td>16.7</td>
<td>15.1</td>
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<tr>
<td>Denileukin diftitox</td>
<td>Prince et al (2010)(^{22})</td>
<td>CD25+</td>
<td>144</td>
<td>3</td>
<td>2</td>
<td>44</td>
<td>26.0</td>
<td>7.8</td>
</tr>
<tr>
<td>Lenalidomide</td>
<td>Querfeld et al (2014)(^{27})</td>
<td>CTCL</td>
<td>32</td>
<td>2</td>
<td>6</td>
<td>28</td>
<td>8.0</td>
<td>10.0</td>
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<tr>
<td>Pembrolizumab</td>
<td>Khodadoust et al (2016)(^{26})</td>
<td>CTCL</td>
<td>24</td>
<td>2</td>
<td>4</td>
<td>38</td>
<td>NR</td>
<td>NR</td>
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<td>Pralatrexate</td>
<td>Horwitz et al (2012)(^{25})</td>
<td>CTCL</td>
<td>54</td>
<td>1/2</td>
<td>4</td>
<td>41</td>
<td>12.7</td>
<td>Not reached</td>
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<tr>
<td>Pralatrexate and</td>
<td>Duvic et al (2017)(^{24})</td>
<td>CTCL</td>
<td>34</td>
<td>1/2</td>
<td>3.5</td>
<td>60</td>
<td>12.8</td>
<td>&gt;29</td>
</tr>
<tr>
<td>bexarotene</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Romidepsin</td>
<td>Whittaker et al (2010)(^{29})</td>
<td>CTCL</td>
<td>96</td>
<td>2</td>
<td>4</td>
<td>34</td>
<td>8.0(^{a})</td>
<td>15.0</td>
</tr>
<tr>
<td>Vorinostat</td>
<td>Olsen et al (2007)(^{26})</td>
<td>CTCL</td>
<td>74</td>
<td>2(^{c})</td>
<td>3</td>
<td>30</td>
<td>1.8(^{b})</td>
<td>&gt;6.1</td>
</tr>
<tr>
<td>Vorinostat</td>
<td>Kim et al (2018)(^{26})</td>
<td>CTCL</td>
<td>186</td>
<td>3</td>
<td>3</td>
<td>5</td>
<td>3.1</td>
<td>9.1</td>
</tr>
</tbody>
</table>

### Note

- CTCL patients only, control arm (physician’s choice) of the ALCANZA trial, bexarotene dose 300 mg/m\(^2\)/d and >300 mg/m\(^2\)/d, respectively, *time to progression reported rather than PFS.

### Abbreviations

- CTCL: cutaneous T-cell lymphoma
- PFS: progression-free survival
- DOR: duration of remission
- NR: not reported
- ORR: overall response rate
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- PTCL: peripheral T-cell lymphoma

### Include

Many cases of PTCL and angioimmunoblastic T-cell lymphoma. Others, like the ALK-positive anaplastic large cell lymphoma and about a third of PTCLs, express CCR4. While immunohistochemical expression in the skin of CTCL patients ranges from 14% to 97%, the proportion of positive cases is significantly larger (90%–100%) in clinical trials enrolling patients with relapsed disease.\(^{26,50,55}\) CCR4 is detectable using flow cytometry in almost all cases of CTCL involving the blood, with percentage of positive cells that vary from 31% to 97%, significantly higher than among healthy individuals (27%, on average).\(^{39,48}\) Furthermore, CCR4 expression is higher on circulating CD4\(^{+}\), CD26– lymphocytes in SS (59%) compared with inflammatory erythroderma (11%) or healthy controls (4%), although the difference is not evident in erythrodermic skin biopsies.\(^{30}\)

In the pivotal MAVORIC trial, out of 290 studied samples of relapsed/refractory MF/SS, 280 (97%) demonstrated CCR4 expression, with median percentage of positive cells of 80% (range 1%–100%), indicating that CCR4 expression in CTCL may actually be on par with ATLL.\(^{36}\) Of note, the percentage of CCR4-positive cells did not correlate with response to mogamulizumab in any of the clinical trials conducted to date.\(^{36,50,55}\)

Genomic profiling studies have further evaluated the role of CCR4 gene alterations in T-cell lymphomas. Activating (gain of function) CCR4 mutations were first identified in 26%–33% of ATLL cases,\(^{57,59}\) and then in 7% of patients with SS.\(^{40}\) Mutations in CCR4, along with TP53, have been described as potential cancer driver mutations in SS.\(^{40}\) CCR4 upregulation can also be used along with other identified genes in making a diagnosis of CTCL over similar appearing dermatoses, and it may be prognostic for progression and survival along with other genes.\(^{61,63}\) At least in ATLL, a small series suggests that CCR4 gain of function mutations are predictive of a better response to mogamulizumab without a difference in response to other treatments.\(^{59}\)

