Expression of genes encoding the pre-TCR and CD3 complex during thymus development

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Abstract

The mature TCR is composed of a clonotypic heterodimer ($\alpha\beta$ or $\gamma\delta$) associated with the invariant CD3 components (γ , δ , ε and ζ). There is now considerable evidence that more immature forms of the TCR-CD3 complex (consisting of either CD3 alone or CD3 associated with a heterodimer of TCR β and pre-T α) can be expressed at the cell surface on early thymocytes. These pre-TCR complexes are believed to be necessary for the ordered progression of early T cell development. We have analyzed in detail the expression of both the pre-TCR and CD3 complex at various stages of adult thymus development. Our data indicate that all CD3 components are already expressed at the mRNA level by the earliest identifiable (CD4^{lo}) thymic precursor. In contrast, genes encoding the pre-TCR complex (pre-T α and fully rearranged TCR β) are first expressed at the CD44^{lo}CD25⁺CD4⁻CD8⁻ stage. Detectable surface expression of both CD3 and TCR β are delayed relative to expression of the corresponding genes, suggesting the existence of other (as yet unidentified) components of the pre-TCR complex.

Introduction

The TCR is a complex, multisubunit structure that consists of at least six chains (α and β , or γ and δ , as well as the CD3 γ , δ , ϵ and ζ chains) that are assembled in the endoplasmic reticulum and transported to the cell surface (reviewed in 1–5). Whereas the α , β , γ and δ chains of the TCR are clonotypic and confer ligand specificity, the CD3 subunits are invariant and function as signal transducing molecules through interactions of their cytoplasmic domains with the protein tyrosine kinases ZAP-70 or p56^{*lck*} (6–8)

Efficient surface expression of a functional TCR–CD3 complex on mature T cells has been shown to require all six chains (9). In the absence of either TCR α or TCR β (10,11), mature T cells bearing functional TCR $\alpha\beta$ –CD3 complexes do not develop. In mice deficient for CD3 ζ (12–15), a similar absence of mature, functional $\alpha\beta$ T cells is observed. While it is generally accepted that surface expression of a functional CD3–TCR complex occurs uniquely on mature T cells, several lines of evidence suggest that low levels of CD3 may be expressed on the surface of immature thymocytes either alone or in association with TCR β . Thymocyte development in TCR $\alpha^{-\!\!\!/}$ mice is arrested at the CD4+CD8+ double-positive (DP) stage (10,11). While no TCR $\alpha\beta$ heterodimer is expressed,

low levels of CD3 and TCR β can be detected on these DP cells by FACS analysis. Furthermore, low levels of CD3 ϵ protein have been detected at even earlier stages of development, both intracellularly in CD44^{lo}CD25⁺ double-negative (DN) thymocytes (16) and on the surface of CD44^{lo}CD25⁻ DN thymocytes (17,18). In addition, studies in the normal human thymus (19) and in acute lymphoblastic leukemias (20,21) have also demonstrated the presence of CD3 ϵ protein in immature T cells, although the exact stage of development was not determined. Finally, biochemical studies have shown that CD3 ϵ , γ and δ (in the presence or absence of TCR β chains), may be detected on the surface of SCID thymocytes (22), SCID cell lines (23) and RAG-1^{-/-} thymocytes (24), all of which are principally of the CD44^{lo}CD25⁺ DN phenotype.

While the exact composition of this immature CD3 complex is not known, it has been shown that signals can be transduced in immature thymocytes by its ligation. Treatment of RAG^{-/-} thymocytes with anti-CD3¢ mAb, either *in vivo* (24,25) or *in vitro* (26), induces expansion and differentiation of CD44^{lo}CD25⁺ DN cells to the DP stage in the absence of TCR rearrangement. RAG^{-/-} or SCID thymocytes can also be

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induced to expand and differentiate to the DP stage under the influence of either a TCR β transgene (24,27,28) or a p56^{*lck*} transgene (29). Taken together, these results in both normal and mutant mice provide strong evidence for the presence of at least some components of the TCR–CD3 complex on immature thymocytes.

