

# CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup>Foxp3<sup>+</sup> T cells are cytotoxic for human neurons

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## ABSTRACT

MS lesions are characterized by destruction of myelin and significant neuronal and axonal loss. Preliminary studies with the use of T<sub>regs</sub> in the mouse model of MS have been extremely encouraging. However, recent studies with human cells have shown the presence of different subpopulations of T cells within the CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cell phenotype, some of which do not have regulatory functions. These findings suggest a potential difference between mouse and human in the regulatory phenotype. Here, we show that human activated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells are neurotoxic *in vitro*. These cells expressed high levels of the cytotoxic molecule GrB and had no suppressive effect. On the contrary, they produced IFN- $\gamma$  and low IL-17, suggesting a shift toward a T<sub>H</sub>1 phenotype. Thus, our data confirm the presence of a nonregulatory cytotoxic subpopulation within the human CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells and suggest further studies on the human regulatory phenotype prior to any potential therapeutic application. *J. Leukoc. Biol.* 89: 927–934; 2011.

## Introduction

Infection and autoimmunity can trigger the activation of the effector arm of the immune system. This activation is characterized by the differentiation of T<sub>H</sub>1, T<sub>H</sub>2, or T<sub>H</sub>17 effector cells. Evidence exists that immune tolerance is maintained or promoted by another subpopulation of T cells, T<sub>regs</sub> [1], a function that has encouraged the potential therapeutic use of T<sub>regs</sub> in autoimmune diseases such as MS. Although the CNS is considered to be an immunologically privileged site, it is now known that this immune privilege is far from absolute [2]. Activated T cells transmigrate across the blood brain barrier, enter the CNS compartment, and accumulate as well as proliferate in response to antigen restimulation [3, 4]. Infiltration of these immune cells characterizes lesions of inflammatory dis-

eases of the CNS such as MS. Within MS lesions, there is an almost complete destruction of myelin, which ensheathes neuronal axons. As a consequence, MS has been classically described as a demyelinating disease of the CNS. The recent re-discovery of axonal and neuronal injury in MS [5] has initiated a shift in the way we think about this disease [3]. Therefore, MS is now considered a demyelinating and a neurodegenerative disease. Little is known about the causes and mechanisms that lead to myelin and/or axonal injury in MS [6]. Pathological studies have shown a correlation among demyelination, axonal damage [7], and infiltrating CD8<sup>+</sup> T cells and macrophages [8]. Furthermore, in the animal model of MS, EAE, the destabilization of a microtubule network, colocalized with the infiltration of immune cells [9]. The presence of CD4<sup>+</sup> and CD8<sup>+</sup> T cells in MS and EAE lesions is proposed to mediate demyelination and axonal injury [10]. In an *in vitro* study, Giuliani et al. [11] reported that human T cells, activated by a soluble anti-CD3 antibody, induce neurotoxicity in allogeneic and syngeneic systems. It has been shown previously that CTLs expressing the serine protease GrB were found frequently in close contact with injured axons within actively demyelinating lesions [12]. Activated T cells release soluble factors such as GrB [13] and proinflammatory cytokines and are thought to mediate axonal and myelin injury [14]. Microarray-based analysis revealed that the expression of GrA and GrB was increased in cerebrospinal fluid of MS patients [15]. However, GrB is best characterized because of its strong proapoptotic activities [13].

In normal conditions, regulatory mechanisms are in place to prevent autoreactive T cells from causing an autoimmune disease. T<sub>regs</sub> have been shown to be crucial for maintenance of self-antigens and tolerance even in post-transplantation [1]. CD4<sup>+</sup>CD25<sup>+</sup> T cells are endowed with suppressor functions and characterized by high expression of Foxp3 [16]. *In vivo*, CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T<sub>regs</sub> are described as neuroprotective after brain ischemia [17]. In the experimental allergic encephalomyelitis, it has been shown that T<sub>regs</sub> may prevent the traffick-

Abbreviations: EAE=experimental autoimmune encephalomyelitis, Foxp3=forkhead box transcription factor, GrB=granzyme B, HFN=human fetal neuron, HPBMC=human PBMC, MAP-2=microtubule-associated protein-2, MS=multiple sclerosis, T<sub>reg</sub>=regulatory T cell

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ing of effector cells into the CNS [18]. On the other hand, under neurodegenerative conditions in the CNS of two mice strains (C57BL/6J and BALB/c/OLA), CD4<sup>+</sup>CD25<sup>+</sup> T<sub>regs</sub> displayed opposite, beneficial and destructive, effects [19]. Indeed, according to other reports, the induction and importance of T<sub>regs</sub> and their regulatory pathways in human disease are not thoroughly understood. Miyara et al. [20] have shown recently that human Foxp3<sup>+</sup> T cells can be dissected in at least three subpopulations, and not all of them have a regulatory phenotype.

In this study, we investigate the potential role of CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells in inflammatory-mediated neurodegeneration. In particular, we examine the differential neurotoxicity and expression of GrB by the different T cell subsets CD4<sup>+</sup>, CD8<sup>+</sup>, and CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup>Foxp3<sup>+</sup> T cells. The results described here show that human CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup>Foxp3<sup>+</sup> T cells, when activated, produce high levels of GrB and IFN- $\gamma$  and induce neuronal killing in vitro. This study shows for the first time that the CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup>Foxp3<sup>+</sup> T cell population includes effector T cells able to induce neurotoxicity. These findings warrant further studies about the human T cell phenotype currently attributed to T<sub>regs</sub>.