Emerging evidence suggests also that CCR4 expression may be regulated by class I HDAC, specifically HDAC2.\(^{64}\)
Table 2  CCR4 expression in subtypes of non-Hodgkin lymphoma

<table>
<thead>
<tr>
<th>Histology</th>
<th>Study/reference</th>
<th>CCR4+ (%)</th>
<th>CCR4+ (N/N total)</th>
<th>Method</th>
</tr>
</thead>
<tbody>
<tr>
<td>MF</td>
<td>Jones et al (2000)⁴⁹</td>
<td>14</td>
<td>1/7</td>
<td>IHC</td>
</tr>
<tr>
<td>MF</td>
<td>Kallinich et al (2003)⁵¹</td>
<td>92</td>
<td>11/12</td>
<td>IHC</td>
</tr>
<tr>
<td>MF</td>
<td>Yagi et al (2006)⁵²</td>
<td>54</td>
<td>14/26</td>
<td>IHC</td>
</tr>
<tr>
<td>Transformed MF</td>
<td>Jones et al (2000)⁴⁹</td>
<td>100</td>
<td>5/5</td>
<td>IHC</td>
</tr>
<tr>
<td>MF and SS</td>
<td>Ferenczi et al (2003)⁵⁶</td>
<td>100</td>
<td>11/11</td>
<td>FC</td>
</tr>
<tr>
<td>MF and SS</td>
<td>Sugaya et al (2015)⁵⁷</td>
<td>57</td>
<td>13/23</td>
<td>FC</td>
</tr>
<tr>
<td>MF and SS</td>
<td>Duvic et al (2012, 2015)⁵⁸</td>
<td>89</td>
<td>31/35</td>
<td>FC and IHC</td>
</tr>
<tr>
<td>MF and SS</td>
<td>Kim et al (2018)⁵⁹</td>
<td>97</td>
<td>280/290</td>
<td>IHC</td>
</tr>
<tr>
<td>SS</td>
<td>Narducci et al (2006)⁵⁹</td>
<td>100</td>
<td>12/12</td>
<td>FC</td>
</tr>
<tr>
<td>MF, SS, or PTCL</td>
<td>Yagi et al (2006)⁵²</td>
<td>100</td>
<td>5/5</td>
<td>IHC</td>
</tr>
<tr>
<td>ATLL</td>
<td>Yoshie et al (2002)⁵²</td>
<td>92</td>
<td>22/24</td>
<td>RT-PCR</td>
</tr>
<tr>
<td>ATLL</td>
<td>Phillips et al (2016)⁵⁴</td>
<td>91</td>
<td>65/71</td>
<td>FC and IHC</td>
</tr>
<tr>
<td>ALCL</td>
<td>Jones et al (2000)⁴⁹</td>
<td>73</td>
<td>8/11</td>
<td>IHC</td>
</tr>
<tr>
<td>ALCL</td>
<td>Yagi et al (2006)⁵²</td>
<td>100</td>
<td>5/5</td>
<td>IHC</td>
</tr>
<tr>
<td>DLBCL</td>
<td>Nakayama et al (2013)⁵⁷</td>
<td>13</td>
<td>10/80</td>
<td>IHC</td>
</tr>
</tbody>
</table>

Abbreviations: ATLL, angioimmunoblastic T-cell lymphoma; ALCL, anaplastic large cell lymphoma; ATLL, adult T-cell leukemia/lymphoma; CCR4, chemokine receptor type 4; DLBCL, diffuse large B-cell lymphoma; FC, flow cytometry; IHC, immunohistochemistry; MF, mycosis fungoides; NOS, not otherwise specified; PTCL, peripheral T-cell lymphoma; SS, Sézary syndrome.

The authors suggested that their findings may influence the order of treatments, as therapy with vorinostat might lower the amount of the target molecule for mogamulizumab. So far, this effect has not been observed in clinical experience, as the responses to mogamulizumab in CTCL were similar among patients crossing over from vorinostat (30%) compared with experimental arm (28%).

Figure 1  A synthesis of clinical data on mogamulizumab in ATLL and CTCL.