A pre-TCR complex, consisting of a rearranged TCR β chain and the recently cloned pT α chain (30), associated with the CD3 complex has recently been described (23). pT α , which is believed to be a key component of the pre-TCR, is expressed in immature DN and DP thymocytes of TCR $\alpha^{-/-}$ and RAG-2^{-/-} mice and not on mature T cells. In this study we have investigated in detail the expression of the genes encoding the pre-TCR and CD3 complex during early adult thymus development.

Methods

Mice and thymocyte subsets

Normal C57Bl/6 female mice were purchased from Harlan/ Olac (Bicester, UK) and used at 4--6 weeks of age. Thymocyte subsets were purified by complement-mediated cytotoxicity and magnetic bead depletion (31,32), and further sorted on a FACStar Plus (Becton Dickinson, Mountain View, CA).

Northern blot analysis and in situ hybridization

Total cell RNA was prepared by CsCl₂ gradient, electrophoresed on formaldehyde-agarose minigels and hybridized with RNA probes as described previously (31). In situ hybridization was performed on single cells FACS-sorted directly onto microscope slides (31). The probes used were as follows: TCR C₆2 and β -actin (31); CD3 ϵ , a 1200 bp Xbal-BamHI insert in pBS.KSII; CD3 y, a 700 bp EcoRI-HindIII insert in pBS.KSII; CD3 δ , a 800 bp Xhol insert in pBS.KSII; CD3 ζ , a 1100 bp EcoRI insert in pBS.SK (these latter four probes were the kind gift of Bernard and Marie Malissen, Marseille-Luminy, France); and pTa which is a 660 bp insert in Bluescript KS⁺ made by PCR amplification of cDNA from CD25⁺ DN thymocytes using the published primer sequences (30). All probes were labeled by RNA transcription with either $\left[\alpha\right]$ ³²P]UTP (for Northern blot) or $[\alpha$ -³⁵S]UTP (for *in situ*). P815 mastocytoma cells were used as a negative control for in situ hybridization. According to the criteria used, ~2-3% cells were positive by in situ hybridization for each of the CD3 probes. Densitometry readings of the exposed X-ray films was performed with the Elscript 400 densitometer (Hirschmann-Getaebau, Unterhachingen, Germany) and the results analyzed with the accompanying software.

FACS analysis

Four-color FACS analysis was performed on a FACStar Plus equipped with a standard argon laser (for FITC, PE and Red-613 in FL1, 2 and 3), and a helium-neon laser for allophycocyanin (APC) (FL4). Fluorescent conjugates were as follows: APC-streptavidin (Molecular Probes, Eugene, OR); control hamster Ig-PE, PE-streptavidin and anti-CD4-APC (Caltag, South San Francisco, CA); anti-CD3-PE, anti-CD4-Red-613, anti-CD8-Red-613 and anti-CD25-Red-613 (Gibco/ BRL, Gaithersberg, MD); anti-TCRβ-PE and HSA-PE (PharM- ingen, San Diego, CA); anti-CD44–FITC, anti-HSA–biotin and anti-CD25–biotin were prepared in this laboratory.

Results

CD3 mRNA expression by immature thymus subsets

During adult thymus development, immature T cells of the $\alpha\beta$ lineage follow a complex developmental program in which differentiation is accompanied by TCR gene rearrangement and expression. The earliest identifiable thymocyte expresses low levels of CD4 (but not CD8 or CD3) and has its TCR genes in germline configuration (33) CD4 is down-regulated and the cells progress through a series of DN (CD4-CD8-) subsets characterized by the differential expression of CD44 and CD25 in the order: CD44^{hi}CD25⁻ to CD44^{hi}CD25⁺ to CD44^{lo}CD25⁺ to CD44^{lo}CD25⁻ (34-39) Transition from the late DN (CD4⁻CD8⁻) to the DP (CD4⁺CD8⁺) stage can occur via an immature single positive (ISP) and is accompanied by expansion (reviewed in 40). Later stages of thymus development involve repertoire selection, up-regulation of a mature TCR-CD3 complex and production of mature single-positive (SP) CD4⁺ or CD8⁺ cells (41-43).