## MATERIALS AND METHODS

### Isolation and culture of HFNs and T cells

Human brain tissue was obtained from legally and medically aborted 15 to 20 week-aged fetuses, according to the guidelines approved by the local institutional ethics committee. HFNs and HPBMCs were isolated and cultured as reported previously [11]. The maturity of HFNs has been described previously [21]. T cell subsets were negatively sorted into CD4<sup>+</sup> or CD8<sup>+</sup>; then, CD4<sup>+</sup>CD25<sup>+</sup> or CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> subpopulations were positively selected from the whole CD4<sup>+</sup> T cells using magnetic cell sorting (MACS separation columns, Miltenyi Biotec, Germany), according to the manufacturer's protocol.

### RNA isolation and real-time RT-PCR

Cultured cells were lysed and homogenized in TRIzol (Invitrogen, Carlsbad, CA, USA). Total RNA was collected and purified using an RNeasy mini kit, according to the stated protocol (Qiagen, Valencia, CA, USA). cDNA was synthesized using oligo-d(T) and Superscript II RT (Invitrogen), according to the manufacturer's recommended instructions. Semiquantitative RT-PCR was carried out using Bio-Rad iQ SYBR Green Supermix (Bio-Rad Laboratories, Hercules, CA, USA) on the iQ5 program or i-Cycler (Bio-Rad Laboratories), according to the manufacturer's protocol. Primers for GrB (forward: GCG GTG GCT TCC TGA TAC AAG, reverse: CCC CCA AGG TGA CAT TTA TGG) and Foxp3 (forward: CAA GTT CCA CAA CAT GCG AC, reverse: ATT GAG TGT CCG CTG CTT CT) amplifications were used at an annealing temperature of 60°C.

### Immunocytochemistry

HFNs and T cells (1:1) were cocultured for 24 h and then fixed with 4% PFA for ~20 min. The viability of neurons was assessed using mouse primary mAb against human MAP-2 as reported previously [11].

### Flow cytometry

The intracellular protein levels of GrB, Foxp3, and IL-17 were assessed using flow cytometry. Activated or unactivated CD4<sup>+</sup>CD25<sup>+</sup> or CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim/-</sup> (CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup>) T cells were incubated with 1  $\mu$ g/ml ionomycin and 10 ng/ml PMA (both Sigma-Aldrich, Oakville, ON, Canada) at 37°C and

5% CO<sub>2</sub> for 2 h. A mixture of transport inhibitors [3  $\mu$ g/ml Brefeldin A (Sigma-Aldrich) and 2  $\mu$ M Monensin solution (eBioscience, San Diego, CA, USA)] was added into each well, and the cell culture was incubated further for 2 h. Cells were harvested and went through several washing steps with PBS, followed by incubation in freshly prepared eBioscience Foxp3 fixation/permeabilization buffer for 30 min at 4°C in dark. After 2 $\times$  washing in permeabilization buffer, cells were incubated with intracellular staining for GrB, Foxp3, and IL-17 (all eBioscience) for 45 min–2 h at 4°C. Stained cells were washed 2 $\times$  in permeabilization buffer, resuspended in PBS + 2% FBS, and analyzed on a FACSCanto flow cytometer using FACSDiva software for acquisition (BD Biosciences, San Diego, CA, USA). Multicolors were compensated using Compbeads (BD Biosciences). Cytometry data were analyzed using FlowJo 7.2.2 software (Tree Star, Ashland, OR, USA).

### Suppression assay

To examine the suppressive function, 50,000 CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim/-</sup> T cells were cocultured for 24 h with 50,000 CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim/-</sup> T cells. AIM-V media (25–30  $\mu$ l), containing 1  $\mu$ Ci <sup>3</sup>H-thymidine, were added into each well. After incubating for 18 h at 37°C, the cells were harvested, and the radioactivity was measured. Finally, the stimulation index of the wells was calculated, and then, the CPM of proliferating cells were analyzed.

### Cytokine-ELISA

The phenotype of CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells was evaluated using human IFN- $\gamma$ -ELISA (Invitrogen), according to the manufacturer's instruction.

