

# Identifying Foxp3-expressing suppressor T cells with a bicistronic reporter

Yisong Y. Wan\* and Richard A. Flavell\*<sup>††</sup>

\*Section of Immunobiology and <sup>†</sup>Howard Hughes Medical Institute, Yale University School of Medicine, New Haven, CT 06520

Contributed by Richard A. Flavell, March 2, 2005

**Regulatory T cells are critical for maintaining self-tolerance and to negatively regulate immune responses. Foxp3 is a regulatory T cell-specific transcription factor that functions as the master regulator of the development and function of regulatory T cells. Here, we report the generation of a mouse model, in which a bicistronic reporter expressing a red fluorescent protein has been knocked into the endogenous Foxp3 locus. Using this mouse model, we assessed Foxp3 expression in various lymphocyte compartments and identified previously unreported Foxp3-expressing cells. In addition, we showed that *de novo* Foxp3 expression along with suppressive function were induced by TGF- $\beta$  in activated CD4 T cells *in vitro*. Finally, we demonstrated that non-Foxp3-expressing CD4 T cells could not be converted into Foxp3-expressing cells upon adoptive transfer into immunodeficient hosts. This Foxp3 bicistronic reporter knockin mouse model should greatly enhance the study of regulation and function of Foxp3-expressing regulatory T cells.**

regulatory T cells | TGF- $\beta$

How the body maintains immunological self-tolerance and negative control of physiological and pathological immune responses has been a long-standing question in immunology. Thirty years ago, it was proposed that subsets of thymus-derived CD4 T cells are essential to actively suppress immune responses and maintain self-tolerance (1–3). Recent studies have identified specific regulatory T cells that likely explain these phenomena (4–7). Subsets of regulatory T cells have been described, and they share common features of being hypoproliferative to T cell antigen receptor (TCR) stimulation *in vitro* and immunosuppressive *in vitro* and *in vivo*. Among regulatory T cells, naturally occurring regulatory T (Treg) cells and antigen-induced IL-10 producing regulatory T cells (Tr1) are the best characterized. Naturally occurring CD4<sup>+</sup>CD25<sup>+</sup> Treg cells comprise 5–10% of peripheral CD4<sup>+</sup> T cells and exert suppressive function through cell–cell contact and cytokine secretion, whereas induced CD4<sup>+</sup>CD25<sup>−</sup> Tr1 cells mediate immunosuppression through IL-10 and TGF- $\beta$  (8–10).

Many attempts in recent times have been made to identify a reliable surface marker for naturally occurring Treg cells. CD5<sup>high</sup>, CD45RB<sup>low</sup>, GITR<sup>high</sup>, and CD25<sup>+</sup> have all been used to enrich for the Treg population (5, 11–14). CD25 has been the most prominent marker used for identifying Treg cells. However, the expression of CD25 on Treg cells is unstable. CD25 is down-regulated when Treg cells are transferred into severe combined immunodeficient mice, whereas the suppressive function of Treg cells is maintained (15). In addition, non-Tr1 CD4<sup>+</sup>CD25<sup>−</sup> peripheral T cells have been found to possess regulatory activity (16, 17). Furthermore, CD25 is ubiquitously expressed by activated T cells, which makes it a marginally useful marker in identifying Treg cells from activated T cells *in vitro* or *in vivo*. Thus, a more reliable and unambiguous marker for Treg cells is needed.

Foxp3 is a transcription factor belonging to the forkhead family. In humans, mutation in *Foxp3* results in immunodysregulation, polyendocrinopathy, enteropathy, X linked syndrome, an X linked immunodeficiency syndrome associated with autoimmune disease in multiple endocrine organs (18–20). A frameshift mutation of Scurfin, the mouse orthologue of Foxp3, results in early lethality