In order to determine at which stage of T cell development in the adult thymus CD3 ε , γ , δ and ζ genes are first expressed, we performed *in situ* hybridization studies at the single cell level on thymus subsets. As shown in Table 1, a substantial proportion (53–90%) of cells in the earliest identifiable thymocyte population (the CD4^{lo} precursor), were positive for each of the CD3 genes. The percent positive cells expressing each of these genes increased gradually in subsequent populations to reach essentially 100% in the CD44^{lo}CD25⁺ DN subset and remained stable throughout further stages of thymus development as well as in mature T cells.

As virtually all cells from the CD44^{lo}CD25⁺ DN subset onwards were positive for CD3 ϵ , γ , δ and ζ chain mRNA, it was interesting to examine whether the level of expression was similar in each subset. To this end, Northern blot analysis was performed on total RNA prepared from purified thymocyte subsets commencing at the CD44^{lo}CD25⁺ DN stage. As shown in Fig. 1 and Table 2, CD3 ϵ and CD3 δ were expressed in approximately equivalent amounts in all subsets. By contrast, CD3 γ was expressed at higher levels in the CD44^{lo}CD25⁺ DN than in the later immature (CD44^{lo}CD25⁻

Table 1. CD3 gene expression in immature thymocyte subsets

Thymocyte subset ^a	Cells positive (%) ^b						
	CD3ε	CD3γ	CD38	СДЗС			
CD4 ⁴⁰ precursor CD44 ^{h1} CD25 ⁻ CD44 ^{h1} CD25 ⁺ CD44 ^{b2} CD25 ⁺ CD44 ^{b2} CD25 ⁻	80 ± 5 90 ± 7 91 ± 3 100 100	88 ± 9 97 ± 3 97 ± 5 99 ± 4 100	53 ± 10 64 ± 7 89 ± 3 98 ± 3 100	69 ± 6 82 ± 8 97 ± 3 100 100			

^aAll more mature subsets and mature T cells were ~100% positive. ^bFor each probe 500–1000 cells were counted on each slide and two to four separate slides analyzed. DN, ISP, DP) and more mature (SP and lymph node T cells) subsets. The levels of CD3 ζ mRNA expression, while substantially lower than those of CD3 ϵ , γ and δ , also varied considerably during development. Maximum expression of ζ was observed at the DP stage, much later than the peak expression of CD3 γ The maximum levels of expression of



Fig. 1. Expression of CD3 genes in thymocyte subsets by Northern blot analysis. Sequential hybridizations of the same blot were performed with probes for CD3 ε , γ , δ , ζ and β -actin. Control cells were polyclonally activated $\gamma\delta$ thymocytes, P815 mastocytoma and $\alpha\beta$ CTL clone. Exposure times were overnight for all probes

CD3 γ at the CD44^{lo}CD25⁺ DN stage, coupled with the fact that virtually all cells at this stage express mRNA from all genes of the CD3 complex, is consistent with the possibility of surface CD3 expression as suggested by both biochemical and functional studies (16,22–24).