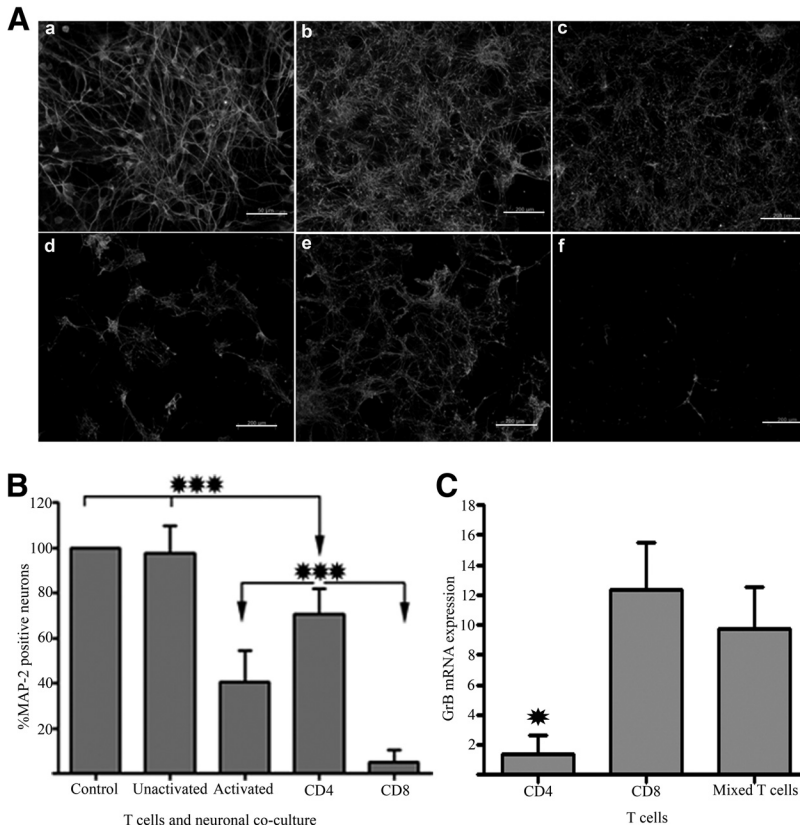
### Statistical analysis

Results were statistically analyzed as mean  $\pm$  SD using GraphPad Prism 5. The groups were compared using one-factor ANOVA followed by Tukey post hoc test for normally distributed data. Two-tailed unpaired *t* test was applied to compare two groups with normally distributed data, and data were presented as mean  $\pm$  SD. *P* values of <0.05 were considered significant. Asterisks correspond to \**P* < 0.05, \*\**P* < 0.01, and \*\*\**P* < 0.001.

## RESULTS AND DISCUSSION

### Activated T cell subsets show different levels of neurotoxicity

We have reported previously that both subpopulations of CD4<sup>+</sup> and CD8<sup>+</sup> T cells display a neurotoxic effect in allogeneic and syngeneic systems [11]. In the present study, we have further evaluated and compared the differential neurotoxic effect and GrB expression of different subsets of human T cells such as CD4<sup>+</sup>, CD8<sup>+</sup>, and CD4<sup>+</sup>CD25<sup>+</sup> T cells. A neuronal killing assay was performed using HFNs cocultured with T cells. T lymphocytes were plated on anti-CD3/anti-CD28-immobilized or control uncoated plates for 3 days before adding them to HFNs. After 72 h in culture, neurons in the control group were treated with only AIM-V medium, without exposure to T cells. Conversely, in the experimental groups, unactivated, activated HPBMCs, and activated CD4<sup>+</sup> or CD8<sup>+</sup> T cells were added to neurons in a 1:1 ratio and kept for 24 h. The neurotoxic effect of T cells was measured by immunocytochemistry using anti-MAP-2 antibody. MAP-2 was chosen as a marker of neuronal viability, as the disappearance of its immunoreactivity has been associated with neuronal damage and death [11, 22]. As indicated in Fig. 1A, a–c, the density of neurons, untreated or treated with unactivated T cells, remained unchanged. In contrast, the density of neurons in the cul-



**Figure 1. Neuronal killing assay by T cells activated after sorting.** Whole human T lymphocytes or subsets sorted using magnetic beads into  $CD4^+$  or  $CD8^+$  T cells were activated using anti-CD3/anti-CD28 antibody for 3 days. HFNs were cocultured with (A, a) only AIM-V media as a control ( $\times 40$ ), (A, b) lower magnification of A, a ( $\times 10$ ), (A, c) unactivated T cells, (A, d) activated mixed population of T cells, (A, e) activated  $CD4^+$  T cells, and (A, f) activated  $CD8^+$  T cells. The culture was immunocytochemically stained with the neuronal marker anti-MAP-2. Data are representative of at least three individual experiments. (B) Manual quantification of MAP-2-positive neurons, demonstrating the differential neurotoxic effects of a mixed T cell population,  $CD4^+$ , or  $CD8^+$  T cells. The most severe neurotoxicity was documented in the group treated with  $CD8^+$  T cells ( $P < 0.001$ ). Data are representative of three individual experiments. (C) Total RNA was extracted from the mixed T lymphocytes,  $CD4^+$ , or  $CD8^+$  T cells to measure GrB mRNA expression. RT-PCR revealed that  $CD4^+$  T cells expressed significantly less GrB transcript than the rest of the groups ( $P < 0.05$ ). Data are representative of at least two individual experiments. Note that the expression of GrB in the unactivated group was negligible so that the scale and comparison were based on  $CD4^+$  T cells [original scale bars: (A, a)  $50\ \mu\text{m}$ ; (A, b-f)  $200\ \mu\text{m}$ ].

ture treated with activated T cells was decreased significantly ( $37.4 \pm 14.5\%$  vs.  $100\%$  in the control group). Despite significant neuronal loss, the group treated with  $CD4^+$  T cells showed a significant amount of surviving neurons ( $69.6 \pm 11.9\%$ ). However,  $CD8^+$  T cells induced severe neuronal injury, resulting in only a few surviving neurons ( $7.3 \pm 5.9\%$ ;  $P < 0.001$ ; Fig. 1A and B). In contrast with our previous findings [11], where  $CD4^+$  and  $CD8^+$  T cells induced comparable neuronal loss, these results show that the majority of the cytotoxic effect was induced by  $CD8^+$  T cells. This discrepancy can be explained by the different activation system that we used in this study (immobilized anti-CD3 vs. soluble OKT3 in the previous work).

