ORIGINAL ARTICLE

Developing in vitro expanded CD45RA⁺ regulatory T cells as an adoptive cell therapy for Crohn’s disease

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ABSTRACT

Background and aim Thymus-derived regulatory T cells (Tregs) mediate dominant peripheral tolerance and treat experimental colitis. Tregs can be expanded from patient blood and were safely used in recent phase 1 studies in graft versus host disease and type 1 diabetes. Treg cell therapy is also conceptually attractive for Crohn’s disease (CD). However, barriers exist to this approach. The stability of Tregs expanded from Crohn’s blood is unknown. The potential for adoptively transferred Tregs to express interleukin-17 and exacerbate Crohn’s lesions is of concern. Mucosal T cells are resistant to Treg-mediated suppression in active CD. The capacity for expanded Tregs, to home to gut and lymphoid tissue is unknown.

Methods To define the optimum population for Treg cell therapy in CD, CD4⁺CD25⁺CD127⁻CD45RA⁺ and CD4⁺CD25⁺CD127⁻CD45RA⁻ Treg subsets were isolated from patients’ blood and expanded in vitro using a workflow that can be readily transferred to a good manufacturing practice background.

Results Tregs can be expanded from the blood of patients with CD to potential target dose within 22–24 days. Expanded CD45RA⁺ Tregs have an epigenetically stable FOXP3 locus and do not convert to a Th17 phenotype in vitro, in contrast to CD45RA⁻ Tregs. CD45RA⁺ Tregs highly express α4β7 integrin, CD62L and CC motif receptor 7 (CCR7). CD45RA⁻ Tregs also home to human small bowel in a C.B-17 severe combined immune deficiency (SCID) xenotransplant model. Importantly, in vitro expansion enhances the suppressive ability of CD45RA⁺ Tregs. These cells also suppress activation of lamina propria and mesenteric lymph node lymphocytes isolated from inflamed Crohn’s mucosa.

Conclusions CD4⁺CD25⁺CD127⁻CD45RA⁺ Tregs may be the most appropriate population from which to expand Tregs for autologous Treg therapy for CD, pavis the way for future clinical trials.

INTRODUCTION

Thymically derived FOXP3⁺ regulatory T cells (Tregs) are key mediators of peripheral tolerance and are likely to have a role in preventing inappropriate mucosal inflammation in response to bacterial, and other, luminal antigens. In mice, Treg depletion impairs oral tolerance.1 Adoptively transferred Tregs prevent the onset of colitis or treat established colitis in a number of murine models.2–7
How might it impact on clinical practice in the foreseeable future?

> These results demonstrate that initial $T_{reg}$ enrichment on the basis of CD45RA+ expression is required to produce a phenotypically stable and suppressive $T_{reg}$ population following in vitro expansion and that these in vitro expanded cells have the capacity to home to mucosal tissue, paving the way for autologous $T_{reg}$ therapy in this therapeutically challenging disease.

FOXP3 mutations lead to multisystem autoimmunity with enteropathy in mice and humans.8 9 Disruption of other key molecules implicated in $T_{reg}$ function, such as transforming growth factor (TGF)-β, Cytotoxic T Lymphocyte-Associated (CTLA)-4, interleukin (IL)-10R subunits, IL-2 or its receptor subunits, is associated with autoimmunity and intestinal inflammation.10

Human peripheral blood (PB) or umbilical cord blood $T_{regs}$ can be expanded in vitro through T cell receptor (TCR) stimulation in the presence of IL-2.11–26 In vitro expanded human $T_{regs}$ prevent transplant rejection,27 28 transplant arteriosclerosis29 and graft versus host disease (GvHD)21 30 in humanised mice. Promisingly, recent phase 1 clinical trials have shown $T_{reg}$ cell therapy to be safe in patients with GvHD12 24 and type 1 diabetes.31 Additional phase 1 studies have started in renal (the ONE study) and liver transplantation (ThRIL study).19 31