Expression of $pT\alpha$ and TCR β mRNA in immature thymocytes

The recently identified $pT\alpha$ chain has been shown to be expressed as a heterodimer with TCR B as part of the CD3associated pre-TCR complex (23,30). In order to determine at which stage during T cell development TCR B and pTa are first expressed, we performed Northern blot analysis of early thymus subsets As shown in Fig. 2, pTa is first detected in CD44^bCD25⁺ DN cells and not in earlier subsets Expression continues through to the DP stage, but no $pT\alpha$ mRNA is detected in either mature SP thymocytes (Fig. 2) or lymph node T cells (data not shown). These results are consistent with those obtained recently by PCR analysis (30). Full length (VDJC) TCR B transcripts are also first detected at the CD44^{lo}CD25⁺ DN stage, although both truncated (DJC) and longer (presumably germline) transcripts are detected earlier in the CD44^hCD25⁺ subset (Fig. 2). Taken together with the CD3 data, these results are consistent with the potential expression of a pre-TCR-CD3 complex by immature thymocytes beginning at the CD44^bCD25⁺ DN stage.

Surface expression of CD3 ϵ and TCR β on immature thymocytes

Although high levels of CD3 ϵ , γ , δ and ζ mRNA are expressed through all stages of thymocyte development (Fig. 1 and Tables 1 and 2), and full length TCR β transcripts are produced at the CD44^{lo}CD25⁺ DN stage (31,35 and Fig. 2), a mature CD3–TCR $\alpha\beta$ complex is not expressed on the surface of immature thymocytes due to the absence of TCR α chains However, intracellular CD3 ϵ protein can be detected in CD44^{lo}CD25⁺ DN thymocytes (data not shown and 16) and evidence described above suggests that a functional CD3 complex may be expressed, albeit at low levels, on this subset We therefore decided to determine at what stage of development CD3 ϵ and TCR β could be detected at the cell surface.

To this end, four-color FACS analysis was performed on thymocyte subsets from the CD44^{lo}CD25⁺DN through to mature SP stages. As shown in Fig. 3, neither CD3 ϵ nor TCR β are detectable in the CD44^{lo}CD25⁺ DN subset; however, surface CD3 ϵ and TCR β are detectable in the CD44^{lo}CD25⁻ DN stage. These results are consistent with the expression

Table 2. (Quantitation	of CD3	gene	expression	in th	vmocvt	e subsets
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Probe	$\gamma \delta^+$ thymocytes	P815	αβ ⁺ CTL clone	CD25 ⁺ DN	CD25 ⁻ DN	ISP	DP	SP	Lymph node T
CD3e	1.1	0	1.1	3.2	1.6	1.0	2.3	25	1.0
CD3y	1.2	0	0.9	18.1	2.8	32	2.3	2.7	1.0
CD38	0.9	0	0.7	3.0	13	18	27	2.3	1.0
CD3ζ	1.0	0	1.2	2.3	2.0	1.8	6.6	4.0	1.0

Densitometry was performed on the autoradiographs in Fig. 1. For each subset densitometric values were divided by the β -actin control. Data for each of the CD3 probes are arbitrarily normalized to a value of 1.0 for lymph node T cells.

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Fig. 2. Expression of TCR β and pT α in thymocyte subsets. Thymocyte subsets and control cells are as in Fig 1, with the addition of the two earlier CD44^{hl} DN subsets. The blot was hybridized sequentially with TCR C $_{\beta}$, pT α and β -actin probes. Exposure times were 3 days for the first TCR β panel, overnight for the second TCR β and for β -actin, and 3 weeks for pT α .

subsets CD25⁺ DN and CD25⁻ DN subsets were further gated to be CD44^{lo} and HSA^{hI} ISP and SP correspond to CD4⁻CD8⁺HSA^{hi} and CD4⁻CD8⁺HSA^b subsets respectively Control staining (indicated by the thin lines) is harmster Ig-PE

of CD3 ϵ and TCR β on immature thymocytes as part of a pre-TCR complex (30).