It is known that effector T cells express GrB upon activation in the presence of costimulatory molecules or cytokines or during pathogenic infections. It has been shown previously that GrB can mediate neuronal apoptosis [13]. In a rat model of cerebral ischemia, GrB mediated post-ischemic neuronal death [23]. Conversely, the absence of GrB has been associated with decreased neuronal injury [24]. We measured the expression of GrB in different subsets of T cells. Subpopulations of T cells ( $CD4^+$  or  $CD8^+$ ) were activated, and the total RNA was extracted. Although all of the cells expressed GrB after activation, RT-PCR revealed that  $CD4^+$  T cells displayed very low levels of GrB ( $1.35 \pm 1.31$ ) compared with  $CD8^+$  and activated HPBMCs, which showed significantly higher levels of GrB ( $12.38 \pm 3.07$  and  $9.70 \pm 2.83$ , respectively; Fig. 1C;  $P < 0.001$ ). The expression of GrB in the unactivated T cells was negligible so that the scale and comparison were based on  $CD4^+$  T cells. The levels of GrB expression correlated directly

with the neurotoxic effect of  $CD4^+$ ,  $CD8^+$ , and the activated whole T cell population (Fig. 1A and B). Neuronal killing by GrB containing conditioned supernatant from  $CD4^+$  and  $CD8^+$  T cells has been reported previously [13]. However, this supernatant-mediated neuronal killing was very mild ( $20\%$ ) compared with the T cell-mediated killing of  $>60\%$  in the present study. Using a transwell system, where neurons and T cells were cocultured in two separate compartments, divided by a porous membrane allowing only the exchange of soluble factors, we have reported previously that no neurotoxicity occurred [11]. These findings underline the importance of cell-cell contact for effective neurotoxicity.

### $CD4^+CD25^+$ T cells induce severe neurotoxicity

$CD4^+CD25^+$  T cells are reported frequently for their role in modulating a diverse array of immunological responses and maintenance of peripheral tolerances by acting on the effector cells. They are characterized by high expression of Foxp3 [16]. Thus, we hypothesized that these cells would suppress the severe neurotoxicity induced by  $CD8^+$  T cells. To validate this hypothesis,  $CD4^+$  and  $CD8^+$  T cells were negatively selected through magnetic separation from HPBMCs and activated for 72 h using an anti-CD3 antibody.  $CD25^+$  T cells were positively isolated from  $CD4^+$  T cells after activation. A mixture of  $CD4^+CD25^+$  and  $CD8^+$  T cells ( $1:1$  ratio) was added to HFN cultures. Cocultures of  $CD8^+$  or  $CD4^+CD25^+$  T cells with HFNs were used as controls. Immunocytochemical analysis using anti-MAP-2 antibody revealed that  $CD4^+CD25^+$  T cells were not able to suppress the cytotoxic effect induced by

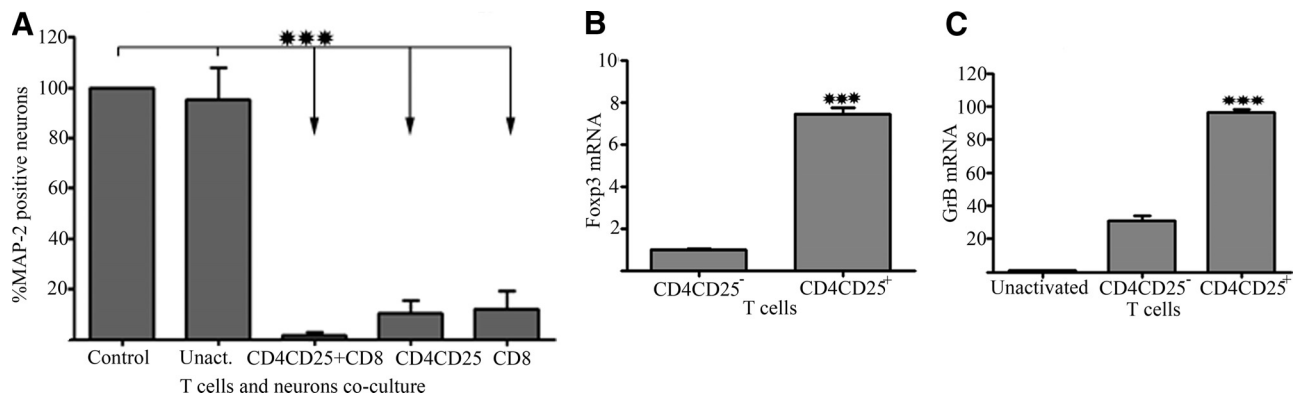
CD8<sup>+</sup> T cells; instead, they induced a neuronal killing comparable with CD8<sup>+</sup> T cells (data not shown). The failure of CD4<sup>+</sup>CD25<sup>+</sup> T cells in preventing neuronal killing can be explained in two ways: first, when CD4<sup>+</sup>CD25<sup>+</sup> and CD8<sup>+</sup> T cells were mixed at the same time of the coculture with neurons, there was not enough time to induce a suppressive response to prevent neuronal killing; second, CD25 is expressed by activated T cells [25], thus activation of CD4<sup>+</sup>CD25<sup>+</sup> T cells could enhance proliferation of non-T<sub>regs</sub> [26]. Therefore, to minimize these confounding effects, CD25<sup>+</sup> T cells were positively isolated from CD4<sup>+</sup> T cells before activation. CD4<sup>+</sup>CD25<sup>+</sup> T cells were activated alone or in a mixture (1:1), with the negatively selected CD8<sup>+</sup> T cells for 72 h and cocultured with HFNs, as stated above. Following 24 h coculture, neurons were stained with MAP-2 for viability assessment. Surprisingly, almost 90% of neurons were killed in the control group treated with CD4<sup>+</sup>CD25<sup>+</sup> T cells only. The count of MAP-2-positive viable neurons resulted in  $10.5 \pm 4.8\%$  in CD4<sup>+</sup>CD25<sup>+</sup> and  $11.9 \pm 7.2\%$  in CD8<sup>+</sup> T cell-treated groups, respectively. Furthermore, the percent of surviving neurons in the coculture with the combination of CD4<sup>+</sup>CD25<sup>+</sup> plus CD8<sup>+</sup> T cells was only  $1.6 \pm 1.3\%$  (Fig. 2A;  $P < 0.001$ ). The results indicate that prior incubation of these two subsets of T cells is not able to inhibit the severe CD8<sup>+</sup>-mediated neurotoxicity. In addition, our findings clearly show that CD4<sup>+</sup>CD25<sup>+</sup> T cells are also neurotoxic after activation.

