

Specific requirement for CD3 ϵ in T cell development

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Edited by Frederick W. Alt, Harvard Medical School, Boston, MA, and approved October 14, 1998 (received for review August 5, 1998)

ABSTRACT T cell antigen receptor (TCR) and pre-TCR complexes are composed of clonotypic heterodimers in association with dimers of signal transducing invariant subunits (CD3 γ , δ , ϵ , and ζ). The role of individual invariant subunits in T cell development has been investigated by generating gene-specific mutations in mice. Mutation of CD3 γ , δ , or ζ results in an incomplete block in development, characterized by reduced numbers of mature T cells that express low levels of TCR. In contrast, mature T cells are absent from CD3 ϵ ^{-/-} mice, and thymocyte development is arrested at the early CD4⁻CD8⁻ stage. Although these results suggest that CD3 ϵ is essential for pre-TCR and TCR expression/function, their interpretation is complicated by the fact that expression of the CD3 γ and CD3 δ genes also is reduced in CD3 ϵ ^{-/-} mice. Thus, it is unclear whether the phenotype of CD3 ϵ ^{-/-} mice reflects the collective effects of CD3 γ , δ , and ϵ deficiency. By removing the selectable marker (PGK-NEO) from the targeted CD3 ϵ gene via Cre/loxP-mediated recombination, we generated mice that lack CD3 ϵ yet retain normal expression of the closely linked CD3 γ and CD3 δ genes. These (CD3 ϵ ^{Δ/Δ}) mice exhibited an early arrest in T cell development, similar to that of CD3 ϵ ^{-/-} mice. Moreover, the developmental defect could be rescued by expression of a CD3 ϵ transgene. These results identify an essential role for CD3 ϵ in T cell development not shared by the CD3 γ , CD3 δ , or ζ -family proteins and provide further evidence that PGK-NEO can influence the expression of genes in its proximity.

Differentiation of thymocytes into mature, functional T cells requires the input of signals delivered through the T cell antigen receptor (TCR) and a precursor form of the TCR, the pre-TCR (1, 2). The TCR complexes expressed on mature T cells are composed of subunits originating from six different genes: TCR α and TCR β (or TCR γ and TCR δ), CD3 γ , CD3 δ , CD3 ϵ , and ζ (or a related family member) (3). The clonotypic (TCR α/β or TCR γ/δ) chains, which specify distinct lineages of T cells, are responsible for ligand recognition and are generated during development by programmed rearrangement of germline [V-(D)-J] gene segments. TCR α/β and TCR γ/δ heterodimers lack inherent signaling activity but associate noncovalently with dimers composed of the invariant signal transducing CD3 and ζ -family subunits. The generally accepted stoichiometry for the TCR complex is TCR α/β or TCR γ/δ /CD3 $\gamma\epsilon$ /CD3 $\delta\epsilon$ / $\zeta\zeta$.

Rearrangement of genes encoding the lineage-specific $\alpha\beta$ - and $\gamma\delta$ -TCR chains is initiated in immature CD4⁻CD8⁻ [or double negative (DN)] thymocytes that constitutively express the CD3 γ , CD3 δ , CD3 ϵ , and ζ subunits (4). Productive rearrangement of both the TCR γ and TCR δ genes results in surface expression of $\gamma\delta$ TCR complexes and commitment of

DN thymocytes to the $\gamma\delta$ -T cell lineage (5). On the other hand, productive rearrangement of the TCR β locus results in surface expression of a “pre-TCR” complex composed of TCR β chain paired with an invariant, pre-T α chain in association with CD3 and ζ -chain dimers (2). Signals transduced by the pre-TCR induce commitment of DN thymocytes to the $\alpha\beta$ -T cell lineage by triggering cell proliferation (resulting in “ β -selection”) and transition to the CD4⁺CD8⁺ [or double positive (DP)] stage, in which TCR α gene rearrangement is initiated (2, 6).

The role of each of the TCR subunits in T cell development has been examined by generating gene-specific mutations in mice. Mice rendered deficient in [V-(D)-J] recombination (Rag1^{-/-} or Rag2^{-/-}) are devoid of T cells and contain only immature DN thymocytes (7, 8) whereas mutations that selectively block expression of the TCR α/β or TCR γ/δ chain heterodimers result in lineage-restricted defects (i.e., absence of α/β or γ/δ T cells, respectively) (6, 9, 10). Moreover, α/β -T cell development is arrested at the DN stage in TCR β ^{-/-} mice and at the DP stage in TCR α ^{-/-} mice (6, 9), consistent with a requirement for TCR β but not TCR α for pre-TCR assembly and function (2).

