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Effects of T3R α 1 and T3R α 2 Gene Deletion on T and B Lymphocyte Development¹

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Thyroid hormones bind to several nuclear receptors encoded by T3R α and T3R β genes. There is now accumulating evidence that thyroid hormones act on the immune system. Indeed, mice deficient for thyroid hormones show a reduction in lymphocyte production. However, the mechanisms involved and, in particular, the role of the different thyroid hormone receptors in lymphocyte development have not been investigated. To address that question, we have studied lymphocyte development in mice deficient for the T3R α 1 and T3R α 2 gene products. A strong decrease in spleen cell numbers was found compared with wild-type littermates, B lymphocytes being more severely affected than T lymphocytes. A significant decrease in splenic macrophage and granulocyte numbers was also found. In bone marrow, a reduction in CD45⁺/IgM⁻ pro/pre-B cell numbers was found in these mice compared with wild-type littermates. This decrease seems to result from a proliferation defect, as CD45⁺/IgM⁻ cells incorporate less 5-bromo-2'-deoxyuridine *in vivo*. To define the origin of the bone marrow development defect, chimeric animals between T3R α ^{-/-} and Rag1^{-/-} mice were generated. Results indicate that for B cells the control of the population size by T3R α 1 and T3R α 2 is intrinsic. Altogether, these results show that T3R α 1 or T3R α 2 gene products are implicated in the control of the B cell pool size. *The Journal of Immunology*, 2000, 164: 152–160.

The role of thyroid hormones in the general development of organisms has been known for a long time. There are also increasing data supporting reciprocal interactions between thyroid hormones and the immune system (1). These interactions have been revealed by experimentally induced fluctuations of thyroid hormones in a number of species, including amphibians (2), avians (3), and rodents (4).

The influence of thyroid hormones on the immune system has been extensively studied in mice deficient for hormones of the pituitary/thyroid axis. Snell dwarf (*dw/dw*) mice are deficient in anterior pituitary hormones, i.e., growth hormone (GH),⁴ prolactin, and thyroid-stimulating hormone (TSH), due to a point mutation in the gene encoding the *Pit-1* transcription factor (5). Lack of GH and TSH production subsequently causes a drop in seric concentrations of insulin-like growth-factor-I (IGF-I) (6) and thyroid hormones, respectively. The numerous studies of the immune system in Snell dwarf mice have revealed a role for thyroid hormones in

B and T cell development (7–13). For example, an increase in thymocyte cellularity is induced by exogenously administered thyroxine (T₄) in Snell dwarf mice, and this effect has been confirmed in the thyroid hormone deficient *hyt/hyt* strain of mice (13) and in normal mice (4). For the B cell lineage, there is a drop in cell numbers at the pro/pre-B cell stage (10, 12, 13), which could result from a proliferation or differentiation defect.

Deficient thyroid hormone production seems the main factor responsible for impaired B cell development in Snell dwarf mice. Indeed, pre-B cell numbers reach normal levels after thyroxine injections, while they are not restored by prolactin, GH, or IGF-I treatment (12, 14). The latter findings are consistent with the observation that B cell development occurs normally in IGF-I, GH, and prolactin-deficient mice (13, 14). The predominant effect of thyroid hormones on B cell development was confirmed by the recovery of the pre-B cell compartment in the *hyt/hyt* thyroid hormone-deficient mice following T₄ injections (13).

In these studies, the target of thyroid hormones was not identified, i.e., it is still unclear whether they regulate the size of the B cell pool by acting directly on B cell progenitors or by optimizing the cellular environment necessary for their development. In addition, the thyroid hormone receptors that influence lymphoid development have not been identified. The thyroid gland produces T₄ that is metabolized by thyroxine 5'-deiodinases to its active form: triiodo-thyronine (T₃) (15, 16). T₃ binds to several nuclear receptors encoded by the T3R α and T3R β loci (17, 18). T3R α (19)-, T3R β (20)-, and T3R α 1 (21)-deficient mice have recently been generated by targeted gene disruption. The T3R α KO mice are deficient for the T3R α 1 and T3R α 2 isoforms (19). However, they are still able to express the T3R Δ 1 and T3R Δ 2 short transcripts (22), which function is unknown. These mice have a general growth defect and die between 4–6 wk of age. A number of defaults, including bone and small intestine development, have been investigated (19).

We have studied the development of lymphocytes in these T3R α KO mice. We have found that the size of the mature B cell

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⁴ Abbreviations used in this paper: GH, growth hormone; BrdU, 5-bromo-2'-deoxyuridine; IGF, insulin-like growth factor; KO, knockout; T₃, triiodo-thyronine; T₄, thyroxine; TSH, thyroid-stimulating hormone.