due to hyperactivation of T cells in *Scurfy* mice (21). Recent studies have unveiled a specific role of Foxp3 in the development and function of Treg cells. Foxp3 is predominantly expressed in CD4<sup>+</sup>CD25<sup>+</sup> thymocytes and CD4<sup>+</sup>CD25<sup>+</sup> peripheral T cells (22, 23). Retroviral-mediated Foxp3 expression in CD4<sup>+</sup>CD25<sup>−</sup> T cells converts them into Treg-like cells phenotypically and functionally (22, 23). Analysis of Foxp3-transgenic or -knockout mice has further established an essential role for Foxp3 in regulatory T cell development. In mice transgenic for  $\approx$ 16 copies of cosmid containing endogenous *Foxp3* gene, the number of CD4<sup>+</sup>CD25<sup>+</sup> Treg cells is increased. Furthermore, in these mice, both CD4<sup>+</sup>CD25 and CD8<sup>+</sup> T cells that express Foxp3 exhibit immunosuppressive activity (21, 24, 25). Like *Scurfy* mice, Foxp3-deficient mice show hyperreactivity of T cells. This finding is due to a deficiency in the development of CD4<sup>+</sup>CD25<sup>+</sup> Treg cells (23). Thus, Foxp3 appears to be a reliable marker for Treg cells as it has been shown to be the master regulator specific for the development and function of these cells. However, as an intracellular protein, Foxp3 cannot be easily detected, which has hampered the study of the biology of Foxp3-expressing cells. Quantitation of Foxp3 mRNA has been used to estimate Treg frequency. There is, however, no guarantee that Foxp3 mRNA levels faithfully reflect Foxp3 protein, although a recent study suggests this method may be reliable (23).

Using a gene targeting approach, we have generated a mouse in which we knocked-in a bicistronic fluorescent reporter into the endogenous *Foxp3* locus, allowing us to identify Foxp3 expressing cells from different lymphocyte lineages and lymphoid organs. Our results indicate that, as expected, Foxp3 is predominantly expressed in CD4<sup>+</sup>CD25<sup>+</sup> peripheral T cells. In addition, we detected Foxp3 expression in a subset of CD4<sup>+</sup>CD25<sup>−</sup> peripheral T cells, a subset of CD4<sup>+</sup> thymocytes and CD4<sup>+</sup>TCR<sup>+</sup> bone marrow cells. Moreover, we show that *de novo* Foxp3 expression in activated CD4 T cells was potently induced by TGF- $\beta$  and conferred suppressive function on these cells *in vitro*. Finally, our results demonstrated that Foxp3<sup>−</sup> CD4 T cells could not be converted into Foxp3<sup>+</sup> cells in immunodeficient hosts.

## Materials and Methods

**Mice.** BALB/c mice (blastocyst donors), CD1 mice (foster mothers), C57BL/6 (B6) mice, Tet-Cre transgenic mice (“deletor” mice, C57BL/6 background), Rag-deficient mice (C57BL/6 background) and Foxp3-IRES-mRFP (FIR) mice (C57BL/6 background) were kept under specific pathogen-free conditions in the animal care facility at the Yale University. All mouse experiments were approved by Institutional Animal Care and Use Committee of Yale University.

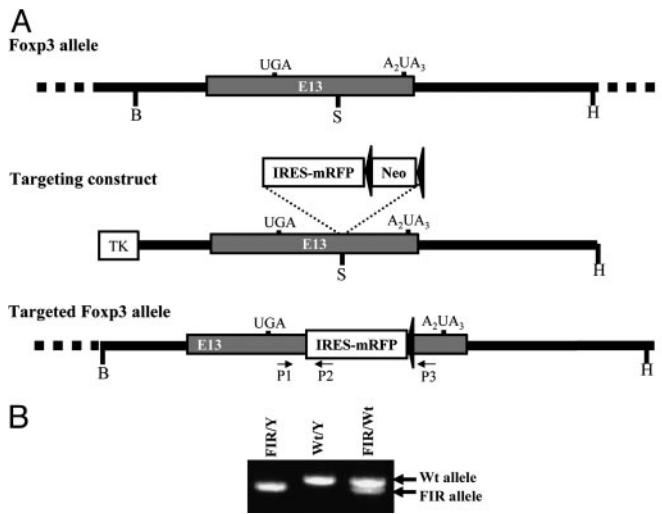
**Generation of FIR Mice.** A BAC clone (RP23–446O15) consisting of *Foxp3* genomic DNA derived from C57BL/6 mice was purchased from BacPac (Oakland, CA). An 11-kb BstZ17I/*Hpa*I fragment comprising exon 13 for *Foxp3* gene was cloned into pEasy-Flox

Abbreviations: APC, antigen-presenting cell; CFSE, 5,6-carboxyfluorescein diacetate succinimidyl ester; IRES-mRFP, internal ribosomal entry site-linked monomeric red fluorescent protein; TCR, T cell antigen receptor; Treg, regulatory T.