Mice lacking all members of the ζ -family (ζ , η , Fc ϵ RI γ) exhibit a partial block in T cell development at both the DN→DP and DP→CD4⁺CD8⁻ or CD4⁻CD8⁺ transitions (11). Small numbers of peripheral T cells are generated in ζ/η /Fc ϵ RI γ ^{-/-} mice despite the fact that they express extremely low levels of TCR (11). Thus, pre-TCR and TCR complexes can still be expressed and are capable of transducing developmental signals, albeit much less efficiently, in the absence of ζ -family dimers (11).

The phenotype of CD3 ϵ ^{-/-} mice has shown that, unlike the ζ -family dimers, CD3 dimers are absolutely required for pre-TCR and TCR assembly and/or signal transduction (12). Both $\alpha\beta$ TCR⁺ and $\gamma\delta$ TCR⁺ T cells are absent from CD3 ϵ ^{-/-} mice, and, although rearrangement of TCR β occurs, thymocyte development is arrested at the immature DN stage (12). In contrast, mutation of either CD3 γ or CD3 δ results in a less severe developmental impairment (13, 14). As in ζ/η /Fc ϵ RI γ ^{-/-} mice, both CD3 γ ^{-/-} and CD3 δ ^{-/-} mice exhibit an incomplete block in development and contain low numbers of peripheral $\alpha\beta$ TCR^{lo} T cells (11, 13, 14). In addition, CD3 δ ^{-/-} mice contain normal numbers of DP thymocytes, and $\gamma\delta$ -T cell development appears unaffected (13). One interpretation of these results has been that CD3 γ and CD3 δ are functionally redundant such that CD3 $\gamma\epsilon$ dimers or CD3 $\delta\epsilon$ dimers alone can, though with different efficiency, promote pre-TCR and TCR assembly. On the other hand, CD3 ϵ is absolutely required because, in its absence, neither heterodimer can be formed.

This paper was submitted directly (Track II) to the *Proceedings* office. Abbreviations: DN, double negative (CD4⁻CD8⁻); DP, double positive (CD4⁺CD8⁺); TCR, T cell antigen receptor; ES cell, embryonic stem cell.

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0027-8424/98/9514909-6\$0.00/0

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However, a potentially confounding factor in the interpretation of these data has been that thymocytes from $CD3\epsilon^{-/-}$ mice were found to express abnormally low levels of $CD3\gamma$ and $CD3\delta$ transcripts (12). Thus, it is has remained unclear whether the phenotype observed in $CD3\epsilon^{-/-}$ mice reflects a specific requirement for $CD3\epsilon$ in early T cell development or the collective effects of $CD3\gamma$, $CD3\delta$, and $CD3\epsilon$ deficiency. In this study, we generated mice that lack expression of $CD3\epsilon$ yet retain normal expression of the closely linked $CD3\gamma$ and $CD3\delta$ genes. Significantly, the developmental defect in these mice was found to be identical to that of the previously reported $CD3\epsilon^{-/-}$ mice and could be rescued by expression of a $CD3\epsilon$ transgene. Collectively, these results identify an absolute requirement for $CD3\epsilon$ in early T cell development not shared by the $CD3\gamma$, $CD3\delta$, or ζ -family proteins.

MATERIALS AND METHODS

Generation of $CD3\epsilon^{N/N}$, $CD3\epsilon^{\Delta/\Delta}$, and $CD3\epsilon^{tg}$ Mice. The $CD3\epsilon$ targeting vector $p\epsilon NEO-loxP$ (Fig. 1A) was constructed by subcloning genomic DNA fragments that contained $CD3\epsilon$ exons 4 and 5 (5' flank; 3.5 kilobases) and exons 7 and 8 (3' flank; 3.0 kilobases) into plasmid pZINIMn, which contains both the $loxP$ -PGK-NEO- $loxP$ and PGK-HSV-TK expression cassettes. Embryonic stem (ES) cell electroporation and selection for homologous recombinants was performed essentially as described (15). $CD3\epsilon^{\Delta/\Delta}$ mice (see allele designations below) were generated by mating $CD3\epsilon^{N/N}$ mice with EIIa-*cre* transgenic mice (16) and then screening offspring for loss of the PGK-NEO cassette. The $CD3\epsilon$ transgene was generated by substituting the murine $CD3\epsilon$ coding sequences for the ζ cDNA sequences in construct ζ -CT108 (17). Four $CD3\epsilon$ transgenic founder lines were generated by zygote injection, and $CD3\epsilon$ protein expression was quantitated by Western blotting. Two founder lines [C7530 (2 \times expression relative to wild-type), and C7705 (0.25 \times expression relative to wild-type)] were used in the present experiments.