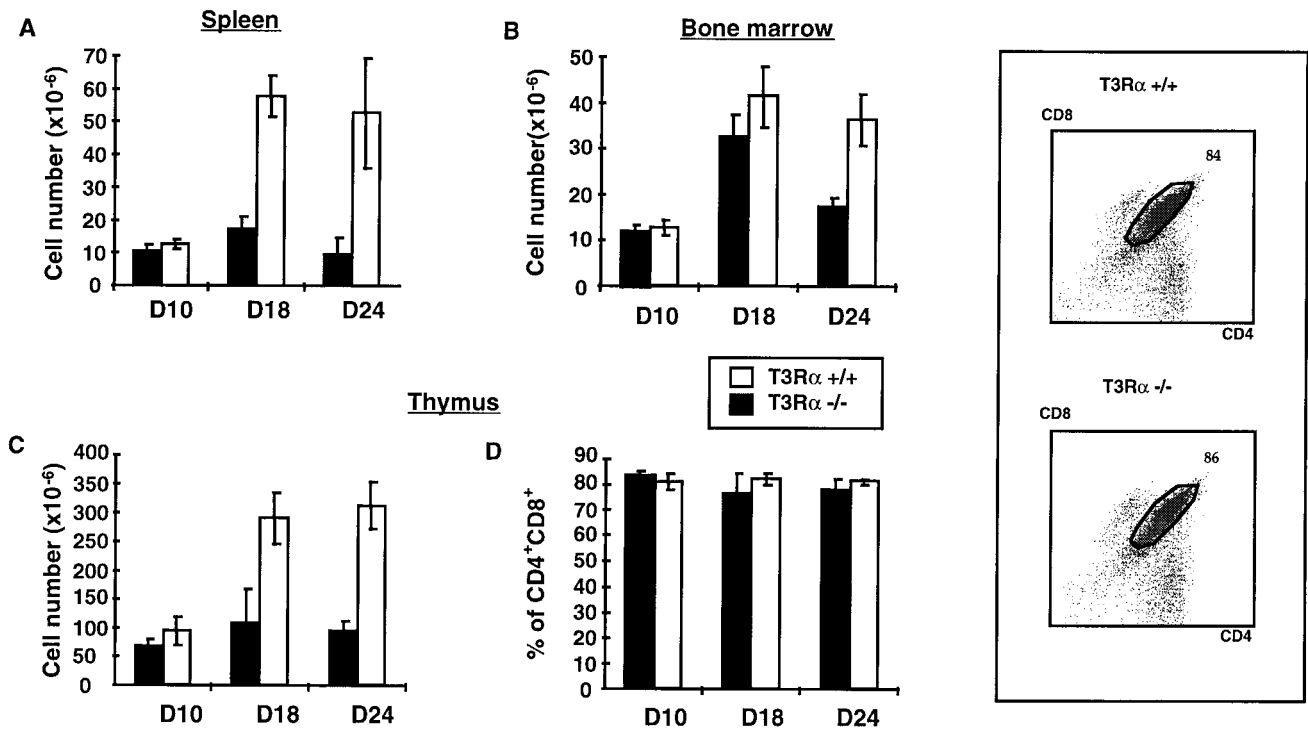


FIGURE 1. Lymphocyte numbers in T3R α KO mice. The numbers of splenocytes (A), bone marrow cells (B), and thymocytes (C), and the percentages of CD4⁺CD8⁺ double-positive thymocytes (D) in 10-, 18-, and 24-day-old T3R α KO (■) and wild-type (□) mice are shown. *Inset*, An example of double anti-CD4/anti-CD8 staining of thymocytes from 18-day-old T3R α ^{+/+} (top) and T3R α ^{-/-} (bottom) mice is shown. The figures represent the percentages of gated double-positive thymocytes.

pool is particularly affected. This default was at least partly due to a lack of B cell progenitor proliferation. Finally, we show that intrinsic expression of T3R α 1 or T3R α 2 gene products is needed for the development of a normal B cell pool.

Materials and Methods

Mice

Rag1^{-/-} mice were purchased from Transgenic Alliance (l'Arbresle, France). T3R α KO mice, developed on the outbred OF₁ background, were generated in our laboratory (19). All mice were bred in the institute's animal facility and kept under pathogen-free conditions. As homozygous T3R α KO mice die between 4 and 6 wk of age, heterozygous animals were routinely crossed to generate the T3R α ^{-/-} and T3R α ^{+/+} littermates used in the experiments. The genotype of the offspring was analyzed by PCR, using a mixture of oligonucleotides specific for T3R α (A and B) and a lacZ oligonucleotide (lacZ-A5) specific for the targeting vector. A PCR product is amplified with the oligonucleotides A and B when the wild-type allele is present, and with the oligonucleotides A and lacZ-A5 when the gene is disrupted (A, 5'-GGACAAGTCTCTGAAGACTTCC-3'; B, 5'-GTCTGACCCACACTCCACCTTG-3'; lacZ A5, 5'-CCTCTTCGCTATTACGCAGCTGG-3'). The amplification conditions used were 20 s at each of the following temperatures: 94°C, 58°C, and 72°C, for 32 cycles.

Fluorescence staining and flow cytometry sorting

For surface marker analysis, spleen cells were stained as previously described (23). The following Abs were used: 2B8-biotin (anti-CD117 (c-kit)), anti-CD4-PE, and anti-CD8 α -Tricolor from Caltag Laboratories (Burlingame, CA); R6-60.2-biotin (anti-IgM), M1/70-biotin (anti-CD11b/Mac-1), S7-biotin (anti-CD43), 7D4-FITC (anti-CD25), and RB6-8C5-biotin (anti-Ly-6G/Gr-1) from Becton Dickinson (Le Pont de Claix, France); RA3 6B2-PE (anti-B220) from Sigma Immunochemicals (St. Quentin Fallavier, France); 145.2C11-FITC (anti-CD3) and M1.93-biotin (anti-CD45) were prepared in house. Avidin-Tricolor (Caltag Laboratories) was used to reveal biotin-conjugated Abs.

BrdU labeling and staining were performed as previously reported (23). Briefly, mice were injected twice i.p. with 1 mg BrdU (Sigma-Aldrich Chimie, St. Quentin Fallavier, France) at a 30-min interval. Six hours later, animals were sacrificed and the lymphocytes were surface stained. Cells

were then fixed and permeabilized, and after partial DNA digestion, BrdU incorporation was revealed by a FITC-conjugated anti-BrdU Ab from Becton Dickinson (Le Pont de Claix, France; clone B44).

Cells were analyzed using the FACScan device (Becton Dickinson, Mountain View, CA) and the CellQuest software (Becton Dickinson Immunocytometry Systems, San Jose, CA). For cell cycle analysis, cells were surface stained and then fixed in 70% cold ethanol. Before analysis, cells were washed in PBS and incubated for 30 min to 2 h in 0.5 μ g/ml Hoechst 33342 (Molecular Probes, Leiden, The Netherlands). Cells were analyzed in the presence of the dye on a FACStar^{Plus} (Becton Dickinson) equipped with an UV argon laser.

Hormone assays

Cortisol levels in sera were assayed by a private laboratory (Laboratoire d'Analyses Médicales Collombel-Orfeuvre, France) using an enzyme-linked fluorescent assay (ELFA).