Foxp3 has been identified as a specific and most up-to-date marker of T<sub>regs</sub> [27] and is required for their development [28]. To characterize further whether CD4<sup>+</sup>CD25<sup>+</sup> T cells acquire a regulatory phenotype, a RT-PCR experiment was performed using a Foxp3 primer. In this case, two sets of cells, CD25<sup>+</sup> and CD25<sup>-</sup> depleted, were selected from CD4<sup>+</sup> T cells

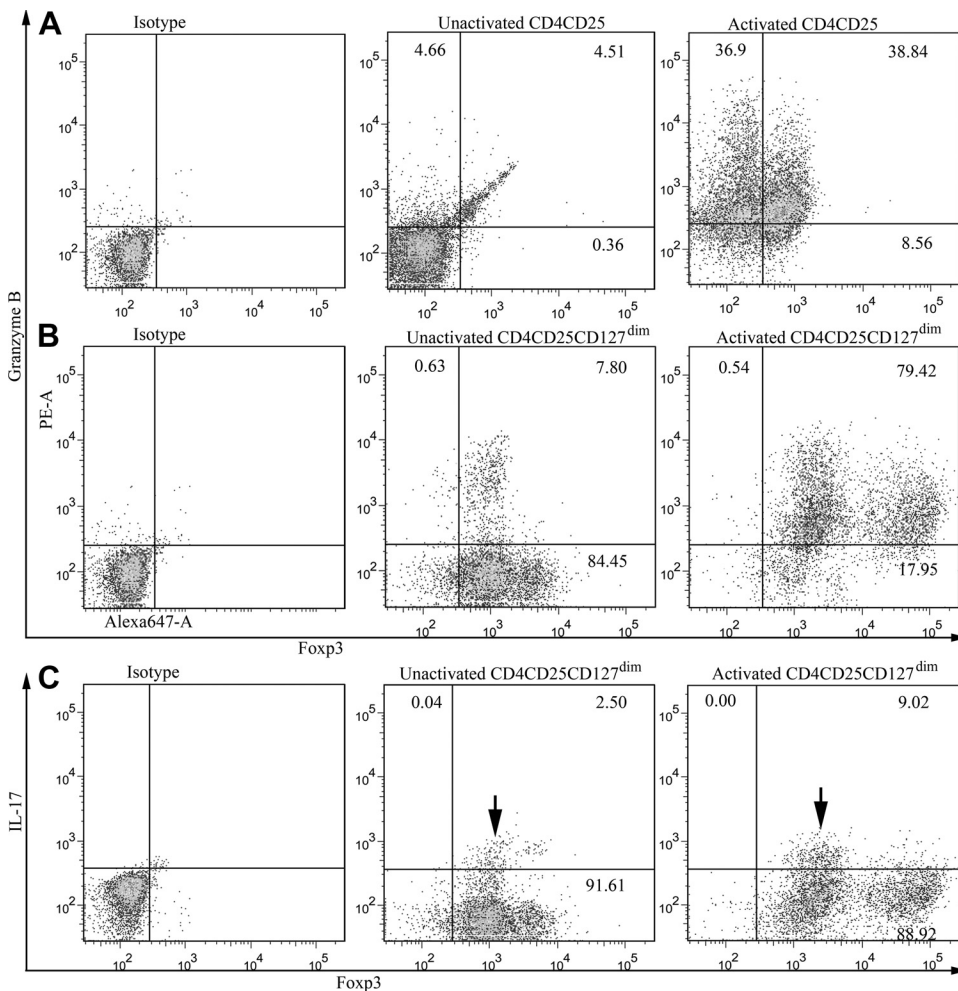
and activated for 72 h. Following RNA extraction, RT-PCR results demonstrated that Foxp3 expression was 7.45-fold increased in activated CD4<sup>+</sup>CD25<sup>+</sup> compared with CD25-depleted CD4<sup>+</sup> T cells (Fig. 2B;  $P < 0.001$ ). This was in line with earlier findings, where CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed high levels of Foxp3 compared with CD4<sup>+</sup>CD25<sup>-</sup> [29]. It has been shown that T<sub>regs</sub> use GrB to kill target effector cells [30]. In our study, we assessed GrB expression by CD25<sup>+</sup> T cell subsets. Interestingly, RT-PCR revealed that activated CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed ~66-fold more GrB than activated CD4<sup>+</sup>CD25<sup>-</sup> T cells and above 96-fold change more than the unactivated T cells (Fig. 2C;  $P < 0.001$ ). GrB expression in these cells was extremely high, and this may account for the extreme neurotoxic effect observed in our experiments.