<sup>††</sup>To whom correspondence should be addressed. E-mail: richard.flavell@yale.edu.

© 2005 by The National Academy of Sciences of the USA





**Fig. 1.** Targeting IRES-mRFP reporter into the mouse *Foxp3* locus. (A) Maps for mouse *Foxp3* locus, targeting DNA construct, and the targeted *Foxp3* locus. An 11-kb mouse genomic DNA, including exon 13 of *Foxp3* gene, was excised by using *Bst*Z171 (B) and *Hpa*I (H) (Top) and cloned into pEasy-Flox vector adjacent to the thymidine kinase (TK) selection marker. A cassette containing IRES-mRFP and *LoxP*-flanked neomycin (Neo) selection marker was inserted into an *Ssp*I (S) site between the translation stop codon (UGA) and the polyadenylation signal ( $A_2UA_3$ ) of *Foxp3* gene (Middle). A correctly targeted ES cell was used to create chimeras and germ-line-transmitted mice. The Neo gene was removed *in vivo* by using deleter mice transgenic for Cre recombinase to generate mice bearing targeted *Foxp3* locus (Lower). (B) PCR genotyping FIR mice. Three primers (P1 to P3 as indicated) were designed to genotype FIR mice. PCR yielded 517-bp product for the wild-type (Wt) *Foxp3* allele and 470-bp product for targeted *Foxp3* allele.

levels of *Foxp3* mRNA were detected in  $CD25^{-}mRFP^{+}$   $CD4$  T cells. Interestingly, the geometric mean fluorescence intensity of mRFP as measured by flow cytometry was also lower in the  $CD25^{-}mRFP^{+}$  ( $\approx 37$ ) as compared with the  $CD25^{+}mRFP^{+}$  ( $\approx 59$ ) population. Thus, in FIR mice, mRFP expression faithfully marks *Foxp3* expression and can be used as a reliable marker for detecting *Foxp3*-expressing cells.

To test whether mRFP expression perturbed the immunological function of Treg cells, T cell proliferation and suppression assays were performed by using FACS-purified  $CD4^{+}mRFP^{+}$  and  $CD4^{+}CD25^{-}mRFP^{-}$  T cells as suppressor cells and responder cells, respectively (Fig. 2B). Suppressor and responder T cells were activated with soluble anti-CD3 and anti-CD28 antibodies in the presence of irradiated APCs either alone or as a combination of the two populations mixed at different ratios. T cell proliferation was measured by the incorporation of [ $^3$ H]thymidine. In contrast to  $CD4^{+}CD25^{-}mRFP^{-}$  T cells, which exhibited extensive proliferation (Fig. 2B, label R),  $CD4^{+}mRFP^{+}$  T cells did not proliferate (Fig. 2B, label S). Furthermore,  $CD4^{+}mRFP^{+}$  T cells inhibited the proliferation of  $CD4^{+}CD25^{-}mRFP^{-}$  T cells in a dose-dependent manner. Therefore, the immunosuppressive function of *Foxp3*-expressing cells is not impaired when mRFP is coexpressed. Thus, in FIR mice, mRFP marks *Foxp3*-expressing cells with high fidelity and without compromising their immunosuppressive functions.