Generation of chimeric animals by morulae aggregation

Rag1^{-/-} and T3R α ^{+/+} animals were crossed independently. On 2.5 days postcoitum, mice were sacrificed, oviducts were dissected out, and morulae were isolated. The zona pellucida was removed from morulae by acid Tyrode's solution. Morulae from the Rag1^{-/-} cross were aggregated to the morulae from the T3R α ^{+/+}, as previously described (24). After overnight in vitro expansion, the resulting blastocysts were transferred to the uteruses of 2.5-day postcoitum pseudo-pregnant females. For each offspring, the T3R α genotype of the morulae derived from the T3R α ^{+/+} cross was determined on mature lymphocytes. Indeed, as mature lymphocytes cannot be derived from the Rag1^{-/-} morulae, this cellular subset is not chimeric. The degree of chimerism was estimated by measuring the ratio between PCR products corresponding to the wild-type and targeted T3R α gene locus, amplified from hepatocytes.

Results

T3R α KO mice show a defect in lymphoid organ cell numbers

To assess the effect of T3R α 1 and T3R α 2 deletion on lymphocyte development, we first determined cell numbers in primary (thymus and bone marrow) and secondary (spleen) lymphoid organs of T3R α KO mice, at different times following birth. As shown in

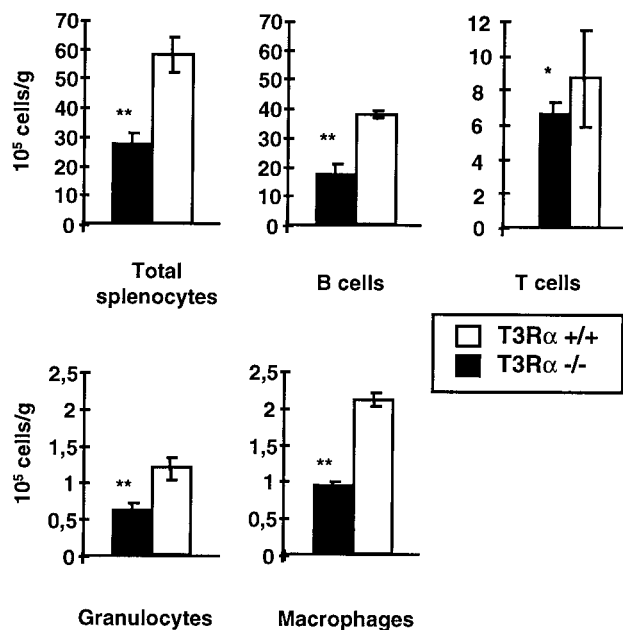


FIGURE 2. Splenocyte subpopulations in T3R α KO mice. Splenocytes from 17- to 19-day-old mice were counted and stained with anti-B220 and anti-CD3 Abs to identify B and T lymphocytes, respectively. Granulocytes and macrophages were identified by anti-Ly-6G and anti-CD11b Abs, respectively. The cellularity index was calculated as the number of cells per organ divided by the weight of T3R α ^{-/-} (■) or wild-type control (□) animals. The cellularity index values are expressed as 10⁵ cells/g. *, $p < 0.1$; **, $p < 0.01$ (Student's t test).

Fig. 1, at 10 days of age, cell numbers of T3R α KO mice are not significantly different when compared with wild-type littermates. In contrast, by day 18 there is a sharp difference in thymocyte and splenic lymphocyte numbers in T3R α KO compared with wild-type animals, while bone marrow cell numbers remain equivalent in both types of mice. By day 24, cell numbers in all three lymphoid organs are drastically reduced in KO mice. This defect is further increased at later time points (data not shown).

However, T3R α KO mice also suffer from stress-inducing defects, including decreased nutritional functions due to a delayed maturation of the small intestine (19). The stress response influences the immune system, notably through glucocorticoid production, which down-regulates immune processes and induces apoptosis of lymphoid cells (25). In particular, the percentage of double-positive thymocytes is a good indicator of stress, as these cells are the most sensitive to glucocorticoid-induced depletion (26). Hence, to assess stress levels, we have measured cortisol

levels in the blood of mice aged 18 or 32 days. At 18 days of age, there is no significant difference in the levels of cortisol measured in T3R α KO and wild-type mice (28 ± 8 nM vs 21 ± 1 nM). However, by day 32, T3R α KO mice show a sharp rise in their levels of circulating cortisol compared with wild-type mice (76 ± 25 nM vs 28 ± 12 nM). Moreover, we have found that the CD4⁺CD8⁺ double-positive subset of thymocytes in T3R α KO mice is not significantly affected up to day 24 of age (Fig. 1), confirming the absence of major stress-induced modifications of the immune system during early life in these mice.

Therefore, to study the impact of T3R α 1 and T3R α 2 deletion on the immune system without potential interference of the stress response, we have performed all further analyses on 17- to 19-day-old animals. Moreover, in all experiments, the percentage of double-positive thymocytes was measured, and the few mice (less than 3%) showing a preferential deletion of double-positive thymocytes, i.e., mice in which the double-positive subset represented less than 80% of total thymocytes, were not included in the analysis.

Finally, as T3R α KO mice are smaller than wild-type littermates (19), cellularity in lymphoid organs was normalized by calculating a cellularity index, which corresponds to the total cell number divided by the weight of the animal (27).

Splenic B lymphocytes are severely affected by T3R α 1 and T3R α 2 deletion

To further characterize splenic subpopulations affected in T3R α KO mice, splenocytes were stained with anti-CD45R (B220), CD3, CD4, and CD8 Abs. As shown in Fig. 2, both B and T lymphocyte numbers were decreased in spleens from T3R α KO animals. However, if T lymphocyte cellularity was reproducibly decreased by about 30% as compared with the control littermates (Table I), the small number of animals analyzed in each group and the wide variability among them led to a poorly significant difference between deficient and wild-type animals in each experiment ($0.05 < p < 0.1$; Student's t test, Fig. 2). It is noteworthy that the numbers of CD4⁺ and CD8⁺ T lymphocytes were reproducibly reduced to the same extent (data not shown).

