T- and B-Lymphocyte Modulation Associated With Teriflunomide Treatment in Patients With Relapsing-Remitting MS: Analysis of the Phase 3b Teri-DYNAMIC Study

Heinz Wiendl,¹ Catharina C Gross,¹ Maren Lindner,¹ Melanie Eschborn,¹ Linda Weisser,¹ Anita Posevitz-Fejfar,¹ Andreas Schulte-Mecklenbeck,¹ Bart Van Wijmeersch,² Sandrine Brette,³ Timothy J Turner,⁴ Alexandre Jagerschmidt,⁵ Amit Bar-Or,⁶ Raymond Hupperts,⁷ Luisa Klotz¹; for the Teri-DYNAMIC Trial Group ¹University of Münster, Münster, Germany; ²Hasselt University, Hasselt, Belgium; ³Lincoln, Boulogne-Billancourt, France; ⁴Sanofi Genzyme, Cambridge, MA, USA; ⁵Sanofi Genzyme, Chilly-Mazarin, France; ⁶Montreal Neurological Institute, Montreal, QC, Canada; ⁷School for Mental Health and Neuroscience, Maastricht University, Maastricht, The Netherlands

OBJECTIVE

To study the effects of teriflunomide on T and B lymphocytes, and their functions, in patients with relapsing-remitting MS (RRMS)

INTRODUCTION

- Teriflunomide, a once-daily oral immunomodulator approved for the treatment of patients with RRMS, selectively and reversibly inhibits dihydroorotate dehydrogenase, a key mitochondrial enzyme in de novo pyrimidine synthesis required by rapidly dividing lymphocytes
- Unlike other currently available oral disease-modifying therapies (DMTs), teriflunomide limits the
 expansion of activated T and B cells underlying the damaging inflammatory processes associated with MS²
- Resting lymphocytes can still undergo homeostatic proliferation as their pyrimidine requirements are met through the salvage pyrimidine pathway, which is unaffected by teriflunomide!
- In clinical trials of teriflunomide, lymphocyte and neutrophil counts were reduced, on average by 10%–15% from baseline over a 6–12-week period and remained stable during treatment, with mean counts staying within the normal range (3.8–10.7 x $10^{9}/L$)^{3,4}
- Long-term safety observations and results from vaccination studies indicate that the effects of teriflunomide on activated T and B cells do not significantly impact protective immunity^{3,5,}
- There was no increased risk of malignancy or serious infection observed in patients receiving teriflunomide
- Individuals treated with teriflunomide mounted an effective immune response to seasonal influenza vaccination,⁵ and had seroprotective responses to rabies neoantigen and memory responses to recall antigens⁶
- In a rat experimental autoimmune encephalomyelitis model of MS, teriflunomide treatment significantly reduced the percentages of T cells in the cervical spinal cord and circulating B cells in the peripheral blood mononuclear cell (PBMC) population during acute attack phases⁷
- Prior to this trial, the effects of teriflunomide on the size and function of lymphocyte subpopulations have not been explored in humans; therefore, investigation of these parameters in patients with RRMS will provide greater insight into the mechanism of action of teriflunomide

METHODS

Study Design

- Teri-DYNAMIC (NCT01863888) is an exploratory, open-label, phase 3b clinical trial that included patients with RRMS, 18–56 years of age, meeting the McDonald 2010 criteria⁸ and either: Naïve to DMT/no DMT for ≥2 years; or
- Received interferon (IFN) B or glatiramer acetate (GA) therapy (with interruption of ≤3 months) and a period of ≥2 weeks without IFNB or GA
- Patients with RRMS were treated with teriflunomide 14 mg for 24 weeks
- Age- and sex-matched healthy individuals (reference population) remained untreated

Sample Collection and Handling

- Blood was collected at baseline, Week 12, and Week 24
- PBMCs were isolated by Ficoll® (Sigma-Aldrich Inc., St Louis, MO, USA) density gradient centrifugation

Sample Analysis

Immune Phenotyping

- Adaptive immune cell subsets were analyzed by flow cytometry as follows:
- CD4* T helper, CD4*; Th1, CD4*IFNY*; Th17, CD4*IL=17*; natural T regulatory [nTreg], CD4*Helios*FoxP3*; induced T regulatory [iTreg], CD4*Helios*FoxP3*; T cytotoxic, CD8*; B, CD19*