Abbreviations: BMT, bone marrow transplantation; CCR4, chemokine receptor type 4; GVHD, graft-vs-host disease; NOS, not otherwise specified; PTCL, peripheral T-cell lymphoma; SS, Sézary syndrome; US/EU/SA, United States, Europe, and South America; ATLL, adult T-cell leukemia/lymphoma; CTCL, cutaneous T-cell lymphoma; Mo., months.

ranged from 5% to 95% dropped to 5% to 20%.”⁶⁴ The authors suggested that their findings may influence the order of treatments, as therapy with vorinostat might lower the amount of the target molecule for mogamulizumab. So far, this effect has not been observed in clinical experience, as the responses to mogamulizumab in CTCL were similar among patients crossing over from vorinostat (30%) compared with experimental arm (28%).⁶⁵
In addition to mogamulizumab, CCR4 may serve as a target for other therapeutic modalities. Earlier attempts at targeting CCR4 were through “chemotoxins,” which fused CCL17 (CCR4 ligand) with neurotoxins or truncated *Pseudomonas* exotoxin released into the cytosol upon binding. More recently, CCR4 has been trialed in vivo as a target for chimeric antigen receptor T-cells.

**Development of mogamulizumab and its role in ATLL**

First approved in Japan for ATLL in 2012, mogamulizumab (KW-0761) is a defucosylated humanized IgG1κ monoclonal antibody. Its approval in Japan was expanded to PTCL and CTCL in 2014, and it gained FDA approval for MF and SS in 2018. Mogamulizumab, like its chimeric predecessor KM2760, binds to the N-terminal domain of CCR4 causing antibody-dependent cellular cytotoxicity (ADCC) rather than complement-mediated killing or direct cytotoxicity. ADCC depends on effector immune cells including macrophages, monocytes, and especially NK cells. Mogamulizumab binds to NK cell Fcγ receptor IIIa. Enhanced ADCC by monoclonal antibodies has been achieved by modifying the oligosaccharides in human IgG, particularly fucose. Defucosylation also allows for improved efficacy with drastically smaller doses of the drug compared with other antibodies. In vitro and murine studies have demonstrated the efficacy of KM2760 in models of ATLL and CTCL. In vivo, KM2760 caused ADCC (executed by peripheral blood mononuclear cells from healthy donors) on both established CTCL lines and tumor cells from patients with aggressive MF and SS. In a murine model, mice inoculated with a human CTCL cell line quickly developed large tumors and died within 3 months, while those treated with KM2760 lived longer without any obvious toxicity from the drug. These findings led to the development of a glycoengineered, fully defucosylated antibody KW-0761 (mogamulizumab), in a process similar to the one used to generate obinutuzumab. Mogamulizumab was first studied in a Phase I clinical trial (NCT00355472) enrolling 16 patients with ATLL (N=13), PTCL-NOS (N=2), and MF (N=1), which established the recommended dose of 1 mg/kg weekly for 4 weeks. No dose-limiting toxicities were observed in the dose escalation phase, and only one patient experienced a dose-limiting toxicity (grade 3 rash and febrile neutropenia) in the expansion cohort. Frequent (44%), although manageable infusion reactions, as well as rare reactivations of viral hepatitis and varicella-zoster virus infection were observed. In a subsequent multicenter Phase II trial (NCT00920790) in 27 subjects with relapsed ATLL, mogamulizumab showed 50% overall response rate with median progression-free survival (PFS) of 5.2 months and OS of 13.7 months. In that experience, infusion reactions were common (89%), but almost entirely grade ≤2. Rash was an additional frequent adverse effect observed in 63% of patients (19% grade 3). In a further follow-up of these Phase I and II trials, long-term survivors were observed, which was very encouraging for a disease associated with poor prognosis like the ATLL. In NCT00355472, four patients have survived over 3 years, and in NCT00920790, six (26%) had PFS exceeding a year, with estimated OS at 3 years of 23%.

The efficacy of mogamulizumab in relapsed/refractory ATLL was confirmed in an international randomized trial (NCT01626664), which enrolled 71 subjects receiving (in a 2:1 ratio) the monoclonal antibody or investigator choice. Overall response rate favored mogamulizumab (28% vs 8% in the control arm), with a median duration of response of 5.7 months, 47% rate of infusion reactions, 43% rate of rash, and 51% rate of infections. No significant differences in PFS (0.9 months in both arms) or OS (4.9 vs 6.9 months, including crossover of 75% of control arm to mogamulizumab) were seen between the study arms, although the trial was not powered to analyze these endpoints. The marked difference in PFS observed in this trial and in the prior Phase II trial may reflect a different mix of ATLL subtypes (acute, lymphomatous, and chronic) but may also suggest a different clinical profile of ATLL patients enrolled in USA, Europe, and Latin America compared with Japan.