In contrast, in all experiments, splenic B (i.e., B220⁺) lymphocyte cellularity was significantly ($p < 0.005$; Student's t test) and reproducibly (Table I) decreased by about 60% in T3R α -deficient mice compared with wild-type littermates (Fig. 2). Similar results were obtained when IgM⁺ splenic cells were measured, indicating that all B cell subsets were affected in the periphery (data not shown). Thus, T3R α 1 and T3R α 2 gene deletion affects splenic B cells more profoundly than splenic T lymphocytes.

As shown in Fig. 2, the numbers of splenic granulocytes and macrophages, identified by anti-Ly-6G (Gr-1⁺) and anti-CD11b

Table I. *In spleen T3R α 1 and T3R α 2 deletion affects more profoundly B than T lymphocytes^a*

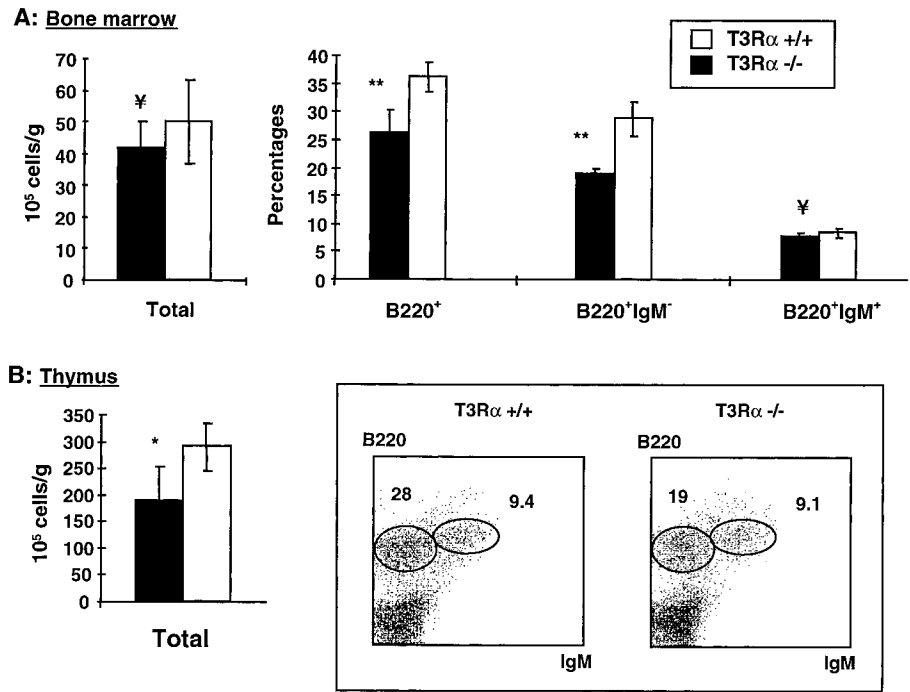
Expt.	<i>n</i> (KO; wt)	Total Splenocytes		B Lymphocytes		T Lymphocytes	
		Cell index (KO; wt) ^b	Ratio, KO/wt (%) ^c	Cell index (KO; wt) ^b	*Ratio, KO/wt (%) ^c	Cell index (KO; wt) ^b	Ratio, KO/wt (%) ^c
1	(4; 1)	(33; 84)	39	(18; 58)	31	(11; 17)	68
2	(3; 3)	(27; 58)	47	(17; 38)	45	(7; 9)	75
3	(2; 1)	(30; 60)	49	(18; 42)	43	(9; 13)	66
4	(2; 2)	(20; 43)	48	(12; 33)	37	(6; 7)	81

^a Seventeen- to 19-day-old T3R α KO (KO) and wild-type (wt) mice were sacrificed. Splenocytes were counted and stained with anti-B220 and anti-CD3 to detect B and T lymphocytes, respectively.

^b Cellularity index was calculated as the number of cells per organ divided by the weight of the animals. The values are expressed as 10⁵ cells/g.

^c For each experiment the percentage shown corresponds to: [(mean of cellular indices in KO mice/mean of cellular indices in wild-type mice) \times 100].

FIGURE 3. Lymphoid development in T3R α KO mice. Bone marrow nucleated cells (A) and thymocytes (B) obtained from 17- to 19-day-old mice were counted and stained. *Inset*, An example of double anti-B220/anti-IgM staining of bone marrow nucleated cells from T3R α ^{+/+} (left) and T3R α ^{-/-} (right) mice is shown. The figures in the upper left and right corners represent the percentages of B220⁺IgM⁻ and B220⁺IgM⁺ bone marrow cells, respectively. The cellularity index was calculated as the number of cells per organ divided by the weight of the animals. The cellularity index values are expressed as 10⁵ cells/g. Results for T3R α ^{-/-} (■) or wild-type control (□) mice are shown. A, Bone marrow lymphocytes were double stained for B220 and IgM expression and analyzed on FACS. As T3R α KO and wild-type animals show no significant differences in bone marrow cellularity indices, the size of the different bone marrow subsets is expressed as a percentage of total cell numbers. †, $p > 0.2$; *, $p < 0.1$; **, $p < 0.01$ (Student's *t* test).



(Mac-1⁺) Abs, respectively, were also significantly reduced by about 50% in T3R α -deficient mice compared with wild-type aged-matched control animals.

T3R α 1 and T3R α 2 deletion affects lymphocytes in primary lymphoid organs

As T3R α 1 and T3R α 2 gene deletion affects both B and T lymphocyte numbers in the periphery, we questioned whether they developed normally in primary lymphoid organs. As shown in Fig. 3A, the total number of cells in bone marrow, when corrected for the weights of mice, was not significantly different in wild-type and T3R α KO mice. However, the numbers and percentages of B lineage (B220⁺) bone marrow cells were significantly (Fig. 3A) and reproducibly (Table II) reduced by about 20% in T3R α -deficient animals. This decrease in the number of B lymphocyte precursors preferentially affects the pro/pre-B cell (B220⁺IgM⁻) stage (Fig. 3A, Table II). Immature/mature (B220⁺IgM⁺) B cell numbers were not reproducibly reduced in bone marrow of T3R α KO mice (Table II). To further pinpoint the B cell progenitor defect, we have used anti-CD43, CD25, and CD117 (*c-kit*) Abs to

look at the different progenitor B cell subsets, as defined by A. Rolink and colleagues (28). Results in Table III show that only the B220⁺CD25⁺ pre-BII cells were significantly decreased in T3R α KO animals. When cells were analyzed according to their size, we found that only the small B220⁺CD25⁺ pre-BII cells were affected (Table III).