By measuring mRFP, we assessed the expression of *Foxp3* in different lymphocyte lineages from different lymphoid organs (Fig. 2C). In peripheral lymph nodes, very few B cells ( $B220^{+}$ ) or  $CD8$  T cells ( $CD8^{+}TCR^{+}$ ) expressed *Foxp3* ( $<1\%$ ), whereas *Foxp3* expression was detected in  $\approx 10\%$  of  $CD4^{+}$  T cells ( $CD4^{+}TCR^{+}$ ) (Fig. 2C Top Left, Top Left Center, and Top Right Center). Immunosuppressive activity in the bone marrow has been documented in ref. 28. We therefore investigated whether *Foxp3*-expressing cells could be found in the bone marrow (Fig. 2C Top Right). Indeed,

$\approx 12\%$  of  $CD4^{+}$  cells residing in the bone marrow expressed *Foxp3*. Based on TCR expression,  $\approx 35\%$  of  $CD4^{+}$  bone marrow cells were also TCR positive ( $CD4^{+}TCR^{+}$ ), and *Foxp3* expression was detected only in the  $CD4^{+}TCR^{+}$  population in the bone marrow. Thus, interestingly,  $>30\%$  of the  $CD4^{+}TCR^{+}$  bone marrow cells expressed *Foxp3*, which was a much higher percentage than that of peripheral  $CD4^{+}TCR^{+}$  cells (Fig. 2C Top Left). The origin and the property of *Foxp3* expressing  $CD4^{+}TCR^{+}$  cells in the bone marrow remain to be elucidated. A small percentage of  $CD4^{+}$  thymocytes have been shown to express *Foxp3* and possess immunosuppressive activities (16, 22). This result prompted us to investigate *Foxp3* expression during T cell thymic development. In the thymus, *Foxp3* was expressed in  $\approx 1\%$  of  $CD4$  single positive and  $\approx 0.37\%$  of  $CD4CD8$  double-positive thymocytes. This result is in accordance with recent work done by Tai *et al.* (29) showing that *Foxp3* expression is induced in double-positive (DP) thymocytes upon TCR engagement *in vitro*. Neither  $CD8$  SP nor  $CD4CD8$  double-negative thymocytes expressed *Foxp3* (Fig. 2C Middle).  $CD4^{+}$  thymocytes were divided into three groups based on the expression levels of TCR,  $TCR^{High}$ ,  $TCR^{Med}$ , and  $TCR^{Low}$ . It appeared that *Foxp3* expression levels correlated with TCR expression levels (Fig. 2C Lower). This finding agrees with the reports showing that, in the thymus, TCR signaling is essential for Treg cell development, and the generation of Treg cells directly correlates with TCR signaling strength (30, 31). We also assessed  $CD25$  expression on *Foxp3*-expressing cells in the thymus and bone marrow. Like  $CD4^{+}$  T cells in the peripheral lymph nodes, a significant fraction of *Foxp3*-expressing  $CD4^{+}$  cells in the thymus and the bone marrow were negative for  $CD25$  (Fig. 2D).

**TGF- $\beta$  Induces de Novo *Foxp3* Expression and Regulatory Activity in  $CD4$  T Cells After Antigenic Stimulation *in Vitro*.**

It has been suggested that TGF- $\beta$  induces *Foxp3* expression in  $CD4$  T cells under *in vitro* and *in vivo* conditions (32–34). However, the question still remains as to whether TGF- $\beta$  does so by inducing *de novo* *Foxp3* expression in these cells. To address this question,  $CD4^{+}CD25^{-}$  T cells that did not express *Foxp3* were purified by FACS. Cells were activated by soluble anti-CD3 and anti-CD28 with irradiated APCs in the absence or presence of increasing amounts of TGF- $\beta$ . Three days later, *Foxp3* expression was analyzed by flow cytometry. *Foxp3* was highly induced in a substantial portion (up to 30%) of  $CD4$  T cells treated with TGF- $\beta$  in a dose-dependent manner after TCR engagement (Fig. 3A). Approximately 2–3% of cells expressed *Foxp3* in the absence of exogenous TGF- $\beta$ . We speculated this finding was due to the effects of endogenous TGF- $\beta$  produced by cultured cells. Thus, TGF- $\beta$  potentially induces *de novo* *Foxp3* expression in activated  $CD4$  T cells. Furthermore, this result suggests that the expression of *Foxp3* is not limited to naturally occurring regulatory T cells; rather, it can also be induced in a high percent of  $CD4$  T cells. To further investigate whether this induction can occur in dividing  $CD4$  T cells,  $CD4^{+}Foxp3^{-}$  T cells were purified by FACS and labeled with CFSE. CFSE is a fluorescent intracellular protein dye, whose fluorescent intensity is reduced to half after every cell division, allowing us to monitor cell division through flow cytometry. In the presence of exogenous TGF- $\beta$ , CFSE-labeled cells were either left inactivated or activated with soluble anti-CD3 and anti-CD28 in the presence of irradiated APCs. After 3 days of stimulation,  $CD4^{+}Foxp3^{+}$  cells were gated and the cell divisions were measured by flow cytometry (Fig. 3B). Our results demonstrate that *Foxp3* expression can be induced in dividing  $CD4$  T cells in the presence of TGF- $\beta$ .