PCR Screening. Mice were genotyped initially by Southern blotting then subsequently by using a competitive PCR reaction that distinguishes between the $CD3\epsilon^+$ (wild-type), $CD3\epsilon^N$ (mutant $CD3\epsilon$ with PGK-NEO insertion), and $CD3\epsilon^{\Delta}$ (mutant $CD3\epsilon$ without PGK-NEO) alleles (Fig. 2). PCR was performed with a mixture of oligonucleotide 1 ($CD3\epsilon$ -exonV/5'-3'; TACAAAGTCTCCATCTCAGG), 2 ($CD3\epsilon$ -exonVII/3'-5'; TGGCCGCTCCTTGTTTTG), 3 ($CD3\epsilon$ -exonV/3'-5'; CTCGAGCTTTCAGGTACAA), and 4 (NEO; GGATTAGATAAATG CTTGCT). PCR parameters were 95°C, 20''; 55°C, 20''; 72°C, 90'' \times 35 cycles. Products were resolved on a 2% agarose gel and were visualized by staining with ethidium bromide.

RNA and Protein Analysis. For Northern blot analysis, RNA was purified from total thymocytes, was electrophoresed, was transferred to membranes, and was hybridized with $CD3\gamma$ -, $CD3\delta$ -, $CD3\epsilon$ -, and ζ -specific probes as described (17). The ef-1 α probe was generated by PCR of total embryo (fetal day 9.5) cDNA with oligos ef-1 α A and ef-1 α B (18). For Western blot analysis, thymocytes were enumerated, were lysed at a concentration of 1×10^8 /ml in SDS-loading buffer plus β -mercaptoethanol, and were resolved by 15% SDS/PAGE. Separated proteins were transferred to poly(vinylidene difluoride) membranes, were blotted with a 1:100 dilution of polyclonal anti- $CD3\epsilon$ (A452; Dako), and were detected by chemiluminescence (ECL, Amersham).

Multicolor Flow Cytometry. Single-cell suspensions of thymocytes or lymph node cells were processed, stained, and analyzed as described (17). mAbs used for flow cytometric analysis (purchased from PharMingen unless noted otherwise) included unlabeled anti- $CD16/CD32$ (2.4G2; blocking antibody), and fluorescein isothiocyanate- or phycoerythrin conjugated anti- $CD4$ (RM4.5), anti- $CD8\alpha$ (53-6.7), anti-TCR β (H57-597), anti-TCR δ (GL-3), anti- $CD3\epsilon$ (145-2C11), anti- $CD25$ (7D4), anti-B220 (RA3-6B2), and anti- $CD44$ (IM7). Quantum-red conjugated anti- $CD4$ and anti- $CD8$ were purchased from Sigma.