As observed for splenic T cells, T3R α 1 and T3R α 2 gene deletion reproducibly (Table II) induces a slightly significant (0.025 < $p < 0.05$; Student's *t* test, Fig. 3B) decrease of about 30% in thymocyte numbers. Interestingly, thymic subpopulations were equally affected as the percentages of CD4⁻CD8⁻, CD4⁺CD8⁺, CD4⁺CD8⁻, and CD4⁻CD8⁺ thymocytes were not modified in 17- to 19-day-old T3R α KO animals (data not shown).

Lymphocyte precursors from T3R α KO mice proliferate less in primary lymphoid organs

To determine whether the reduction in lymphocyte numbers in both primary and secondary lymphoid organs of T3R α KO mice was due to a decreased proliferative capacity of these cells, we have measured their proliferation in vivo. For that purpose, T3R α

Table II. The size of the B220⁺IgM⁻ pre/pro B cells subset is decreased in T3R α KO mice^a

Expt.	Thymus				Bone Marrow						
	n (KO; wt)	Cell Index (KO; wt) ^a	Ratio, KO/wt (%) ^b	Cell index (KO; wt) ^a	% B220 ⁺		% B220 ⁺ IgM ⁻		% B220 ⁺ IgM ⁺		
					Ratio, KO/wt (%) ^b	Ratio, KO/wt (%) ^d	Ratio, KO/wt (%) ^d	Ratio, KO/wt (%) ^d	Ratio, KO/wt (%) ^d	Ratio, KO/wt (%) ^d	
1	(3; 3)	(19; 29)	66	(43; 41)	104	(24; 33)	74	(17; 23)	72	(8; 10)	79
2	(3; 3)	(15; 24)	64	(42; 36)	116	(26; 38)	70	(19; 31)	62	(8; 9)	79
3	(2; 2)	(20; 38)	53	(46; 34)	136	(45; 54)	84	(28; 33)	83	(18; 19)	97
4	(3; 3)	(24; 34)	69	(42; 41)	102	(34; 47)	72	(19; 28)	68	(13; 19)	69
5	(3; 3)	ND	ND	ND	ND	(37; 42)	89	(23; 30)	77	(14; 12)	121
6	(3; 3)	ND	ND	ND	ND	(39; 45)	86	(19; 27)	71	(20; 19)	104

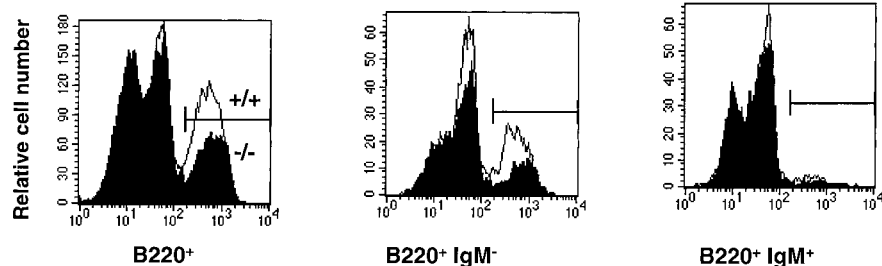
^a Mononucleated cells were counted and cellularity index was calculated as the number of cells per organ divided by the weight of the animals. Values are expressed as 10⁵ cells/g.

^b The percentage shown corresponds to: [(mean of cellular indices in KO mice/mean of cellular indices in wild-type (wt) mice)] × 100.

^c Bone marrow cells were stained with anti-B220 and anti-IgM Abs, and the percentages of each subset was determined using flow cytometry.

^d The percentage shown corresponds to: [(mean values for KO mice/mean values for wt mice)] × 100.

A: Bone marrow



B: Thymus

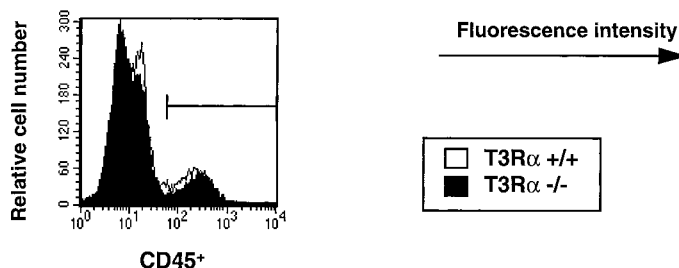


FIGURE 4. In vivo proliferation of bone marrow cells and thymocytes from T3R α KO mice. T3R α KO (filled histograms) or wild-type (open histograms) mice (17–19 days old) were injected twice with 1 mg BrdU at a 30-min interval. Six hours later, mice were sacrificed, and bone marrow (A) and thymic (B) cells were surface stained for B220 and IgM, or CD45 expression, respectively. BrdU-positive cells were then revealed, as described in *Materials and Methods*. Results for one representative experiment of three are shown.

KO mice and wild-type littermates were injected twice i.p. at a 30-min interval with 1 mg BrdU. This thymidine analogue is incorporated in the DNA of cycling cells. Incorporation of BrdU in bone marrow and thymic cells was measured 6 h after the last injection. Cells were first stained for surface markers to identify lymphocyte subsets. BrdU⁺ cells were then revealed by a FITC-conjugated anti-BrdU Ab, as described in *Materials and Methods*. PBS-injected animals were used as a negative control for background staining.

As shown in Fig. 4A and Table IV, the proportion of bone marrow B220⁺ cycling cells is decreased by 40% in T3R α KO mice. When looking at bone marrow subpopulations, as expected (29, 30), most of the cycling B lymphocyte precursors were contained in the pro/pre-B cell subset (Table IV). However, in T3R α KO animals, the percentage of BrdU⁺ cells was decreased in both the pro/pre-B cell (B220⁺IgM⁻) and the immature/mature B cell (B220⁺IgM⁺) subset by 30% and 50%, respectively, compared with wild-type controls.