*Foxp3* expressing  $CD25^{+}CD4^{+}$  naturally occurring Treg cells are hyporesponsive toward TCR stimulation and suppress T cell effector functions (22, 23). To address whether TGF- $\beta$ -induced *Foxp3*-expressing T cells generated *in vitro* share these characteristics of naturally occurring Treg cells, we first purified *Foxp3* $^{+}$  and *Foxp3* $^{-}$   $CD4$  T cells after 3 days of activation in the presence of TGF- $\beta$ . Purified T cells were restimulated by soluble anti-CD3 and anti-





CD4<sup>+</sup>CD25<sup>-</sup>Foxp3<sup>-</sup> T cell transferred hosts developed IBD as manifested by morbidity, hunched back, diarrhea, and an enlarged spleen, intestine, and colon (data now shown), whereas bone marrow transplanted hosts showed no sign of any disease even 6 weeks after transfer. Thus, in the immunodeficient hosts, mature non-Foxp3 expressing CD4 T cells cannot be converted into Foxp3-expressing CD4 T cells, and the failure to generate Foxp3<sup>+</sup> Treg cells may contribute to the development of IBD in this model. It would be interesting to know whether CD4<sup>+</sup>Foxp3<sup>-</sup> T cells can become CD4<sup>+</sup>Foxp3<sup>+</sup> T cells in the sublethally irradiated wild-type hosts.

### The FIR Mouse Is a Promising Model to Study the Biology of Foxp3 Expressing Treg Cells.

It has been widely accepted that Treg cells are critical for maintaining self-tolerance and actively suppressing immune responses. Attempts have been made to manipulate Treg populations to treat various human pathologies, enhance antitumor immunity, and maintain allograft tolerance after organ transplantation (38–40). Further understanding the biology of Treg cells clearly has important clinical implications. However, the lack of reliable markers has been a major hindrance to the study of Treg cells. With few exceptions, e.g., Tr1 cells (41–43), the expression of Foxp3 goes hand in hand with suppressive T cell function. Ample studies had been done to establish Foxp3 as an essential gene that plays specific roles in regulating the development and function of Treg cells. Thus, the expression of Foxp3 appears to specifically

mark Treg cells. By a knockin approach, we have inserted a bicistronic red fluorescent reporter into the endogenous *Foxp3* locus to generate FIR mice without compromising the expression of Foxp3 and the function of Foxp3-expressing Treg cells. Using this FIR mouse, we were able to identify previously reported and some unreported Foxp3-expressing cells in mouse. Using FIR cells, we isolated TGF- $\beta$ -induced Foxp3-expressing functional suppressor T cells after TCR stimulation *in vitro*, which could have not been done by relying on other traditional markers for Treg cells. Furthermore, we demonstrated that this model is of value to study the generation of Foxp3-expressing Treg cells *in vivo*. The fact that these mice were generated on the inbred B6 background will substantially facilitate their further use particularly for adoptive transfer studies to other mice. In summary, FIR mice greatly enhanced the ability to study the biology of Foxp3-expressing Treg cells and will be a valuable tool to test both scientific hypotheses and clinical applications of Treg cells in mice.

We thank Klaus Rajewsky (Harvard Medical School, Boston) for the gift of Bruce4 ES cells (44), Roger Tsien for providing mRFP cDNA, Linda Evangelisti and Cindy Hughes for generating ES cells and chimeric mice, respectively, Judy Stein for the Southern screening, Elizabeth Eynon for managing the mouse program, and Fran Manzo for help with manuscript preparation. Y.Y.W. was supported by a grant from the Cancer Research Institute. This work was supported by a grant from the National Institutes of Health (RO1 DK51665) and the American Diabetes Association. R.A.F. is an investigator of the Howard Hughes Medical Institute.