### CD4<sup>+</sup>CD25<sup>+</sup> T cells express GrB and Foxp3 but low IL-17

As described above, RT-PCR revealed that CD4<sup>+</sup>CD25<sup>+</sup> T cells express Foxp3 and GrB at a transcript level (Fig. 2B and C, respectively). To confirm whether these high expressions translated into high-protein production, cells underwent intracellular staining for Foxp3 and GrB. Flow cytometric analysis indicated that only 9% of unactivated CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed GrB, whereas 4.51% were positive for GrB and Foxp3 (Fig. 3A, middle panel). However, after activation, ~76% of CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed GrB. In addition, 38.84% of activated CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed GrB and Foxp3, and ~47% were positive for Foxp3. On the other hand, >50% of activated CD4<sup>+</sup>CD25<sup>+</sup> T cells were negative for Foxp3 expression (Fig. 3A, right panel). As the neurotoxic effect could have been related to the Foxp3-negative fraction, we further purified Foxp3<sup>+</sup> cells. The absence of a surface marker for Foxp3



**Figure 2. Neuronal killing assay by postselection-activated CD4<sup>+</sup>CD25<sup>+</sup> T cells.** Human T lymphocytes were positively sorted into CD4<sup>+</sup>CD25<sup>+</sup> subsets of T cells using magnetic-conjugated antibody and activated for 3 days by anti-CD3/anti-CD28 antibody prior to coculture with neurons. (A) HFNs were treated with the coculture media as a control (left-most bar) or cocultured with unactivated (Unact.), activated in a mixture of CD4<sup>+</sup>CD25<sup>+</sup> and CD8<sup>+</sup> T cells (1:1), postselection-activated CD4<sup>+</sup>CD25<sup>+</sup> T cells, or CD8<sup>+</sup> T cells (only 50%). After 24 h in culture, cells were immunocytochemically stained using MAP-2 and manually quantified. As demonstrated in the graph, CD4<sup>+</sup>CD25<sup>+</sup> T cells failed to reverse the CD8<sup>+</sup> T cell-induced neurotoxicity ( $P < 0.001$ ). Data are representative of three individual experiments. (B) Expression of the transcription factor Foxp3 by CD4<sup>+</sup>CD25<sup>+</sup> T cells. Total RNA was extracted from each group (activated CD4<sup>+</sup>CD25<sup>+</sup> or activated CD4<sup>+</sup>CD25<sup>-</sup>) to measure the Foxp3 mRNA expression. RT-PCR revealed that CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed above sevenfold more Foxp3 in comparison with CD4<sup>+</sup>CD25<sup>-</sup> T cells ( $P < 0.001$ ). Data are representative of three individual experiments. (C) GrB expression was measured, and activated CD4<sup>+</sup>CD25<sup>+</sup> T cells expressed more than 96- and 60-fold increases compared with unactivated whole T cell populations and activated CD4<sup>+</sup>CD25<sup>-</sup> T cells, respectively ( $P < 0.001$ ). Data are representative of at least two individual experiments.



**Figure 3. Showing the expressions of intracellular GrB, Foxp3, and IL-17.** (A) Flow cytometry data showing CD4<sup>+</sup>CD25<sup>+</sup> T cells expressing GrB and Foxp3. Only 4.51% of unactivated CD4<sup>+</sup>CD25<sup>+</sup> T cells were positive for GrB and Foxp3 (middle panel), whereas the expression of GrB and Foxp3 was increased significantly in the activated CD4<sup>+</sup>CD25<sup>+</sup> T cells, but still more than half of the cells were negative for Foxp3 (right panel, Activated CD4CD25). (B) Foxp3<sup>+</sup> T cells were enriched further by depletion of CD127<sup>high</sup> T cells so that close to 97% of the cells were Foxp3-positive (right panel). However, the Foxp3 expression is bimodal. In activated CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells, 79.42% of the cells were positive for GrB and Foxp3. (C) Micrograph showing expression of IL-17 and Foxp3 by CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells. Among activated CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells, ~98% expressed Foxp3, whereas only 9.02% were positive for the intracellular IL-17 staining (right panel). In unactivated and activated CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells, the majority of IL-17-expressing cells was Foxp3<sup>low</sup> (arrows). Data are representative of three individual experiments.

limits the isolation of pure Foxp3<sup>+</sup> T cells. It has been shown previously that CD127 (surface marker) is down-regulated in human T<sub>regs</sub> and inversely correlated to Foxp3 expression [31]. Consequently, the CD127<sup>high</sup> cells were eliminated from the CD4<sup>+</sup>CD25<sup>+</sup> T cell population using magnetic separation. Flow cytometry showed that 92% unactivated and 98% activated CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells were Foxp3<sup>+</sup>, and 79.42% were positive for GrB and Foxp3 (Fig. 3B). By selecting CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup>, the purity of Foxp3<sup>+</sup> T cells reached 98% (Fig. 3B and C, right panels). However, the distribution of Foxp3-expressing cells was bimodal. This dual expression of Foxp3 has been described recently by Miyara et al. [20], who showed that three subpopulations of cells were present within the Foxp3<sup>+</sup>CD127<sup>dim</sup> population. Nevertheless, ~80% of both subpopulations (Foxp3 low or high) of activated CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells expressed a high level of GrB (Fig. 3B, right panel).