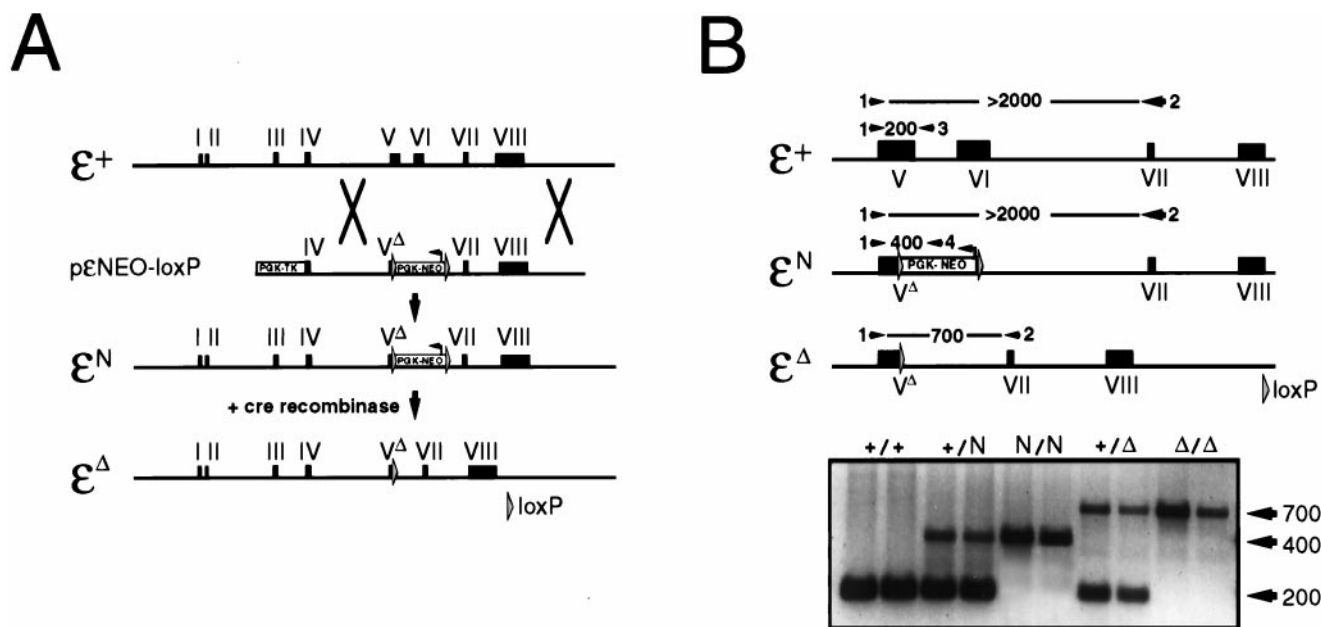


FIG. 1. Generation of $CD3\epsilon^{N/N}$ and $CD3\epsilon^{\Delta/\Delta}$ mice. (A) Representations of the $CD3\epsilon^+$ (wild-type), $CD3\epsilon^N$ (PGK-NEO⁺ mutant), and $CD3\epsilon^{\Delta}$ (PGK-NEO⁻ mutant) alleles. The $CD3\epsilon^N$ mutation was generated in ES cells by homologous recombination with the vector $p\epsilon NEO-loxP$ as shown. The $CD3\epsilon^{\Delta}$ allele subsequently was generated from the $CD3\epsilon^N$ allele by Cre/ $loxP$ -mediated recombination as described in *Materials and Methods*. (B) Competitive PCR strategy for distinguishing between the $CD3\epsilon^+$, $CD3\epsilon^N$, and $CD3\epsilon^{\Delta}$ alleles. Mouse tail DNA was amplified with a mixture of oligonucleotides 1-4, and the products were visualized by gel electrophoresis. Allele-specific products were distinguished on the basis of size: $CD3\epsilon^+$ (200-bp product of oligos 1+3); $CD3\epsilon^N$ (400-bp product of oligos 1+4); and $CD3\epsilon^{\Delta}$ (700-bp product of oligos 1+2). Although oligos 1+2 also can amplify a product from both the $CD3\epsilon^+$ and $CD3\epsilon^N$ alleles, the large size of the product (>2,000 bp) precludes its accumulation under the competitive PCR conditions. Results shown are from two independent DNA samples obtained from mice of the indicated genotypes.