- Nishizuka, Y. & Sakakura, T. (1969) *Science* **166**, 753–755.
- Penhale, W. J., Farmer, A., McKenna, R. P. & Irvine, W. J. (1973) *Clin. Exp. Immunol.* **15**, 225–236.
- Penhale, W. J., Irvine, W. J., Inglis, J. R. & Farmer, A. (1976) *Clin. Exp. Immunol.* **25**, 6–16.
- Powrie, F. & Mason, D. (1990) *J. Exp. Med.* **172**, 1701–1708.
- Sakaguchi, S., Fukuma, K., Kuribayashi, K. & Masuda, T. (1985) *J. Exp. Med.* **161**, 72–87.
- Sugihara, S., Izumi, Y., Yoshioka, T., Yagi, H., Tsujimura, T., Tarutani, O., Kohno, Y., Murakami, S., Hamaoka, T. & Fujiwara, H. (1988) *J. Immunol.* **141**, 105–113.
- McKeever, U., Mordes, J. P., Greiner, D. L., Appel, M. C., Rozing, J., Handler, E. S. & Rossini, A. A. (1990) *Proc. Natl. Acad. Sci. USA* **87**, 7618–7622.
- Maloy, K. J. & Powrie, F. (2001) *Nat. Immunol.* **2**, 816–822.
- Sakaguchi, S. (2004) *Annu. Rev. Immunol.* **22**, 531–562.
- Roncarolo, M. G., Bacchetta, R., Bordignon, C., Narula, S. & Levings, M. K. (2001) *Immunol. Rev.* **182**, 68–79.
- Powrie, F., Leach, M. W., Mauze, S., Caddle, L. B. & Coffman, R. L. (1993) *Int. Immunol.* **5**, 1461–1471.
- Morrissey, P. J., Charrier, K., Braddy, S., Liggett, D. & Watson, J. D. (1993) *J. Exp. Med.* **178**, 237–244.
- Shimizu, J., Yamazaki, S., Takahashi, T., Ishida, Y. & Sakaguchi, S. (2002) *Nat. Immunol.* **3**, 135–142.
- Sakaguchi, S., Sakaguchi, N., Asano, M., Itoh, M. & Toda, M. (1995) *J. Immunol.* **155**, 1151–1164.
- Gavin, M. A., Clarke, S. R., Negrou, E., Gallegos, A. & Rudensky, A. (2002) *Nat. Immunol.* **3**, 33–41.
- Stephens, L. A. & Mason, D. (2000) *J. Immunol.* **165**, 3105–3110.
- Apostolou, I., Sarukhan, A., Klein, L. & von Boehmer, H. (2002) *Nat. Immunol.* **3**, 756–763.
- Chatila, T. A., Blaeser, F., Ho, N., Lederman, H. M., Voulgaropoulos, C., Helms, C. & Bowcock, A. M. (2000) *J. Clin. Invest.* **106**, R75–R81.
- Wildin, R. S., Ramsdell, F., Peake, J., Faravelli, F., Casanova, J. L., Buist, N., Levy-Lahad, E., Mazzella, M., Goulet, O., Perroni, L., et al. (2001) *Nat. Genet.* **27**, 18–20.
- Bennett, C. L., Christie, J., Ramsdell, F., Brunkow, M. E., Ferguson, P. J., Whitesell, L., Kelly, T. E., Saulsbury, F. T., Chance, P. F. & Ochs, H. D. (2001) *Nat. Genet.* **27**, 20–21.
- Brunkow, M. E., Jeffery, E. W., Hjerrild, K. A., Paepfer, B., Clark, L. B., Yasayko, S. A., Wilkinson, J. E., Galas, D., Ziegler, S. F. & Ramsdell, F. (2001) *Nat. Genet.* **27**, 68–73.
- Hori, S., Nomura, T. & Sakaguchi, S. (2003) *Science* **299**, 1057–1061.
- Fontenot, J. D., Gavin, M. A. & Rudensky, A. Y. (2003) *Nat. Immunol.* **4**, 330–336.