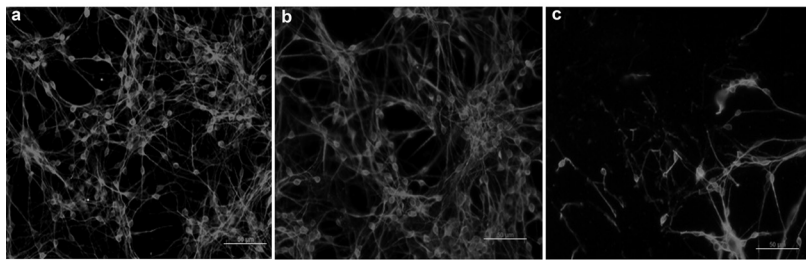
Plasticity is a characteristic of CD4<sup>+</sup> T cell subsets [32], and there is increasing evidence that certain subsets of T cells, defined by their function and transcription factors, are not stable. It has been shown previously that T cells with a T<sub>reg</sub> phenotype can shift to a T<sub>H</sub>1, T<sub>H</sub>2 [32], and T<sub>H</sub>17 phenotype [33]. Furthermore, we have reported previously that T<sub>H</sub>17 cells express high levels of GrB and induce neuronal killing

[34]. In addition, T<sub>H</sub>17 and T<sub>regs</sub> share a common lineage so that they may demonstrate plasticity toward each other in the presence of IL-6 or IL-10, respectively. Therefore, an intracellular staining for IL-17, a cytokine expressed by T<sub>H</sub>17 T cells, was performed on unactivated or activated CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells. Out of 98% activated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells, 9.02% expressed IL-17. Interestingly, IL-17-expressing cells were mainly from the Foxp3<sup>low</sup> subpopulations (Fig. 3C, arrows). CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim/-</sup> T cells are enriched for Foxp3, so we will refer to them as CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup>.

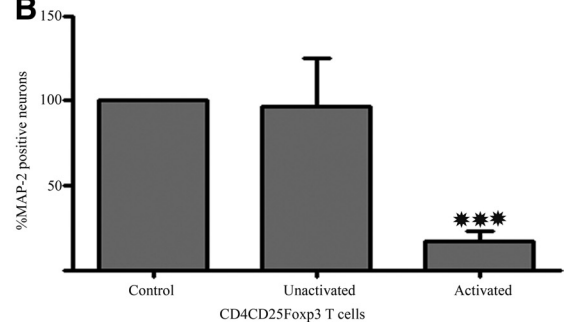
### Purified CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells are neurotoxic

To exclude the possibility that the neurotoxic effect induced by CD4<sup>+</sup>CD25<sup>+</sup> T cells could be caused by the Foxp3-negative fraction of the cells shown in Fig. 3A, right panel, a neuronal killing assay was performed using purified CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells displayed in Fig. 3B and C. HFNs were treated with AIM-V serum-free media (untreated control), cocultured with unactivated or activated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells. After 24 h in culture, immunocytochemical staining for MAP-2 revealed that neurons treated only with media or unactivated T cells remained healthy and dense (Fig. 4A, a and b, respectively). In contrast, the group treated with activated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells showed

A



B



**Figure 4. Micrographs showing that  $CD4^+CD25^+Foxp3^+$  T cells induced neuronal killing.** After 3 days activation,  $CD4^+CD25^+Foxp3^+$  T cells were cocultured with HFNs for 24 h, followed by immunocytochemical staining with MAP-2. HFNs were treated with: (A, a) AIM-V media (control), (A, b) unactivated, or (A, c) activated  $CD4^+CD25^+Foxp3^+$  T cells. (B) MAP-2-stained neurons were enumerated, and the viability of neurons in A, c (treated with activated  $CD4^+CD25^+Foxp3^+$  T cells) was significantly affected ( $P < 0.001$ ). Data are representative of three individual experiments (original scale bars: 50  $\mu$ m).

severe neuronal loss (Fig. 4A, c). Quantification of the above qualitative observation confirmed that only  $16.6 \pm 6.5\%$  MAP-2-positive, viable neurons were present in the group treated with activated  $CD4^+CD25^+Foxp3^+$  T cells (Fig. 4B;  $P < 0.001$ ). The loss of  $>80\%$  of neurons clearly showed that these cells were significantly neurotoxic, although they expressed a  $T_{reg}$  profile.

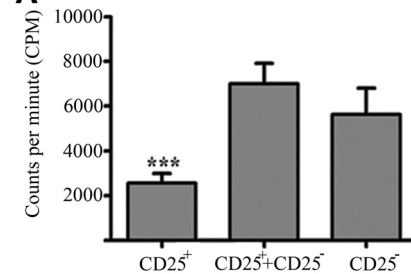
As indicated in Fig. 3B, the high level of GrB expression was reproduced in the  $CD4^+CD25^+Foxp3^+$  T cell population. The GrB released by activated T cells has been shown to induce cytotoxicity on human neurons [35]. GrB and perforin are the most crucial effectors of  $T_{reg}$ -mediated cytolysis. In a GrB-deficient mouse model, the function of  $T_{regs}$  was reduced compared with NK and  $CD8^+$  T cells [36]. Up-regulation of GrB expression in murine  $CD4^+CD25^+$   $T_{regs}$  correlated with induction of regulatory activity [37]. This reflects the importance of GrB for  $T_{reg}$  cytolysis activity. Given that GrB induces neurotoxicity [13], and  $T_{regs}$  use GrB to suppress their target cells [36] using the granular pathway [38] of cytotoxicity, we can speculate that in this study, GrB plays a role in the  $CD4^+CD25^+Foxp3^+$  T cell-mediated killing of human cortical neurons. This possibility is supported by the finding that the  $CD4^+CD25^+Foxp3^+$  T cells showed above 96-fold increases of GrB expression. In addition,  $\sim 80\%$  of the cells were positive for GrB and Foxp3.