Lamina propria (LP) $T_{regs}$, are increased in the mucosa of patients with active Crohn’s disease (CD) and decreased in blood, compared with healthy controls.32–34 LP $T_{regs}$ obtained from inflamed CD mucosa suppress proliferation of conventional CD4+CD25lo/int T cells (Tcon) obtained from blood but not LP $T_{cons}$ suggesting that mucosal $T_{cons}$ in active CD may be resistant to $T_{reg}$-mediated suppression. LP $T_{cons}$ from CD mucosa over-express Smad7, an inhibitor of TGF-β signalling, which confers resistance to $T_{reg}$-mediated suppression.35 36 Activated $T_{cons}$ also have an effector-memory phenotype, conferring relative resistance to $T_{reg}$-mediated suppression.35 However, $T_{cons}$ expanded in vitro in the presence of rapamycin from the PB of patients with CD. These cells are resistant to Th17 plasticity, express lymphoid and gut homing markers, and home to human gut following adoptive transfer to a SCID mouse bearing subcutaneously implanted human small bowel (SB). In vitro expansion also enhances the suppressive ability of these cells, licensing them to suppress activation of LP and mesenteric lymph node (MLN) $T_{cons}$ obtained from inflamed CD resection specimens. These data suggest that CD PB CD4+CD25loCD127hiCD45RA+ cells may be the most appropriate population from which to expand $T_{reg}$s in vitro for forthcoming clinical trials of autologous $T_{reg}$ cell therapy in CD.

**Materials and Methods**

**Patient samples**

Following Institutional Review Board (IRB) approval (SE London REC 2; 10/H0804/65 and East London REC 2 (10/H0704/74)), patients with CD attending Guy’s & St Thomas’ National Health Service (NHS) Foundation Trust and Bart’s Health NHS Trust were invited to donate blood and/or resected tissue. Prospective written consent was obtained. Demographic details are shown in table 1.

**$T_{reg}$ enrichment and sorting**

Online supplementary figure S1 illustrates the experimental design. Peripheral blood mononuclear cells (PBMCs) were isolated by density gradient centrifugation over lymphocyte separation medium (LSM) 1077 and CD4+ lymphocytes enriched to >95% by positive magnetic activated cell separation (MACS) selection (Miltenyi, Bergisch-Gladbach, Germany). Lymphocytes were labelled using the ‘Human Regulatory T Cell Sorting Kit’ (BD Biosciences, San Diego, California, USA), as described previously,25 and sorted to CD4+CD25hiCD127loCD45RA− $T_{cons}$ (a$T_{cons}$) and CD4+CD25hiCD127loCD45RA+ $T_{cons}$ subsets, and autologous CD4+CD25hiCD127loCD45RA+ $T_{cons}$ on a FACSaria (BD; see online supplementary figure S2A–D). Median (IQR) postsort purity was 86.3% (80.8–91.6%); n=13 for CD4+CD25hiCD127loCD45RA− $T_{cons}$ and 92.7% (87.7–94.9%); n=13 for CD4+CD25hiCD127loCD45RA− $T_{cons}$. Autologous $T_{cons}$ were stored at −80°C.
In vitro generation of T<sub>reg</sub> lines
Precursor T<sub>reg</sub> populations were expanded in vitro as described previously<sup>21, 25</sup> and described in detail in online supplemental methods.

Cell surface and intracellular stains
Fluorochrome-conjugated antibodies, buffers and experimental technique are listed in online supplemental methods.

Assessment of the in vitro suppressive ability of putative T<sub>reg</sub>
Assays to determine T<sub>reg</sub> function in vitro were performed as described previously<sup>25, 49</sup> and described in detail in the online supplemental methods.

rtPCR
Following total RNA extraction from Trisure (Bioline, London, UK), cDNA was synthesised using the RevertAid First Strand cDNA Synthesis Kit and multiplex rtPCR performed in duplicate using the Maxima Probe/ROX qPCR Master Mix (both Thermo Fischer Scientific) on a BioRad C1000 Thermal Cycler. Primers are listed in online supplemental methods.

Estimation of cytokine concentrations
Cytokine concentrations were estimated in culture supernatants using the Cytometric Bead Array (CBA) Human Th1/Th2/Th17 Cytokine Kit (BD) or sandwich ELISAs (R&D), as indicated.

Assessment of IL-17 production under proinflammatory conditions
In vitro generated T<sub>reg</sub> were activated with anti-CD3/anti-CD28 beads at a 1:1 ratio and cultured at 10<sup>6</sup> cells/mL in complete Roswell Park Memorial Institute (RPMI) for 5 days at 37°C/5% CO₂, supplemented with the following cytokine cocktails, as previously described<sup>21, 23, 39</sup> (A) IL-2 (10 IU/mL, Proleukin); (B) IL-2, IL-1 (10 ng/mL), IL-6 (4 ng/mL) and TGF-β (5 ng/mL); (C) IL-2, IL-21 (25 ng/mL), IL-23 (25 ng/mL) and TGF-β (all R&D Systems). Supernatant IL-17 concentrations were measured by ELISA.