- Khattari, R., Cox, T., Yasayko, S. A. & Ramsdell, F. (2003) *Nat. Immunol.* **4**, 337–342.
- Khattari, R., Kasprovicz, D., Cox, T., Mortrud, M., Appleby, M. W., Brunkow, M. E., Ziegler, S. F. & Ramsdell, F. (2001) *J. Immunol.* **167**, 6312–6320.
- Campbell, R. E., Tour, O., Palmer, A. E., Steinbach, P. A., Baird, G. S., Zacharias, D. A. & Tsien, R. Y. (2002) *Proc. Natl. Acad. Sci. USA* **99**, 7877–7882.
- Jang, S. K. & Wimmer, E. (1990) *Genes Dev.* **4**, 1560–1572.
- Muraoka, S. & Miller, R. G. (1980) *J. Exp. Med.* **152**, 54–71.
- Tai, X., Cowan, M., Feigenbaum, L. & Singer, A. (2005) *Nat. Immunol.* **6**, 152–162.
- Jordan, M. S., Boesteanu, A., Reed, A. J., Petrone, A. L., Hohenbeck, A. E., Lerman, M. A., Naji, A. & Caton, A. J. (2001) *Nat. Immunol.* **2**, 301–306.
- Suto, A., Nakajima, H., Ikeda, K., Kubo, S., Nakayama, T., Taniguchi, M., Saito, Y. & Iwamoto, I. (2002) *Blood* **99**, 555–560.
- Chen, W., Jin, W., Hardegen, N., Lei, K. J., Li, L., Marinos, N., McGrady, G. & Wahl, S. M. (2003) *J. Exp. Med.* **198**, 1875–1886.
- Peng, Y., Laouar, Y., Li, M. O., Green, E. A. & Flavell, R. A. (2004) *Proc. Natl. Acad. Sci. USA* **101**, 4572–4577.
- Green, E. A., Gorelik, L., McGregor, C. M., Tran, E. H. & Flavell, R. A. (2003) *Proc. Natl. Acad. Sci. USA* **100**, 10878–10883.
- Liang, S., Alard, P., Zhao, Y., Parnell, S., Clark, S. L. & Kosiewicz, M. M. (2005) *J. Exp. Med.* **201**, 127–137.
- Curotto de Lafaille, M. A., Lino, A. C., Kutchukhidze, N. & Lafaille, J. J. (2004) *J. Immunol.* **173**, 7259–7268.
- Singh, B., Read, S., Asseman, C., Malmstrom, V., Mottet, C., Stephens, L. A., Stepankova, R., Tlaskalova, H. & Powrie, F. (2001) *Immunol. Rev.* **182**, 190–200.
- Baecher-Allan, C. & Hafler, D. A. (2004) *J. Exp. Med.* **200**, 273–276.
- Sutmoller, R. P., van Duivenvoorde, L. M., van Elsas, A., Schumacher, T. N., Wildenberg, M. E., Allison, J. P., Toes, R. E., Offringa, R. & Melief, C. J. (2001) *J. Exp. Med.* **194**, 823–832.
- Edinger, M., Hoffmann, P., Ermann, J., Drago, K., Fathman, C. G., Strober, S. & Negrin, R. S. (2003) *Nat. Med.* **9**, 1144–1150.
- Groux, H., O'Garra, A., Bigler, M., Rouleau, M., Antonenko, S., de Vries, J. E. & Roncarolo, M. G. (1997) *Nature* **389**, 737–742.
- Levings, M. K., Sangregorio, R., Sartirana, C., Moschin, A. L., Battaglia, M., Orban, P. C. & Roncarolo, M. G. (2002) *J. Exp. Med.* **196**, 1335–1346.
- Levings, M. K., Gregori, S., Tresoldi, E., Cazzaniga, S., Bonini, C. & Roncarolo, M. G. (2005) *Blood* **105**, 1162–1169.
- Kontgen, F., Suss, G., Stewart, C., Steinmetz, M. & Bluethmann, H. (1993) *Int. Immunol.* **5**, 957–964.