T-Cell Function

- The effects of teriflunomide treatment on T-cell proliferation and cytokine production were assessed following stimulation in culture as follows:
- T-cell proliferation was measured by flow cytometry using eFluor670 dye dilution following 72-hour stimulation with anti-CD3/anti-CD28
- Cytokine production was measured by Luminex assay® following 24-hour stimulation with anti-CD3/anti-CD28.
- In vitro assay with exogenous teriflunomide: PBMCs collected at baseline, with or without the addition of exogenous teriflunomide (100 μM)
- Ex vivo assay without exogenous teriflunomide: PBMCs collected at baseline and Week 24

Statistical Methods

- Data for patients with RRMS using comparisons of Week 12 and Week 24 with baseline are presented • The change from baseline to Week 24 was analyzed using a mixed-effect model with repeated measures, adjusted for baseline
- Adjusted least squares means (standard error) at Week 24 of changes from baseline are presented - Where data did not follow a Gaussian distribution, rank-transformed data were used and median
- changes from baseline (with interguartile ranges) are presented - Owing to the exploratory nature of the trial, no adjustment of P values for multiplicity was performed
- The per-protocol population comprised all patients with no maior/critical pharmacodynamic-related
- protocol deviations and with evaluable samples at baseline and Week 24

CONCLUSIONS

- The effects of teriflunomide on various immune cell types enhance our understanding of the teriflunomide unique mechanism of action, involving both activated T and B cells - Treatment with teriflunomide in patients with RRMS reduced the absolute numbers and frequencies of total lymphocytes, CD8+ T cells, and CD19+ B cells, consistent with observations from the phase 3 teriflunomide trials, TEMSO, TOWER, and TOPIC.⁴ However, total lymphocyte counts remained within normal range throughout the study
- Teriflunomide exerted a selective effect on different CD4⁺ T-cell subsets, suggesting a shift in T-cell populations from proinflammatory to regulatory, potentially anti-inflammatory, T cells
- The contrast between effects on proliferation ex vivo vs in vitro supports a reversible effect of teriflunomide
- The results of the Teri-DYNAMIC trial further support the efficacy of teriflunomide as an immunomodulatory agent to treat patients with RRMS

RESULTS

Participants

- The safety population (n=50) consisted of patients with RRMS with a median (min, max) Expande Disability Status Scale score of 1.5 (0, 4.0) at baseline and a median time since first symptoms of MS of 6.67 (0.3, 37.0) years
- Most patients (68%) had a recent history of IFNB-1 or GA treatment prior to the protocol-mandated treatment discontinuation (≥2 weeks) and study enrollment
- The per-protocol pharmacodynamic population consisted of 39 patients

Immune Phenotyping

Overall Lymphocyte Populations

- Total lymphocyte counts remained within the clinically normal range after 24 weeks of teriflunomide treatment, although treatment significantly altered absolute cell counts and percentages of immune cell subsets, as follows:
- Percentage and absolute numbers of lymphocytes were reduced compared with baseline (**Table 1**). However, decline in total lymphocytes was not clinically significant, as lymphocyte counts remained within normal range for adults (1000-4800 cells/µL)⁹
- Percentage of CD19⁺ B cells declined compared with baseline (Table 1)
- in an elevated CD4⁺/CD8⁺ ratio (median [interquartile range]: 0.6 [-0.2, 3.9], P=0.008)

Table 1. Total Lymph24 Weeks of Teriflund	ble 1. Total Lymphocytes and Lymphocyte Subsets in Patients With RRMS After Weeks of Teriflunomide 14 mg (n=39) ¹⁰				
Absolute counts (cells/µL)	Total Lymphocytes	CD4 ⁺ T Cells	CD8 ⁺ T Cells	CD19 ⁺ B Cells	
Change from baselineª	-284.94 (47.3) ^c	-101.14 (30.91) ^d	-34.7 (-55.5, -5.3) ^d	N/A	
P value ^b	<0.001	0.002 ^c	0.005 ^c		
Percentages	Total Lymphocytes	CD4 ⁺ T Cells	CD8 ⁺ T Cells	CD19 ⁺ B Cells	
	(% of PBMCs)	(% of CD3 ⁺ Cells)	(% of CD3 ⁺ Cells)	(% of Lymphocytes)	
Change from baseline ^b	-3.05 (1.43) ^d	1.6 (-0.5, 3.7) ^d	-1.0 (-3.5, 0.4) ^d	-2.0 (-4.1, 0) ^d	
<i>P</i> value ^c	0.040 ^d	0.006 ^d	0.009 ^d	0.001 ^d	

Represented as LSM change (SE) or median change (IQR); ^bP value from MMRM on change from baseline to Week 24, adjusted or baseline: fn=38; fn=37, IQR, interguartile range: LSM, least squares mean; MMRM, mixed-effect model for rep PBMC, peripheral blood mononuclear cell; RRMS, relapsing-remitting MS; SE, standard error.