Assessment of FOXP3 promoter demethylation
Genomic DNA was isolated using a ‘DNeasy kit’ (Qiagen, Manchester, UK). Bisulfite conversion and assessment of the methylation status of the FOXP3 T<sub>reg</sub>-specific demethylated region (TSDR) was performed by Epiontis<sup>50, 51</sup> The genomic locations of FOXP3 and GAPDH CpG-rich regions probed have been reported.<sup>51</sup>

Isolation of LP mononuclear cells and MLN mononuclear cells
LP mononuclear cells (LPMCs) and MLN mononuclear cells (MLNMCs) were isolated as described previously<sup>52</sup> and listed in online supplemental methods.

C.B-17 SCID mouse human intestinal xenotransplant model
Experimental design is illustrated in figure 3C. The C.B-17 SCID mouse human intestinal xenotransplant model has been described previously<sup>11, 13, 15, 21, 23, 54</sup> and is described in detail in online supplemental methods. IRB and IACUC approvals were obtained prospectively (Ethics Committee for Animal Experimentation, Hebrew University of Jerusalem; MD-11-12692-4 and the Helsinki Committee of the Hadassah University Hospital; 81-23/04/04). Techniques for the detection of adoptively transferred T<sub>reg</sub> are also described in detail in online supplemental methods.

Statistical analysis
Statistical analysis was carried out using GraphPad Prism 5 (GraphPad Software Inc, La Jolla, California, USA) and the methods used are described in detail in the online supplemental methods.

RESULTS
T<sub>reg</sub> can be expanded from the blood of patient with CD using GMP-compatible protocols
Hoffmann <i>et al</i><sup>17</sup> showed that initial T<sub>reg</sub> enrichment on the basis of CD45RA<sup>+</sup> expression was required to expand homogenous and stable T<sub>reg</sub> lines from healthy donors in the absence of supplemental rapamycin. Rapamycin prevents the outgrowth of contaminating T<sub>cons</sub> in T<sub>reg</sub> cultures, and may make the requirements for the starting population less stringent.<sup>11, 13, 15, 21, 23, 55</sup> However, the optimum precursor population from which to expand a homogenous, suppressive and epigenetically stable T<sub>reg</sub> population from CD PB is currently unknown. In previous studies, we accomplished in vitro expansion of in vitro suppressive T<sub>reg</sub> from healthy controls<sup>21</sup> and renal transplant candidates.<sup>26</sup> We sought to determine if T<sub>reg</sub> could be expanded in vitro from the blood of patients with CD.

Freshly isolated CD4<sup>+</sup> lymphocytes from 13 patients with CD were fluorescence-activated cell sorting (FACS)-sorted into CD4<sup>+</sup>CD25<sup>hi</sup>CΔ127<sup>lo</sup>CΔ45RA<sup>+</sup> (median (IQR) of 2200 cells/mL PB (860–4400)) and CD4<sup>+</sup>CD25<sup>hi</sup>CΔ127<sup>lo</sup>CΔ45RA<sup>+</sup> subsets (3700 cells/mL (2000–4500)), then expanded in vitro in the presence of high-dose IL-2, rapamycin and anti-CD3/<i>-</i>anti-CD28 beads. Active disease, evidenced by a Harvey Bradshaw Index >5 (n=4) or elevated C reactive protein (n=1), was not associated with a significantly reduced yield (see online supplementary figure S2E). Donor clinical characteristics are given in table 1.

Table 1 Demographic details of study patients

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<th>Characteristics</th>
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<td>Penetrating (B3)</td>
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GI, gastrointestinal; HBI, Harvey Bradshaw Index.


http://gut.bmj.com/ on May 25, 2022 by guest.
Every CD45RA\(^+\) T\(_{\text{reg}}\) line proliferated, to a median (IQR) of 175-fold (66–531; n=13) at D24 (figure 1A). In contrast, 3 of 13 (23%) CD45RA\(^-\) T\(_{\text{reg}}\) lines did not proliferate and were discontinued. CD45RA\(^-\) T\(_{\text{reg}}\) expanded 130-fold (8–209; n=10). Expanded T\(_{\text{reg}}\) were exclusively CD4\(^+\) lymphocytes. Expression of CD25 and FOXP3 was comparable in D24 CD45RA\(^+\) and CD45RA\(^-\) T\(_{\text{reg}}\) (figure 1B), but a greater proportion of CD45RA\(^+\) T\(_{\text{reg}}\) maintained a CD4\(^+\)CD25\(^{\text{hi}}\)CD127\(^{\text{lo}}\)FOXP3\(^+\) phenotype (p=0.037; figure 1C).