To determine which subsets among pro/pre-B cells were affected in their cycling capacity, we have performed cell cycle analysis using Hoechst 33342 labeling on cells that were stained for B220, IgM, CD25, CD43, or *c-kit*. In T3R α KO mice, a small (15%) but significant ($p \sim 0.05$) reduction in the percentage of cycling cells was observed among all B cell progenitor subsets defined by these markers (Fig. 5, data not shown). Thus, T3R $\alpha 1$ and T3R $\alpha 2$ gene deletion affects the proliferation of all subsets of progenitor B cells. This effect was particularly obvious in large B220⁺CD25⁺ pre-BII cells, which represent the major subset of cycling cells (28) (Fig. 5B). This reduced proliferation might ex-

plain the defect in the number of small B220⁺CD25⁺ pre-BII cells observed in T3R α KO mice.

Thymocytes from T3R α KO mice also proliferate less in vivo than thymocytes from wild-type animals. Under the experimental conditions, using BrdU, the percentage of cycling thymocytes is decreased by about 20% (Fig. 4B).

T3R $\alpha 1$ or T3R $\alpha 2$ gene products control the size of the B cell pool by intrinsic mechanisms

As T3R $\alpha 1$ and T3R $\alpha 2$ are widely expressed among murine tissues (19, 31, 32), we questioned whether the deficient lymphocyte development was intrinsic to these cells or whether it was due to a defect in environmental factors. To address this question, we produced T3R $\alpha^{-/-}$ \times Rag1^{-/-} chimeric animals. T3R $\alpha^{+/-}$ mice were crossed and the resulting morulae were aggregated to morulae from Rag1-deficient mice. Rag1 is necessary for TCR and Ig gene recombination (33), and mice deficient for this enzyme are not able to develop mature T or B lymphocytes (34). Thus, in T3R $\alpha \times$ Rag1^{-/-} chimera, all mature B and T lymphocytes can only derive from the T3R α morulae, while all other cell types are a mix of cells derived from T3R α or Rag1^{-/-} morulae. The degree of chimerism was analyzed by PCR analysis on liver cells, as described in *Materials and Methods*. For the experiment described, it was estimated that Rag1^{-/-} T3R $\alpha^{-/-}$ chimeric animals contained about 60% to 75% of liver cells from Rag1^{-/-} (i.e., T3R $\alpha^{+/-}$ genotype) origin. This implies that in T3R $\alpha^{-/-} \times$ Rag1^{-/-} chimeric animals, mature lymphocytes that do not ex-

Table III. Analysis of bone marrow progenitor B cell subsets of T3R α KO and wild-type mice^a

Subpopulations	B220 ⁺ <i>c-kit</i> ⁺	B220 ⁺ <i>c-kit</i> ⁻	B220 ⁺ CD43 ⁺	B220 ⁺ CD43 ⁻	B220 ⁺ CD25 ⁺	B220 ⁺ CD25 ⁻	Larges, B220 ⁺ CD25 ⁺	Petites, B220 ⁺ CD25 ⁺
% (KO; wt) ^b	(1.2; 1.0)	(39.8; 46.6)	(3.2; 3.2)	(43.8; 50.0)	(12.1; 17.0)	(35.9; 37.2)	(2.1; 2.0)	(10.0; 15.0)
Ratio KO/wt ^c	118 [¥]	85*	102 [¥]	88*	71**	96 [¥]	107 [¥]	67**

^a Results are representative of one experiment out of two, including seven mutant and seven wild-type animals.

^b Bone marrow cells were stained with anti-B220, CD117 (*c-kit*), CD25 and CD43 antibodies and the percentages of each subset was determined using flow cytometry.

^c The percentage shown corresponds to: [(mean values for KO mice/mean values for wild-type mice)] \times 100.

¥, $p > 0.2$; *, $p < 0.05$; **, $p < 0.005$ Student's *t* test).

Table IV. Lymphocyte precursors in primary lymphoid organs from *T3Rα KO* mice proliferate less^a

Cell Types	% of BrdU ⁺ T3Rα KO	% of BrdU ⁺ Wild Type
Bone marrow		
B220 ⁺	10.9	18.9
B220 ⁺ IgM ⁻	20.9	29
B220 ⁺ IgM ⁺	2.3	4.9
Thymus		
CD45 ⁺	12.8	15.3

^a T3Rα KO and wild-type mice were injected twice with 1 mg BrdU. Six hours later, they were sacrificed and bone marrow and thymic cells were surface stained for B220 and IgM or CD45 expression. After permeabilization and DNA digestion, BrdU⁺ cells were revealed with an anti-BrdU FITC-conjugated Ab using flow cytometry. Results for one representative experiment out of three are shown.

press T3Rα1 and T3Rα2 have developed in a cellular environment in which a significant proportion of cells express these proteins. Hence, any decrease in lymphocyte cell numbers generated in these chimeric mice would indicate an intrinsic involvement of T3Rα1 or T3Rα2 gene products in the control of the development of these cells.

As shown in Fig. 6B, there is no significant difference in thymocyte numbers between T3Rα^{-/-} × Rag1^{-/-} and T3Rα^{+/+} × Rag1^{-/-} mice. In contrast, splenic cellular indices are significantly reduced by about 55% in T3Rα^{-/-} × Rag1^{-/-} mice compared with the T3Rα^{+/+} × Rag1^{-/-} control chimera. This decrease results mainly from a strong reduction in splenic B cell numbers (Fig. 6A). Indeed, splenic T cells, granulocytes, and macrophages are found in equivalent numbers in T3Rα^{-/-} × Rag1^{-/-} and T3Rα^{+/+} × Rag1^{-/-} mice. This indicates that the defect in T cell numbers observed in T3Rα KO mice was not intrinsic to T3Rα^{-/-} T cells and, more importantly, that the cellular environment provided in chimeric animals was able to support T lymphocyte, granulocyte, and macrophage development. Thus, these results indicate that B cells, in contrast to T cells, are direct targets for thyroid hormone regulation through the T3Rα receptor.