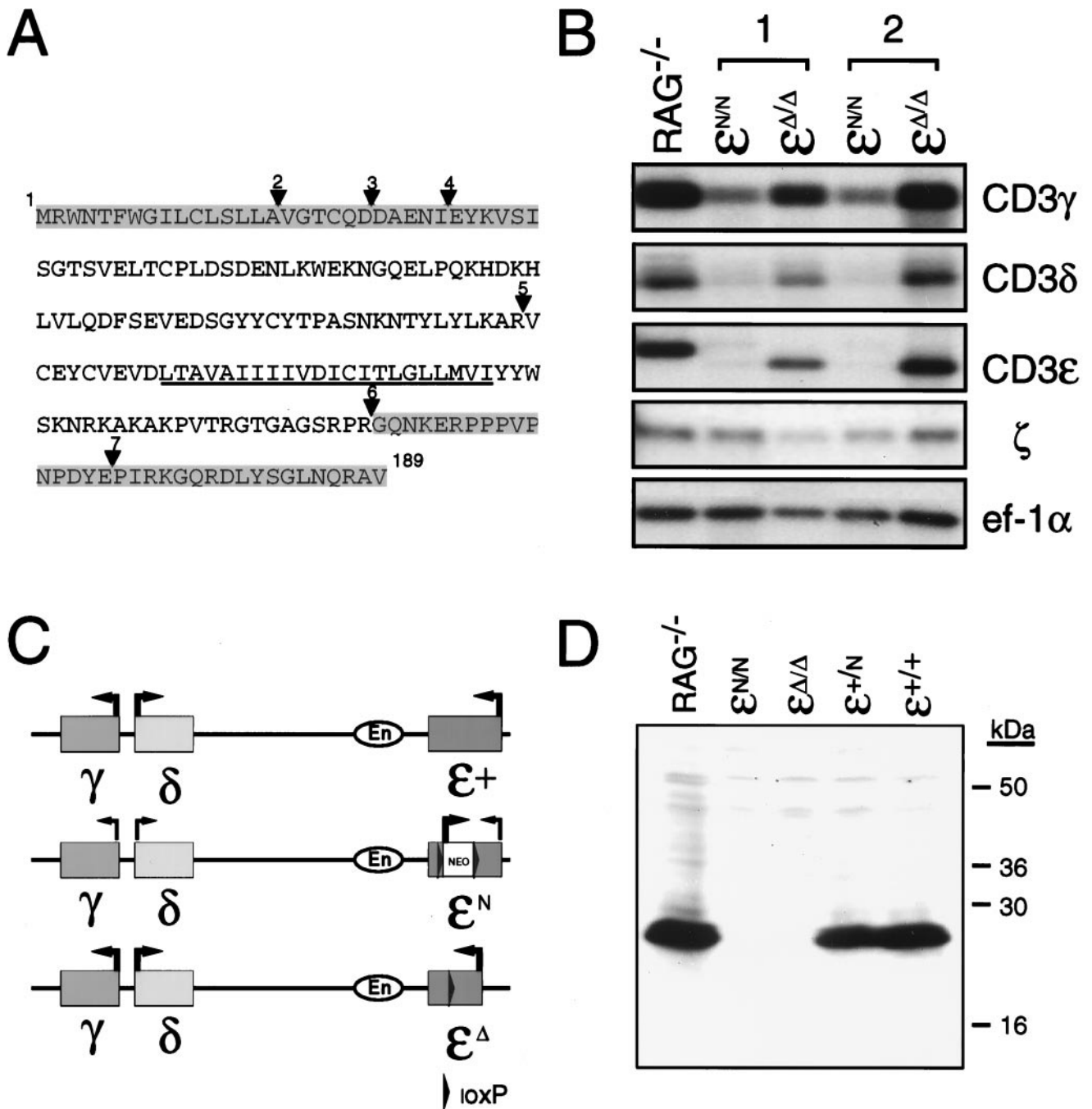


FIG. 2. (A) Predicted amino acid sequence of CD3 ϵ^{Δ} . Amino acid sequence of the fully processed CD3 ϵ protein is shown. The transmembrane domain is underlined. Triangles indicate the location of introns (2–7) within the CD3 ϵ gene. Shaded areas represent the protein that potentially could be expressed in CD3 $\epsilon^{\Delta/\Delta}$ mice based on the coding sequences that are retained within thymocyte transcripts from these mice. (B) Northern blot analysis of thymocyte total RNA from Rag $^{-/-}$ mice and from CD3 $\epsilon^{N/N}$ and CD3 $\epsilon^{\Delta/\Delta}$ mice that had been generated from two independently derived ES cell clones (1 and 2). Blots were hybridized with probes corresponding to CD3 ϵ , CD3 γ , CD3 δ , and ζ and then were hybridized with a probe corresponding to the ubiquitously expressed ef-1 α to assess the integrity and quantity of RNA in each lane. (C) Representation of the CD3- γ , - δ , - ϵ cluster on mouse chromosome 9. Location of the CD3 enhancer (En) element 3-prime of CD3 ϵ is shown. Arrows indicate the direction of gene transcription. Thin lines indicate reduced gene expression. Depicted are the relative transcription levels observed from the (CD3) ϵ^{+} , (CD3) ϵ^N , and (CD3) ϵ^{Δ} alleles. (D) Western blot of thymocyte protein from Rag $^{-/-}$, (CD3) $\epsilon^{N/N}$, (CD3) $\epsilon^{\Delta/\Delta}$, (CD3) $\epsilon^{N/N}$, and (CD3) $\epsilon^{+/+}$ mice. Extracts from 7×10^6 thymocytes were run under reducing conditions on a 15% SDS-polyacrylamide gel, were transferred to poly(vinylidene difluoride) membrane, and were analyzed for CD3 ϵ expression by using a rabbit antiserum raised against a peptide corresponding to amino acids within the C terminus of the CD3 ϵ . Positions of the molecular mass standards are indicated in kilodaltons.