Proliferation assays were performed to determine if in vitro expanded T\(_{\text{reg}}\) retained the ability to suppress proliferation of autologous CD4\(^+\)CD25\(^{\text{lo}}\) T\(_{\text{con}}\). CD45RA\(^+\) and CD45RA\(^-\) T\(_{\text{reg}}\) suppressed T\(_{\text{con}}\) proliferation to an equivalent degree (figure 1D–E), demonstrating specific suppression (vs the 2X cell density control) above an 8:1 T\(_{\text{con}}\):T\(_{\text{reg}}\) ratio. CD45RA\(^+\) and CD45RA\(^-\) T\(_{\text{reg}}\) reduced IL-2 expression in 96 h co-culture supernatants (see online supplementary figure S3A). CD45RA\(^+\) T\(_{\text{reg}}\) also suppressed IFN-\(\gamma\) expression in 96 h co-culture supernatants (see online supplementary figure S3B).

In vitro expanded CD45RA\(^+\) T\(_{\text{reg}}\) are resistant to IL-17 induction and stably express FOXP3
The ‘inflammatory potential’ of in vitro expanded T\(_{\text{reg}}\) from patients with CD was examined. Genes important in...
Th17 biology, including RORC, AHR and IL-17, were significantly overexpressed in CD45RA− Tregs, in comparison with expression in paired CD45RA+ Tregs (p<0.05 for each comparison, figure 2A). IL-17 secretion was also significantly different in these Treg subsets. IL-17 expression was below the limit of detection in 10/11 (91%) CD45RA+ Tregs and significantly higher in CD45RA− Tregs (p=0.02; figure 2B).

The potential of in vitro expanded Tregs to turn on an inflammatory programme following exposure to Th17-inducing cytokines, as occurs in vitro in Treg subsets isolated from blood, was examined. D24 Tregs were washed and cultured for a further 5 days in the presence of IL-2 alone, a cocktail of IL-2, IL-1, IL-6 and TGF-β or a cocktail of IL-2, IL-21, IL-23 and TGF-β, n=17, bar at median. % FOXP3 TSDR demethylation; n=15, bar at median. *p<0.05, **p<0.01, ***p<0.001. Tregs, thymus-derived regulatory T cells; IL, interleukin; IFN, interferon; TNF, tumour necrosis factor; TGF, transforming growth factor; TSDR, Treg-specific demethylated region; NS, not significant.

Th17-inducing cytokines failed to induce IL-17 production by CD45RA+ Tregs. In contrast, IL-17 production by CD45RA− Tregs was 3-fold higher than CD45RA+ Tregs in neutral conditions (IL-2 alone) and 10-fold higher in skewing conditions (p<0.001 each comparison).

To ensure that phenotypic stability of CD45RA+ Tregs correlated with an epigenetically stable FOXP3 locus, we determined the methylation status of the FOXP3 ‘TSDR’ (figure 2D). We found the TSDR to be completely demethylated in all CD45RA+ Treg lines tested (100%; n=9), suggesting an epigenetically stable FOXP3 locus in CD45RA+ Tregs even after 24d of in vitro expansion. In contrast, variable degrees of TSDR demethylation were seen in CD45RA− Treg lines (median (IQR) of 90.6% (36.6%–100%); n=6; p=0.008).

In vitro expanded CD45RA+ Treg express homing receptors for gut and lymphoid tissue

The ability of in vitro expanded Treg to home to relevant immune niches, where they may suppress inflammation, is thought to be critical for cell therapy. Consequently, the expression of gut homing receptors on in vitro expanded Tregs was examined by FACS (figure 3A, B). We found that D24 CD45RA− Tregs modestly expressed α4β7 integrin and CCR6 (20.8%±7.8% and 12.2%±7.9%, respectively) and did not express CCR9. Both CD62L (84.8%±20.6%; p=0.04 vs CD45RA+) and CCR7 (92.1%±12.8%; p=0.03) required for lymph node homing were more highly expressed in CD45RA− Tregs than CD45RA+ Tregs, CCR4 (95.4%±4.2%) was also highly expressed.