Discussion

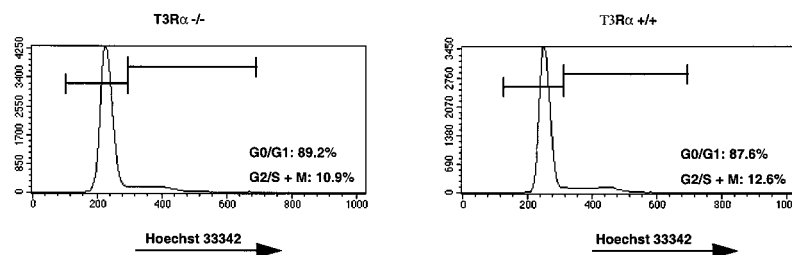
Defective lymphocyte development in T3Rα KO mice

The aim of this study was to determine the role of T3Rα receptor isoforms in the immune system. Mice deficient for T3Rα1 and T3Rα2 isoforms show a reduced T and B lymphocyte development that results in a decreased cellularity. However, lymphocyte development seems qualitatively normal, as no difference was observed in terms of surface phenotype and function. Indeed, in vitro, spleen lymphocytes from T3Rα KO mice show a normal proliferative response following LPS or anti-CD3 activation (data not shown). Furthermore, we show that T3Rα1 and T3Rα2 gene products are involved in the proliferation control of progenitor cells in primary lymphoid organs. This does not exclude a role for these proteins in the differentiation of the T or B lymphocyte precursor cells. The reduced lymphocyte cellularity observed in T3Rα KO mice could also result from an increased sensibility to programmed cell death (35). This does not seem to be the case, at least for bone marrow cells, as we did not find sub-G₁ apoptotic cells following Hoechst cell cycle analyses on freshly isolated bone marrow cells. Similarly, no detectable increase in apoptosis after in vitro bone marrow cell culture was found (data not shown).

Multiple levels of T₃ actions on the immune system development

Previous observations in Snell dwarf mice strongly supported a direct effect of T₃ on B cell development, but they could not completely exclude a possible indirect effect on GH/IGF-I production by the cellular environment. Indeed, thyroid hormones regulate GH/IGF-I production by enhancing GH gene transcription in a pituitary cell line (36). Furthermore, IGF-I has been shown to promote B lymphopoiesis (37, 38). Our results show that T₃ seems to act directly on B lymphocytes. This could be explained by the direct control by T3Rα1 or T3Rα2 of genes involved in B cell progenitor proliferation and/or differentiation. Indeed, it has been shown that bone marrow-derived B lineage cell lines express these isoform of the T₃ nuclear receptors (39). Still, one cannot exclude that T₃ acts only to up-regulate GH/IGF-1 secretion by B lineage cells (40, 41). However, this seems unlikely, as it has been shown

A: B220+



B: Large B220+CD25+

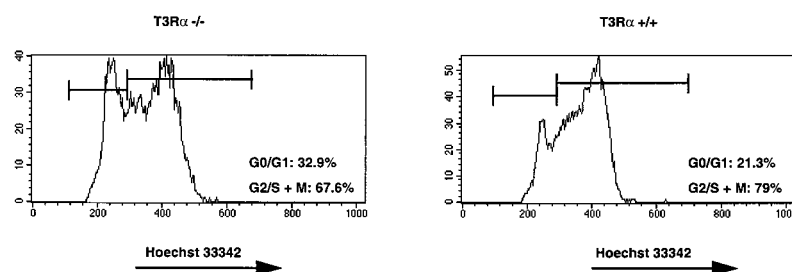
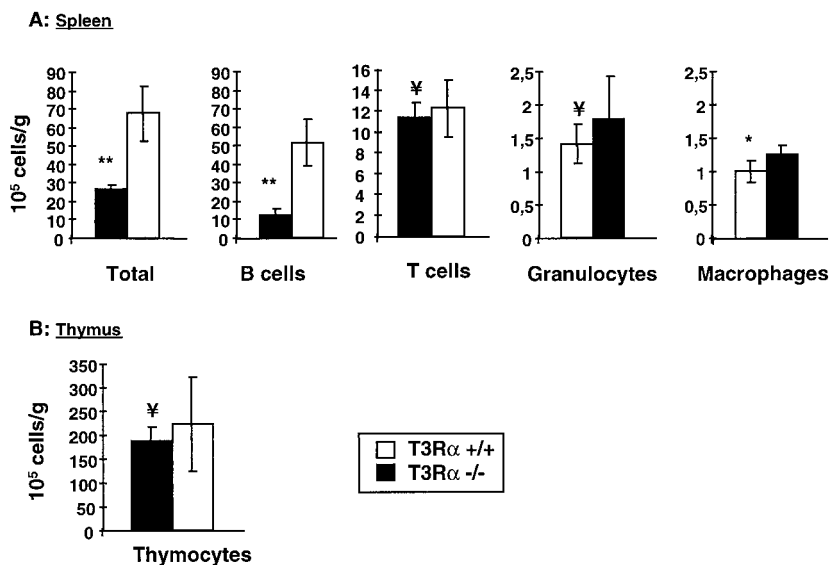


FIGURE 5. Cell cycle analysis of freshly isolated bone marrow nucleated cells from T3Rα KO mice. Bone marrow nucleated cells were first stained for B220 and CD25 expression. After ethanol fixation, cells were washed in PBS, incubated with Hoechst 33342 dye, and analyzed on FACs. Hoechst stainings of B220⁺ (A) and large B220⁺CD25⁺ (B) cells from T3Rα^{-/-} (left) and T3Rα^{+/+} (right) mice are shown. Representative of two experiments.