Regulatory and T-Helper Cells

- · Within the CD4+ T-cell population, teriflunomide treatment resulted in significant changes from baseline through:
- Increased percentage of iTregs (Helios-FoxP3+, Figure 1A)
- Decreased absolute counts of nTregs (Helios*FoxP3+, Figure 1D)
- Decreased absolute counts of Th1 cells (Figure 1G)
- No significant change in percentage or absolute count of Th17 cells (Figures 1F and 1H)
 - Elevated iTreg/Th1 ratio (Figure 1I)

T-Cell Function

Proliferation

- Inclusion of teriflunomide in the in vitro proliferation assays led to significant reductions in the proliferation of both CD4* and CD8* T cells (Table 2)
- However, ex vivo assays demonstrated that there were no significant changes in prolife isolated T cells following teriflunomide treatment when comparing baseline with Week 24 (Table 2)

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ios-FoxP3+; 'n=38; 'Helios+FoxP3+; 'n=32; 'n=30. Bar graphs represent mean and error bars show SD (Figures A ents with RRMS treated with teriflunomide 14 mg. P value calculated from MMRM on change from baseline, perfo surser, inTegs, natural T-regulatory cells; RRMS, relapsing-remiting MS; SD, standard deviation; SE, standard dero ror bars show SD (Figures A–D, G, H), Box plots IOR with ures E. F. I. J). Perned on raw or rank data. Δ, change; IQR, interquartile range; iTregs, inducible T-regulatory cells; LSM, least square

Table 2. Proliferation of T Cells From Patients With RRMS In Vitro at Baseline and Ex Vivo After 24 Weeks of Teriflunomide 14 mg (n=39) ¹⁰		
Experimental Conditions	CD4 ⁺ T Cells ^a	CD8 ⁺ T Cells ^a
In vitro assays (n=25)		
Change with addition of exogenous teriflunomide 100 μM		
LSM (SE)	-32.84 (1.84)	-32.15 (2.15)
P value ^b	< 0.0001	< 0.0001
Ex vivo assays (n=33)		
Change from baseline to Week 24		
LSM (SE)	-2.1 (1.36)	-1.53 (1.01)
P value ^b	0.132	0.138
Cells were stimulated with anti-CD3/CD28 for 72 hours. *As % of total proli	ferating cells; ^b P value calculated	d from MMRM on

med on raw data. LSM, least squares mean: MMRM, mixed-effect model with repeated measure: nge from baseline inerfor RRMS, relapsing-remitting MS; SE, standard error

Cvtokine Production

- In vitro exposure to teriflunomide significantly reduced T-cell production of pro-inflammatory cytokines IL-2, TNFα, GM-CSF, IFN-γ, and IL-1β (Table 3)
- Compared with baseline, after 24 weeks of treatment with teriflunomide 14 mg, T-cell production of IL-2 and TNF α were suppressed, whereas production of GM-CSF, IFN- γ , and IL-1 β did not differ (Table 3)

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- Relative size of CD4⁺ population increased, while CD8⁺ population decreased (Table 1), resulting

	<0.001	0.002°	0.005°	IN/A
ages	Total Lymphocytes (% of PBMCs)	CD4 ⁺ T Cells (% of CD3 ⁺ Cells)	CD8+ T Cells (% of CD3+ Cells)	CD19 ⁺ B Cells (% of Lymphocytes)

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Experimental Conditions	IL-2	TNFα	GM-CSF	IFN-γ	IL-1
In vitro assays (n=19)					
Change with addition of					
100 µM					
Median (IQR)	-310.0 (-493, -151)	-1252.9 (17.9)ª	-235.0 (-313, -94)	-624.0 (-1531, -73)	-3.0 (-3
P value ^b	<0.0001	<0.0001	<0.0001	<0.001	<0.0
Ex vivo assays (n=26)					
Change from baseline to Week 24					
Median (IQR)	-147.0 (-505, 106)	-459.6 (106.8)°	-60.0 (-224, 76)	-200.0 (-2172, 92)	0 (–10,
P value ^b	0.002	<0.001	0.305	0.099	0.95