Adoptively transferred CD45RA+ Treg home to inflamed human small intestine in a C.B-17 SCID human SB xenotransplant model

In view of the favourable phenotype of CD45RA+ Tregs as a candidate cell therapy, we next sought to determine whether these cells could home to inflamed human SB in vivo. D24 CD45RA+ Tregs were administered to a C.B-17 SCID mouse bearing human small intestinal xenotransplants and homing assessed.
24 h later (figure 3C, D). Intraluminal injection with enteropathogenic *Escherichia coli* was used to induce mucosal inflammation (see online supplementary figure S4A). Following adoptive transfer, human CD45+CD3+CD4+ cells were detected in mouse spleen and inflamed human SB LP by FACS (see figure 3E; gating strategy online supplementary figure S4B),
indicating that adoptively transferred CD45RA+ Tregs homed to inflamed human SB LP in this model. This was confirmed by the detection of human CD45+CD3+ cells in inflamed human SB LP by immunofluorescence (figure 3F). We previously showed that human fetal SB contains a population of CD3−CD7+ cells that persist following xenotransplantation.53 Human CD45+CD3− events were also detected in non-inflamed human SB LP in both mice that received intravenous PBS and intravenous Tregs (figure 3E), suggesting that a population of long-lived human immune cells was co-transferred with the human SB transplant.

In vitro expansion enhances the in vitro suppressive ability of CD45RA+ Tregs

LP Tcons from inflamed CD mucosa are resistant to in vitro suppression by autologous LP Tregs.35 36 Consequently, it is possible that in vitro expanded Tregs will need an enhanced suppressive function in order to be successful as a future cell-based therapy. Expansion with supplemental rapamycin enhances the in vitro suppressive function of Tregs from patients with ESRF, SLE, RA, MS and asthma.26 38 In order to determine if in vitro expansion enhanced Treg function in patients with CD, freshly isolated CD45RA+ Tregs or D24 CD45RA+ Tregs that were expanded in vitro from these FACS-sorted CD4−CD25hiCD127loCD45RA+ Tregs, or D24 CD45RA+ Tregs that were expanded in vitro from these FACS-sorted CD4−CD25hiCD127loCD45RA+ Tregs (n=3 independent experiments; cells from the same lot of single-donor, freeze-thawed Tcons for each experiment). D24 CD45RA+ Tregs suppressed Tcon proliferation to a greater degree than the freshly isolated CD4−CD25hiCD127loCD45RA+ Tregs from which they were expanded, at both a 4:1 and 8:1 Tcon:Treg ratio (p<0.01 and p<0.001, respectively; figure 4A). This suggests that in vitro expansion enhances the suppressive ability of D24 CD45RA+ Tregs.

In vitro expanded CD45RA+ Tregs suppress proliferation and activation of MLN and LP T cells in active CD

We next wished to determine if D24 CD45RA+ Tregs could suppress activation and proliferation of Tcons taken from the MLN and LP of patients with CD (figure 4B, C). MLNMCs were co-cultured with Treg and CD3+ proliferation assessed at 96 h. Dose-dependent Treg-mediated suppression of MLN CD3+ proliferation was seen at each MLNMC:Treg ratio (figure 4D). We were unable to demonstrate in vitro suppression of LPMCs CD3+ proliferation with this technique, as both freshly isolated and freeze-thawed LPMCs obtained from inflamed CD mucosa died prior to acquisition at 96 h (n=4 independent experiments; see online supplementary figures S5 and S6).

We recently validated a novel co-culture assay for the assessment of in vitro expanded Treg function. This takes advantage of Treg-mediated suppression of the early activation marker CD154 (CD40 L) on Tcons at 7 h, which correlates with Treg-mediated suppression of CFSE dilution and cytokine expression in Tcons.
**Figure 4** In vitro expanded CD45RA^+ T\(_{\text{reg}}\) suppress CD3^+ T cell responses from inflamed Crohn’s MLN and LP.