FIGURE 6. Lymphocyte numbers in T3R α KO \times Rag1 KO chimeric animals. T3R $\alpha^{-/-}$ and T3R $\alpha^{+/+}$ morulae were fused to Rag1 $^{-/-}$ morulae (see text for more details). The splenocytes (A) and thymocytes (B) from 17- to 19-day-old T3R $\alpha^{-/-}$ \times Rag1 $^{-/-}$ (■) and T3R $\alpha^{+/+}$ \times Rag1 $^{-/-}$ (□) chimeric offsprings were counted. A, Splenocytes stained with anti-B220 and anti-CD3 to identify B and T lymphocytes, respectively. Granulocytes and macrophages were identified by anti-Ly-6G anti-CD11b Abs, respectively. Cellularity indices were calculated as the number of cells per organ divided by the weight of the animals. The cellularity index values are expressed as 10⁵ cells/g. ¥, $p > 0.2$; *, $p < 0.1$; **, $p < 0.01$ (Student's t test).



that GH is unable to fully restore the pre-B cell compartment in Snell dwarf mice (12).

Conversely, defective T cell development in T3R α KO mice was shown to depend only on cellular environment, as it is fully restored in T3R $\alpha^{-/-}$ \times Rag1 $^{-/-}$ chimeric animals. This is in accordance with previous reports showing that thymic cellularity is enhanced by treatment with IGF-I (42) or GH (11, 43). Furthermore, these hormones are locally produced in the thymus (25, 44). T₃ could also influence thymocyte development through the regulation of other thymic-origin hormonal factors, such as thymulin (Zn-FTS), which induces thymocyte proliferation and differentiation (45). Indeed, the production of thymulin is correlated to seric concentrations of T₃ and T₄ (46, 47). However, T3R α 1 or T3R α 2 expression seems to influence the immune system development at multiple levels, as it acts directly on B lymphocytes and through the cellular environment on T lymphocytes.

In the case of myeloid cell development, we have found a deficit in macrophage (Mac-1⁺) and granulocyte (Gr-1⁺) splenic cell numbers in T3R α KO mice. However, T₃ again seems to act at another level, as it is unlikely that a defective myeloid central development accounts for this decreased macrophage and granulocyte splenic cellularity. Indeed, we did not observe any significant differences in the number of Gr-1⁺ and Mac-1⁺ bone marrow myeloid precursors (data not shown), as previously reported in Snell dwarf mice (12). As myeloid cells do not need Rag1 enzyme expression during their development, our experiments with chimeric animals did not allow us to identify the thyroid hormone target. However, a direct effect of T₃ on macrophages and granulocytes has been reported in a number of systems. Indeed, thyroid hormones modulate the metabolism and the functions of purified rat macrophages (48, 49) and stimulate phagocytosis by purified human polymorphonuclear granulocytes (50). Moreover, the production of TSH by cultured monocytes following thyrotropin-releasing factor stimulation is inhibited by T₃ (51).

The role of the different isoforms in the immune system development

The immune deficiencies observed in T3R α KO mice are directly attributable to the absence of the T3R α 1 and T3R α 2 gene products. Gene transcriptional regulation by T₃ nuclear receptor isoforms is complex. The first degree of complexity is given by the numerous isoforms of T₃ nuclear receptors, some of which do not

bind T₃ nor act as transcription factors. Furthermore, these isoforms act as homodimers or heterodimers together with other members of the nuclear receptor family (52, 53) and, depending on the targeted genes and the presence of their ligand, coactivators or corepressors either activate or repress gene transcription (17, 53–55). T₃ nuclear receptor isoforms can also inhibit one another's transcriptional activities (22, 56). Finally, T3R isoforms specifically retro-control thyroid hormone production in a complex fashion. Indeed, although thyroid hormones negatively retro-control pituitary TSH production, T3R β -deficient mice have elevated seric levels of T₃ and T₄, together with high TSH serum concentrations (20, 57), while the level of all three hormones is reduced after 3–4 wk of age in T3R α KO mice compared with wild-type littermates. (19). In T3R α 1-deficient mice (21), seric T₃ is unaffected, while T₄ seric concentrations are reduced only in male animals. It is important to note that in our experiments using 17- to 19-day-old animals, T₃ and T₄ seric concentrations were always found in the normal range, definitely establishing a specific role for T3R α 1 and T3R α 2 gene deletion in the observed phenotype.

Given the complexity of the regulation of gene transcription by nuclear hormone receptors, it would be of particular interest to correlate the hormonal status and immune deficiencies in different mice deficient for T3R isoforms and other members of the nuclear receptor family. We have preliminary data showing a decreased granulocyte, macrophage, and T and B lymphocyte cellularity in spleens from T3R β KO mice (data not shown). Our preliminary experiments also show drastically reduced numbers of these cells in the spleens of mice deficient for all T3R isoforms, indicating an additive effect of T3R α and T3R β gene deletions (data not shown). This further supports a role for T₃ in B lymphocyte development and suggests a partial redundancy between thyroid hormone α and β receptors. It has also been shown that T3R α plays a major role in the control of the proliferation of the crypt cells in the intestine, and that partial redundancy between T3R α and T3R β also occurs in this system (58). Moreover, additive effect of T3R α and T3R β gene deletions excludes the possibility that in absence of T3R α 1, T₃ would act through T3R β to reduce cell cycle progression. Indeed, it has been shown that thyroid hormones regulate proliferation arrest and differentiation of oligodendrocyte precursor cells and that the level of T3R β 1 receptor expression in these cells is involved in determining the time when they will withdraw from cell cycle and differentiate (59).

In conclusion, this study shows that T3R α 1 and T3R α 2 gene products are involved in immune system development. It also shows for the first time that expression of these receptors in bone marrow B lineage cells is intrinsically needed for a proper B lymphopoiesis. Finally, it supports increasing amounts of data demonstrating the importance of cross-talks between neuro/endocrine and immune systems. In terms of functional consequences of these receptors' defect, we have found that T3R α -deficient lymphocytes proliferate after polyclonal activation (data not shown), but we have not been able to test Ag-specific responses in T3R α KO mice that die within a few weeks after birth. However, this needs to be investigated in the different thyroid hormone-deficient mice. Indeed, in humans with thyroid hormone resistance due to mutations in T3R β receptors, a cohort study showed a high frequency of nose, ear, and throat infections, which could indicate a partial impairment of their immune system (60).

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