Discussion

The importance of the CD3 complex for T cell development has been firmly established by experiments using transgenic or knockout mice. Indeed, overexpression of a human CD3 ɛ transgene in mice inhibits T cell development in a copy number dependent fashion (44) At low copy numbers, T cell development is blocked at the CD44^{lo}CD25⁺ DN stage (similar to RAG^{-/-} mice). In contrast, at high copy number, the block is earlier at the CD44^{hi}CD25⁻ stage, raising the possibility that CD3 ϵ may play a role even earlier in T cell development than either the recombinase genes or TCR β . However, in CD3 ɛ deficient mice, T cell development is not arrested until the CD44^{lo}CD25⁺ DN stage (Malissen, personal communication), rather suggesting that this is the first stage where the presence of functional CD3 c protein is required. This apparent discrepancy could be explained if overexpression of the human CD3 ɛ transgene interfered with development indirectly (e.g. by competing for kinases such as p56^{kck})

In contrast to CD3 ϵ , T cell development in CD3 ζ^{-1} mice (12–15) is arrested later at the DP stage. In addition, when CD3 ζ is overexpressed, development is also blocked at the

DP stage in a copy-number-dependent manner (45). In this latter study, the authors suggested that the developmental block was due to premature termination of RAG-1 and RAG-2 expression preventing productive rearrangements of TCR α and TCR β It is of interest that the DP stage of development where the effects of overexpression or absence of CD3 ζ are manifest is correlated with the maximum expression of CD3 ζ (Fig. 1 and Table 2). These results suggest that while CD3 ζ is essential for the DP to SP transition (most likely due to its role in signal transduction), it is less important for the earlier DN to DP transition. In the absence of CD3 ζ, DP cells are produced but their numbers are decreased compared with normal mice. In addition, the CD44^{lo}CD25⁻ DN subset, which is the immediate precursor of the ISP (pre-DP) subset, and the major cycling DN population in normal mice, is absent in these mice (46). These latter data suggest that CD3 ζ may have a specific role in the control of proliferation of DN thymocytes.

At the mRNA level, CD44^bCD25⁺ DN thymocytes are the most immature subset that expresses all known CD3 components as well as pT α and full length TCR β . Yet this subset has no easily detectable surface CD3 or TCR β expression. In fact, clear expression of CD3 (measured by anti-CD3 ϵ mAb) and TCR β occur later at the CD25⁻ DN stage. Several explanations for the delayed appearance on the surface of expressed CD3 ϵ and TCR β chains can be entertained. For example, it is possible that limitations in posttranscriptional processing or assembly of CD3 or pre-TCR complexes may delay their membrane insertion Alternatively, the pre-TCR complex may (by analogy with pre-B cells) contain other as yet unidentified components, such as the recently proposed V pre-T (30). In either case it is obvious that the fully assembled (CD3 or pre-TCR) complexes could not be expressed at the cell surface until all components became available.

Finally it is informative to consider our data in the context of other studies of TCR $\alpha\beta$ transgenic mice. Thus we (32) and others (36) have shown that surface TCR $\alpha\beta$ (or CD3) expression in such mice is only apparent at the CD25⁻ DN stage, despite the fact that the TCR α and β transgenes are presumably expressed much earlier at the mRNA level. This surprising result, which was obtained independently of the nature of the TCR transgenic construct (cDNA in our case and rearranged genomic DNA in the other), suggests that surface expression of a mature TCR $\alpha\beta$ -CD3 complex is likewise developmentally regulated. It therefore appears that neither the pre-TCR nor the mature TCR $\alpha\beta$ complex can be efficiently expressed on the cell surface prior to the CD25-DN stage. To explain this paradox, one could speculate that the CD3 complex exists in two alternate configurations, an early form which is only permissive for CD3 expression (perhaps in association with another unknown molecule) and a later form which can associate with either the pre-TCR or mature TCR. Such a model would imply that the differential expression of pre-TCR and TCR complexes during development depends upon the regulated expression of $pT\alpha$ and TCR a genes.

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Abbreviations

APC	allophycocyanin
DP	double positive
DN	double negative
ISP	immature single positive
SP	single positive
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