(A) Suppression of proliferation of a single lot of freeze-thawed, allogeneic T\(_{\text{cons}}\), by freshly isolated PB CD4^+CD25^hiCD127^loCD45RA^+ T\(_{\text{reg}}\) or D24 CD45RA^+ T\(_{\text{reg}}\) that were expanded in vitro from these freshly isolated precursors. Pooled data from three sets of freshly isolated PB T\(_{\text{reg}}\) and subsequently expanded T\(_{\text{reg}}\) populations. Data points are mean ±SEM. (B) Fresh ileal resection specimen opened longitudinally to show ileal stricture (marked ‘S’) and proximal inflamed, haemorrhagic mucosa with deep ulceration (arrows). Scale bar: 2 cm. (C) Representative microscopic image from this resection showing mucosal distortion, ulceration (marked ‘U’) and transmural inflammation, including lymphoid aggregates (arrows). 12.5× H&E. Scale bar: 2 mm. (D) Representative FACS plots gated on live CD3^+ events, showing proliferation of MLN T\(_{\text{cons}}\) cultured alone (top left panel) or with T\(_{\text{reg}}\) at a 1:1 MLNMC:T\(_{\text{reg}}\) ratio (bottom left panel). Pooled data showing T\(_{\text{reg}}\)-mediated suppression of MLN CD3^+ proliferation (right panel, n=5). Box and whisker plot shows median, IQR and range. (E) Representative FACS plots gated on live MLN CD3^+ events showing CD154 expression on MLN T\(_{\text{cons}}\) cultured alone (top left panel) or with T\(_{\text{reg}}\) at a 1:1 MLNMC:T\(_{\text{reg}}\) ratio (bottom left panel). Pooled data showing T\(_{\text{reg}}\)-mediated suppression of CD154 expression in live MLN CD3^+ cells (right panel, n=5). (F) Representative FACS plots gated on live LP CD3^+ events showing CD154 expression on LP T\(_{\text{cons}}\) cultured alone (top left panel) or with T\(_{\text{reg}}\) at a 1:1 LPMC:T\(_{\text{reg}}\) ratio (bottom left panel). Pooled data showing T\(_{\text{reg}}\)-mediated suppression of CD154 expression in live LP CD3^+ cells (right panel, n=5). (D–F) Dotted line shows non-specific suppression from ‘2X control’.

Comparisons between observed suppression and non-specific suppression (tp<0.05, ttp<0.01, ttt tp<0.001, tttt tp<0.0001) and observed suppression and no suppression (zero, *p<0.05, **p<0.01, ***p<0.001 and ****p<0.0001) are shown. T\(_{\text{reg}}\), thymus-derived regulatory T cells; MLN, Mesenteric lymph node; LP, lamina propria; T\(_{\text{cons}}\), conventional CD4^+CD25^lo/int T cells; PB, peripheral blood; FACS, fluorescence-activated cell sorting; MLNMC, MLN mononuclear cell; LPMC, LP mononuclear cell; CTV, Cell Trace Violet.
at 96h.\textsuperscript{25,49} Significant dose-dependent suppression of CD154 expression in MLN and LP T cells was observed (figure 4E, F), demonstrating that in vitro expanded D24 CD45RA\textsuperscript{+} T\textsubscript{regs} suppress early activation of MLN and LP T\textsubscript{cons} in vitro.

**DISCUSSION**

There remains an unmet need to develop novel therapies for CD, as current drug treatments frequently fail to maintain long-term remission and may be complicated by significant side effects. Cellular therapies are emerging as potentially attractive therapeutic strategies. T\textsubscript{regs} are effective in preclinical models of colitis\textsuperscript{6} and phase 1 clinical trials suggest that in vitro expanded T\textsubscript{regs} are safe in the prophylaxis and treatment of GvHD\textsuperscript{12,24} and type 1 diabetes.\textsuperscript{18} We built on recent work to describe a method for isolation and expansion of T\textsubscript{regs} from Crohn’s blood that is readily transferable to a GMP background and addresses several barriers to the use of expanded T\textsubscript{regs} as an autologous cell-based therapy in this important disease.

T\textsubscript{regs} can be selected and expanded in vitro to clinically useful numbers under both R&D-grade\textsuperscript{11,13,16,21,23,26} and GMP conditions\textsuperscript{12,18,24} retaining an in vitro suppressive function before infusion into humans. We showed that it is feasible to do the same using T\textsubscript{regs} obtained from Crohn’s blood, including patients receiving thiopurines or anti-tumor necrosis factor (TNF) medications. Even after prolonged culture, these T\textsubscript{regs} maintained FOXP3 expression and suppressed activation of autologous T cells.