**Clinical experience with mogamulizumab in CTCL**

Given the efficacy of mogamulizumab in ATLL and high CCR4 expression in MF and SS, mogamulizumab was naturally well-suited for a trial in CTCL. Assessment of drug efficacy in CTCL demonstrates unique challenges. While OS and PFS are valid endpoints, they fail to capture the benefit of symptom management, particularly pruritus, pain, and erythroderma, which have significant impact on patients’ quality of life. A standardized response assessment for use in clinical trials was proposed in 2011. Skin burden is assessed using the Severity Weighted Assessment Tool in original or modified form, SWAT or mSWAT, respectively, which uses body surface area and type of lesion (patch, plaque, or tumor), to calculate a score. Complete response (CR) in the skin is defined as clearance of disease in all areas, partial response as >50% regression, and stable disease as between 50% decrease and 25% increase according to mSWAT assessment. A Global Response Score, which incorporates responses within every disease compartment: skin, nodes,
viscera, and blood, has been adopted by many studies in CTCL to report the overall response rate.

The first published Phase II study of mogamulizumab in CTCL was a multicenter study in Japan (NCT01192984), in which 38 patients with CTCL and PTCL, selected based on CCR4-positive status (after screening 65 candidates), received mogamulizumab 1 mg/kg weekly for eight treatments only. Of the 37 treated subjects, 13 (35%) responded and five (14%) achieved a CR according to International Working Group criteria suitable more for PTCL. Median PFS was relatively brief at 3.0 months (95% CI, 1.6–4.9). In the subset of patients with CTCL, the response rate was 50% (4 out of 8) using the standardized CTCL Global Response Score. The authors also noted a pronounced and prolonged decrease in Treg cells. Like in other studies, percentage of CCR4-positive cells had no correlation with response. Adverse events included lymphopenia (81%, 73% grade 3/4), neutropenia (38%, 19% grade 3/4), infusion reactions (24%, no grade 3/4 events), and skin disorders (51%, 11% grade 3/4). Additionally, one patient developed grade 3 polymyositis, two cytomegalovirus retinitis, and one second primary malignancy. The response rate in CTCL was considered encouraging, particularly considering selection by CCR4 expression, which may correlate with more advanced disease.

In another Phase I/II study (NCT00888927), Duvic et al also demonstrated promising safety and efficacy for mogamulizumab in CTCL, including 22 subjects with MF and 19 with SS. Median age was 66 years (35–85 years), median number of prior systemic therapies was 3, and 63% of patients had stage IV disease. Similar to the ATLL experience, the researchers observed no dose-limiting toxicities, and in the Phase II they used 1.0 mg/kg weekly for 4 weeks, followed by every-2-week dosing until progression. The reported global response rate was 37% with more responders in the SS subgroup (47%) than in MF (29%). Skin-based responses occurred in 42% of patients, whereas responses in lymph nodes occurred in 25%. Furthermore, among patients with blood involvement by the lymphoma, the response rate in the blood was 95% (18 out of 19), with 58% (11) achieving a CR in that compartment. Three patients had a global CR. Median PFS was 11.4 months, impressive for relapsed CTCL, and median duration of response was 10.4 months. Most adverse events were grade 1/2, including nausea (31%), infusion reactions (21%), chills (24%), headache (21%), fever (19%), fatigue (17%), and rash (17%). There were no grade 4 events in the study, but the overall rate of serious adverse effects was 24%. Lymphopenia occurred in 41% and was considered an expected on-target effect.