### Activated $CD4^+CD25^+Foxp3^+$ T cells lack suppressive functions and express IFN- $\gamma$

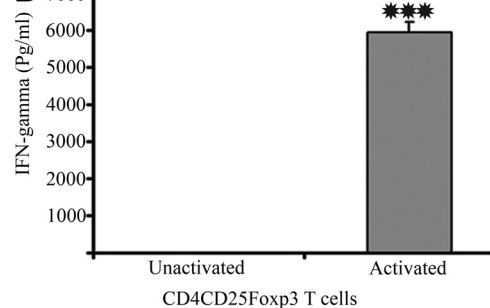
$T_{regs}$  maintain tolerance by suppressing the immune reaction of effector cells. The suppressive action of  $T_{regs}$  occurs by different mechanisms, such as inhibitory cytokines (e.g., IL-10, IL-35, and TGF- $\beta$ ), cytolysis, metabolic disruption, and modulation of DC maturation or function [39]. To address whether  $CD4^+CD25^+Foxp3^+$  T cells acquire a suppressive function,  $CD4^+CD25^+CD127^{dim}$  were cocultured with  $CD4^+CD25^-CD127^{dim}$  in a 1:1 ratio. Control groups were  $CD4^+CD25^+CD127^{dim}$  or  $CD4^+CD25^-CD127^{dim}$  alone. After 18 h incubation, quantification of  $^3H$ -thymidine-incorporating cells showed that  $CD4^+CD25^+CD127^{dim}$  cells did not increase in

number, and the effector cells,  $CD4^+CD25^-CD127^{dim}$ , underwent significantly increased proliferation. More importantly, the proliferation of the effector cells was not inhibited by  $CD4^+CD25^+CD127^{dim}$  T cells (Fig. 5A;  $P < 0.001$ ), which demonstrates that acti-

A



B



**Figure 5. Showing  $CD4^+CD25^+Foxp3^+$  T cell suppression assay and expression of IFN- $\gamma$ .** (A) To evaluate the suppressive function, if any, of  $CD4^+CD25^+Foxp3^+$  T cells, 50,000  $CD4^+CD25^+CD127^{dim}$  T cells were cocultured with 50,000  $CD4^+CD25^-CD127^{dim}$  effector T cells (denoted as  $CD25^+ + CD25^-$  in the middle bar). Controls are  $CD4^+CD25^+CD127^{dim}$  or  $CD4^+CD25^-CD127^{dim}$ . Proliferation of  $^3H$ -thymidine-incorporated cells was quantified. Data are representative of two individual experiments. (B) IFN- $\gamma$ -ELISA showing that activated  $CD4^+CD25^+Foxp3^+$  T cells released significantly higher levels of IFN- $\gamma$  ( $P < 0.001$ ). Data are representative of two individual experiments.

vated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells had no suppressive function.

To further characterize the profile of this T cell subpopulation, supernatants were collected from unactivated and activated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells to evaluate the secretion of IFN- $\gamma$ , a marker for cytotoxic T<sub>H</sub>1 cells. Interestingly, IFN- $\gamma$  ELISA showed that activated CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells released high amounts (6000 pg/ml) of IFN- $\gamma$  into the culture media (Fig. 5B;  $P < 0.001$ ). In line with these findings, Zhou et al. [40] have shown that Foxp3<sup>+</sup> T<sub>regs</sub> lose their Foxp3 expression and take on an effector-memory T<sub>H</sub>1 phenotype producing high IFN- $\gamma$  and in some instances, have the potential to cause rapid pancreatic cell destruction and immune-mediated diabetes [40, 41]. In our study, we found that upon activation, human CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells did not lose the expression of Foxp3<sup>+</sup> and produced high levels of IFN- $\gamma$  and low IL-17. To explain these findings, two hypotheses are possible. First, there is a shift to a T<sub>H</sub>1 effector phenotype without the loss of Foxp3 or with transient Foxp3 expression. This hypothesis is in contrast with what has been reported previously in the mouse [40], which may highlight another difference between the mouse and human immune system. In addition, as suggested by Osorio et al. [42], Foxp3-expressing T<sub>regs</sub> may play an unpredicted proinflammatory role on some occasions. They are involved in the rejection of certain organ transplants [1], which may depend on the microenvironment in which the cells were activated [16] or be affected by the local inflammatory cytokines in the target tissue [43].

Second, Miyara et al. [20] have reported within the population of CD4<sup>+</sup>CD25<sup>+</sup>CD127<sup>dim</sup> T cells a fraction of Foxp3<sup>+</sup> non-T<sub>regs</sub> producing IL-17, IFN- $\gamma$ , and IL-2. Thus, there is the possibility that after activation, we have expanded this non-regulatory effector population.

In conclusion, this study demonstrates that upon activation, human CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> T cells become neurotoxic and present a shift toward a T<sub>H</sub>1 phenotype. Our study provides important insight about the potential limitations of human T<sub>regs</sub> for therapeutic applications and the requirement of serious caution in the use of these cells for disease-intervention strategies until we reach a better definition of the different subpopulations within the human CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> phenotype.

## AUTHORSHIP

Y.H. participated in the conception and design of the project, conducted most of the experiments, and wrote the manuscript. D.P. prepared and cultured human cortical neurons. D.T. conducted the suppression assay. R.C.B. participated in the conception and design of the project and revised the manuscript. F.G. supervised the entire study, participated in the conception and design of the project, and wrote the manuscript.

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## KEY WORDS:

granzyme B · axonal injury · cytotoxicity · T<sub>regs</sub> · neurodegeneration