T cell lineage plasticity is well described. A major potential barrier to T\textsubscript{reg} therapy is the possibility that these cells might adopt an inflammatory phenotype and worsen inflammation on adoptive transfer. Freshly isolated thymus-derived T\textsubscript{regs} from both mice and humans can express proinflammatory cytokines and transcription factors (TF) canonical to effector CD4\textsuperscript{+} lineages, including IL-17\textsuperscript{39-44} and IFN-\gamma\textsuperscript{44} both of which are implicated in CD pathogenesis. Indeed, IL-17\textsuperscript{+} FOXP3\textsuperscript{+} T\textsubscript{regs} have been identified in non-inflamed human blood and lymphoid tissue,\textsuperscript{40} and inflamed Crohn’s mucosa.\textsuperscript{41} While there is some evidence that plastic cytokine and TF expression may license efficient T\textsubscript{reg} homing to, and suppression of, TH1-mediated and TH17-mediated inflammation,\textsuperscript{44,56} this may also lead to the generation of T\textsubscript{regs} with an effector phenotype that contribute to inflammation.

We and others have demonstrated that in vitro expanded T\textsubscript{regs} cultured in the presence of rapamycin have enhanced phenotypic stability.\textsuperscript{13,21} We show that as well as retaining their suppressive capacity, CD45RA\textsuperscript{+} T\textsubscript{regs} expanded from the blood of patients with CD in the presence of rapamycin do not express IL-17A or other TH17-related genes, even following exposure to proinflammatory cytokines that they would likely meet in inflamed intestinal mucosa. These data corroborate data from Hoffmann et al.\textsuperscript{47,48} in healthy controls, showing that expanded CD45RA\textsuperscript{+} T\textsubscript{regs} are resistant to the induction of proinflammatory cytokines on stimulation and highly express CD62L and CCR7, which are associated with phenotypic stability.

Freshly isolated CD45RA\textsuperscript{+} T\textsubscript{regs} have an epigenetically stable FOXP3 locus with extensive TSDR demethylation.\textsuperscript{56} TSDR demethylation correlates with stable FOXP3 expression in vitro\textsuperscript{56,57} and TSDR-mediated protection from autoimmunity in vivo\textsuperscript{57} in humans. However, the significance of TSDR demethylation for in vitro expanded T\textsubscript{regs} is poorly understood. Barzaghi et al.\textsuperscript{58} recently described a cohort of patients with ‘Immune dysregulation, Polyendocrinopathy, Enteropathy, X-linked syndrome (IPEX)-like syndrome’, severe multisystem autoimmunity in the absence of identifiable mutations in molecules implicated in T\textsubscript{reg} function, with decreased TSDR demethylation despite normal T\textsubscript{reg} numbers and in vitro suppression. This suggests that ex vivo expanded CD45RA\textsuperscript{+} T\textsubscript{regs} with incomplete TSDR demethylation, may have suboptimal biological activity in vivo, despite suppressive function in vitro. These data also suggest that CD45RA\textsuperscript{−} T\textsubscript{regs} are more likely to retain phenotypic stability and are less likely to acquire an effector phenotype than CD45RA\textsuperscript{−} T\textsubscript{regs} consistent with a more favourable safety profile of this T\textsubscript{reg} subset as a cell-based therapy for CD.

In order to be therapeutically effective, adoptively transferred T\textsubscript{regs} may need to traffic to intestinal lymphoid tissue or LP. Some groups have taken advantage of TCRs specific for luminal antigens to direct T\textsubscript{regs} to the intestinal mucosa, such as IL-10-producing T cell clones with ovalbumin-specific TCRs,\textsuperscript{59} or T cells with transgenic Cblr1 flagellin-specific TCRs.\textsuperscript{60} Alternatively, T\textsubscript{reg} expansion in the presence of ATRA induces αβ integrin expression but also increases effector cytokine expression, such as IL-17 and IFN-γ, potentially limiting its use in GMP cell expansion.\textsuperscript{13,21} We show that CD45RA\textsuperscript{+} T\textsubscript{regs} expanded in the presence of IL-2 and rapamycin highly express CD62L and CCR7, allowing homing to, and anatomical orientation within lymphoid tissue.\textsuperscript{60,61} T\textsubscript{reg} CD62L expression is also required for T\textsubscript{reg}-mediated cure of GvHD.\textsuperscript{30} CD45RA\textsuperscript{+} T\textsubscript{regs} also expressed CCR4, required for T\textsubscript{reg}-mediated prevention of CD45RB\textsuperscript{62} colitis. Interestingly, murine T\textsubscript{regs} do not need to home to intestinal LP to prevent CD45RB\textsuperscript{63} adoptive transfer colitis. β integrin-null T\textsubscript{reg} home to MLN and prevent colitis in this model, despite almost undetectable LP homing.\textsuperscript{63} Consequently, the ability to home to MLN is highly desirable in potentially therapeutic cells.