Just over 10 years since the early in vivo studies of mogamulizumab in CTCL, the drug has received FDA approval for the treatment of MF or SS relapsing after ≥1 line of therapy. The approval followed a release of results of the largest randomized trial performed in CTCL, the Phase III international MAVORIC trial (NCT01728805). In MAVORIC, 372 patients were randomized to either mogamulizumab (four weekly doses at 1.0 mg/kg, followed by every-2-week dosing until progression) or vorinostat 400 mg daily—an FDA-approved oral HDAC inhibitor. Patient characteristics were well balanced between groups, with stage of disease ranging from IB to IVB, and slightly more MF (55%) than SS. All patients had received ≥1 prior line of systemic therapy (median 3) and had performance status of 1 or less on the Eastern Cooperative Oncology Group scale. The primary endpoint was PFS. As neither patients nor clinicians were blinded to treatment, the investigators used a blinded independent review to assess response and progression (including mSWAT evaluations and radiology scans), which were determined according to the global composite response score. Crossover was allowed from vorinostat to mogamulizumab and 136 of 186 subjects in the vorinostat arm crossed over (109 for progression, 27 for toxicity). The overall response rate for mogamulizumab was 28% compared with 5% for vorinostat, which in earlier Phase II study had a response rate of 30%. Stage IV MF or SS patients had again a higher response rate with mogamulizumab at 36% and 37%, respectively. However, only five patients achieved a global CR. PFS was superior in the mogamulizumab arm with median of 7.7 months (95% CI, 5.7–10.3) vs 3.1 months (95% CI, 2.9–4.1) for vorinostat (hazard ratio, 0.53; 95% CI, 0.41–0.69), sustained on independent review. Considering crossover, no significant difference in OS was observed (not reached for mogamulizumab vs 43.9 months for vorinostat, P=0.94). The PFS advantage for mogamulizumab persisted in all predefined subsets with the exception of stage I/IIB disease, where there was no difference (hazard ratio, 0.88; 95% CI, 0.58–1.35). Median time to response with mogamulizumab was 3.3 months, and median duration of response was 14.1 months. Responses were higher in the blood compartment (68%) than in the skin (42%), lymph nodes (17%), or viscera (0%). Among patients from the control arm who crossed over to receive mogamulizumab upon progression, response rate was 31% and median PFS was 8.9 months. Uniquely, MAVORIC researchers have also shown improvement in some aspects of quality of life, including skin pain and fatigue, among patients treated with mogamulizumab.
In the Phase III trial, as in prior experience, infusion-related reactions (35%) and rash (24%) were the most common adverse events, together with diarrhea (24%) and fatigue (24%). Most adverse events were grade 2 or lower, with rare (1%–2%) rates of higher grade constipation, nausea, diarrhea, fatigue, fever, cellulitis, pneumonia, sepsis, infusion reaction, hepatitis, weight loss, anorexia, hypertension, and rash. The rate of grade 3/4 events was 41%, equal in both arms, and rates of serious adverse events were 20% for mogamulizumab and 16% for vorinostat. Nineteen percent of patients discontinued mogamulizumab because of toxicity. Two treatment-related deaths occurred in the mogamulizumab arm: one related to sepsis and one related to polymyositis.

While mogamulizumab has been consistently relatively safe in clinical trials, serious rare adverse events have been noted in clinical experience. One rare, yet potentially fatal toxicity is Steven–Johnson syndrome or toxic epidermal necrolysis so far reported in less than ten cases, all among patients with ATLL. Another important risk results from depletion of CCR4-expressing nonmalignant Treg lymphocytes in patients who subsequently undergo allogeneic bone marrow transplantation, putting them at increased risk of graft-vs-host disease (GVHD). ATLL patients exposed to mogamulizumab have 1.8 times increased risk of grade 3/4 acute GVHD and 2.1 times increased risk of steroid-refractory GVHD, resulting in 44% nonrelapse mortality at 1 year. In a preliminary report, one of eight CTCL patients undergoing allogeneic bone marrow transplantation developed a severe acute GVHD. For any patient treated with mogamulizumab, transplantation should be delayed for at least 50 days from the last dose, or longer, and Treg counts may be assessed prior to transplant.

Conclusion

Mogamulizumab is a useful novel tool in the management of CTCL, and considering a favorable risk/benefit ratio, it is likely to be widely used for patients with relapsed MF/SS in USA. Unfortunately, its efficacy as a single agent remains modest, particularly in comparison with highly active monoclonal antibodies used in B-cell lymphomas. Mogamulizumab offers some unique benefits over alternative therapies in CTCL: the relatively long duration of remission confirmed in a large trial, high response rates within the blood compartment, and in SS. It may also prove useful in combination with other systemic agents, both as a direct antineoplastic agent and as immune modulator. For example, several ongoing trials combine mogamulizumab with checkpoint inhibitors (NCT03309878, NCT02476123) as a means of depleting undesirable Treg cells and enhancing their immune effect. Disappointingly, the first trial of combined chemotherapy and mogamulizumab in ATLL did not show improved rates of response or survival over chemotherapy alone.

Clinicians using mogamulizumab for CTCL and ATLL should be aware of the associated risks, particularly infusion reactions and rash (which rarely may become severe or even fatal), as well as the increased risk of GVHD, given that allogeneic bone marrow transplantation remains an important curative modality for both advanced CTCL and ATLL.

Disclosure

AJO reports research funding from Spectrum Pharmaceuticals, TG Therapeutics, and Genentech, and consulting from Spectrum Pharmaceuticals. The authors report no other conflicts of interest in this work.

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