CD45RA\textsuperscript{−} T\textsubscript{regs} also express αβ integrin and CXC motif receptor 3 (CXCR3), indicating an ability to home to LP and sites of inflammation, respectively. Moreover, we used a human small intestinal xenotransplant model to show, for the first time, that in vitro expanded CD45RA\textsuperscript{+} T\textsubscript{regs} from patients with CD home to inflamed human gut in vivo. Xenotransplanted SB segments develop into tissue that is morphologically and functionally identical to normal gut and is capable of peristalsis and nutrient absorption.\textsuperscript{63,64} The xenografts also possess a chimeric endothelium that expresses human MadCAM-1.\textsuperscript{64} This is the first demonstration that this model can be used in the assessment of immune cell homing.

Xenograft-bearing mice received rhIL-2 in order to support survival of adoptively transferred human T\textsubscript{regs},\textsuperscript{23} as murine IL-2 is less efficient at promoting proliferation of human T cells than rhIL-2, despite cross-reactivity.\textsuperscript{65} As recent phase 1 trials of in vitro expanded T\textsubscript{regs} in GvHD and type 1 diabetes mellitus showed signs of clinical efficacy without supplemental rhIL-2, it is likely that this is a feature of the experimental system and will not be required in clinical trials in Inflammatory bowel disease (IBD).\textsuperscript{65}

Future work will include ‘humanising’ xenograft-bearing mice and developing additional techniques to induce xenograft inflammation, thus allowing us to assess the functional impact of CD45RA\textsuperscript{+} T\textsubscript{regs} on gut inflammation. The percentage of LP human T cells that could be recovered from human bowel transplants was relatively modest compared with the percentage of T cells recovered from the spleen. Given that the expression of the gut homing integrin α5β\textsubscript{7} was only expressed on ~20% of the purified T\textsubscript{regs}, future work may need to address methods to increase α5β\textsubscript{7} expression, such as the use of retinoic acid, as we have previously shown.\textsuperscript{21}

An additional barrier to T\textsubscript{reg} therapy in CD is that effector T cells from the diseased mucosa of patients with CD may be resistant to the suppressive action of T\textsubscript{regs}. Indeed, we previously
showed that Tcons isolated from inflamed Crohn’s mucosa are relatively resistant to Treg-mediated suppression, due to overexpression of Smad7, an inhibitor of TGF-β signalling. In this study, we utilised Tregs cultured in the presence of rapamycin, which has been shown to enhance the suppressive ability of in vitro expanded Tregs, compared with Tregs freshly isolated from the same donor and shown that in vitro expansion enhances the suppressive ability of Tregs obtained from CD PB. Rapamycin-expanded CD45RA+ Tregs effectively suppress both MLN and PT cells obtained from inflamed Crohn’s resection specimens. These data suggest that in vitro expanded CD45RA+ Tregs may modulate immune responses in niches directly relevant to the pathogenesis of CD. Tregs use multiple mechanisms to suppress in vitro and in vivo, including contact-dependent mechanisms (CTLA-4, perforin-granzyme B) and contact-independent mechanisms (IL-10, TGF-β, extracellular ATPase activity via CD39/CD73 etc). Sakaguchi et al has proposed a multistep model of in vitro suppression that initially requires cell-cell contact but is subsequently contact independent. The mechanism of suppression of erstwhile ‘resistant’ mucosal Tcons by in vitro expanded Tregs is currently unknown and will be the subject of further study. In addition, not all of the patients in this study had active disease, so it will be important to extend these data further to broaden the therapeutic relevance of these findings. However, a substantial proportion of the patients in this study did have evidence of disease activity (n = 5/13), which did not affect either Treg expansion or function.

In conclusion, we have shown that in vitro expanded CD45RA+ Tregs are likely to be the most suitable Treg subset for cellular therapeutics in CD. This subset is readily expandable to sufficiently high numbers under conditions that are readily transferable to GMP, for clinical use. They express an appropriately therapeutically challenging disease.

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REFERENCES


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