RESULTS AND DISCUSSION

Generation of CD3 $\epsilon^{N/N}$ and CD3 $\epsilon^{\Delta/\Delta}$ Mice. The vector p ϵ NEO-loxP, which contains a PGK-NEO expression cassette flanked by loxP sites situated within a portion of the CD3 ϵ gene, was used to generate targeted mutations of CD3 ϵ in mouse ES cells (Fig. 1A). ES cell clones in which the targeting

construct had integrated by homologous recombination contained a mutated CD3 ϵ allele (ϵ^N) in which PGK-NEO was substituted for most of exon V and all of exon VI, which encode most of the extracellular domain and the entire transmembrane domain of CD3 ϵ , respectively (Figs. 1A and 2A). Two positive ES clones were used to generate chimeric mice, which subsequently transmitted the mutated allele to their offspring.

Because it previously had been reported that mutation of $CD3\epsilon$ alters the expression of the $CD3\gamma$ and $CD3\delta$ genes (12), thymocytes from mice homozygous for the $CD3\epsilon$ mutant allele ($CD3\epsilon^{N/N}$) were screened for expression of $CD3\gamma$ and $CD3\delta$. Indeed, $CD3\epsilon^{N/N}$ mice generated from both of the independently derived ES clones exhibited a phenotype identical to that previously described [i.e., a reduction in the level of $CD3\gamma$ transcripts and a near absence of $CD3\delta$ transcripts (ref. 12 and Fig. 2*B*)]. $CD3\epsilon$ transcripts, which were of aberrant size, were also nearly undetectable in $CD3\epsilon^{N/N}$ mice whereas expression of the unlinked ζ gene was unaffected (Fig. 2*B*).

It has been shown that the PGK-NEO cassette can negatively influence the expression of genes in its proximity, presumably by competing for regulatory sequences (19). Because the $CD3\gamma$, $CD3\delta$, and $CD3\epsilon$ genes are linked closely, residing within 50 kilobases on mouse chromosome 9 (20), and are thought to be coregulated by an enhancer element located directly 3' of $CD3\epsilon$ (ref. 21 and Fig. 2*C*), it was conceivable that such a mechanism could account for the phenotype of $CD3\epsilon^{N/N}$ mice. Consistent with this idea, multiple thymocyte-specific NEO transcripts were detectable in both $CD3\epsilon^{+/N}$ and $CD3\epsilon^{N/N}$ mice (data not shown). To ascertain whether the presence of PGK-NEO was in fact responsible for the decrease in $CD3\gamma$ and $CD3\delta$ expression, the *loxP*-PGK-NEO-*loxP* cassette was removed from $CD3\epsilon^{N/N}$ mice by Cre/*loxP*-mediated

recombination. $CD3\epsilon^{N/N}$ mice were mated with EIIa-*cre* transgenic mice, which express Cre recombinase in all cells at the preimplantation stage of embryogenesis (16). Offspring of these matings initially were screened for loss of PGK-NEO (designated $CD3\epsilon^{\Delta}$) by Southern blotting and subsequently were tracked by a competitive PCR reaction that scores for the reduction in length of a product generated by primers corresponding to sequences within exon V and exon VII (Fig. 1*B*).

Analysis of thymocyte mRNA from $CD3\epsilon^{\Delta/\Delta}$ mice by Northern blotting revealed a single abundant $CD3\epsilon$ transcript of reduced length compared with the wild-type $CD3\epsilon$ transcript (Fig. 2*B*). Sequencing of cDNAs generated from $CD3\epsilon^{\Delta}$ transcripts by RT-PCR demonstrated that these mRNAs contained the intact coding sequences from exons I-IV and VII-VIII of $CD3\epsilon$ and thus represented a precise deletion of the exon V and exon VI coding sequences (Fig. 2*A*). Because these transcripts could conceivably direct the synthesis of a mutant $CD3\epsilon$ protein, thymocyte extracts from $CD3\epsilon^{\Delta/\Delta}$ mice were screened by Western blotting with an antibody directed against a peptide corresponding to amino acids 156-168 of $CD3\epsilon$, which should be retained within the intact $CD3\epsilon^{\Delta}$ product (Fig. 2*A*). Immunoreactive protein was undetectable in $CD3\epsilon^{\Delta/\Delta}$ mice, indicating that either the protein encoded by the $CD3\epsilon^{\Delta}$ transcripts is highly unstable or that the epitope recognized by the antibody has been deleted (Fig. 2*D*). Sig-

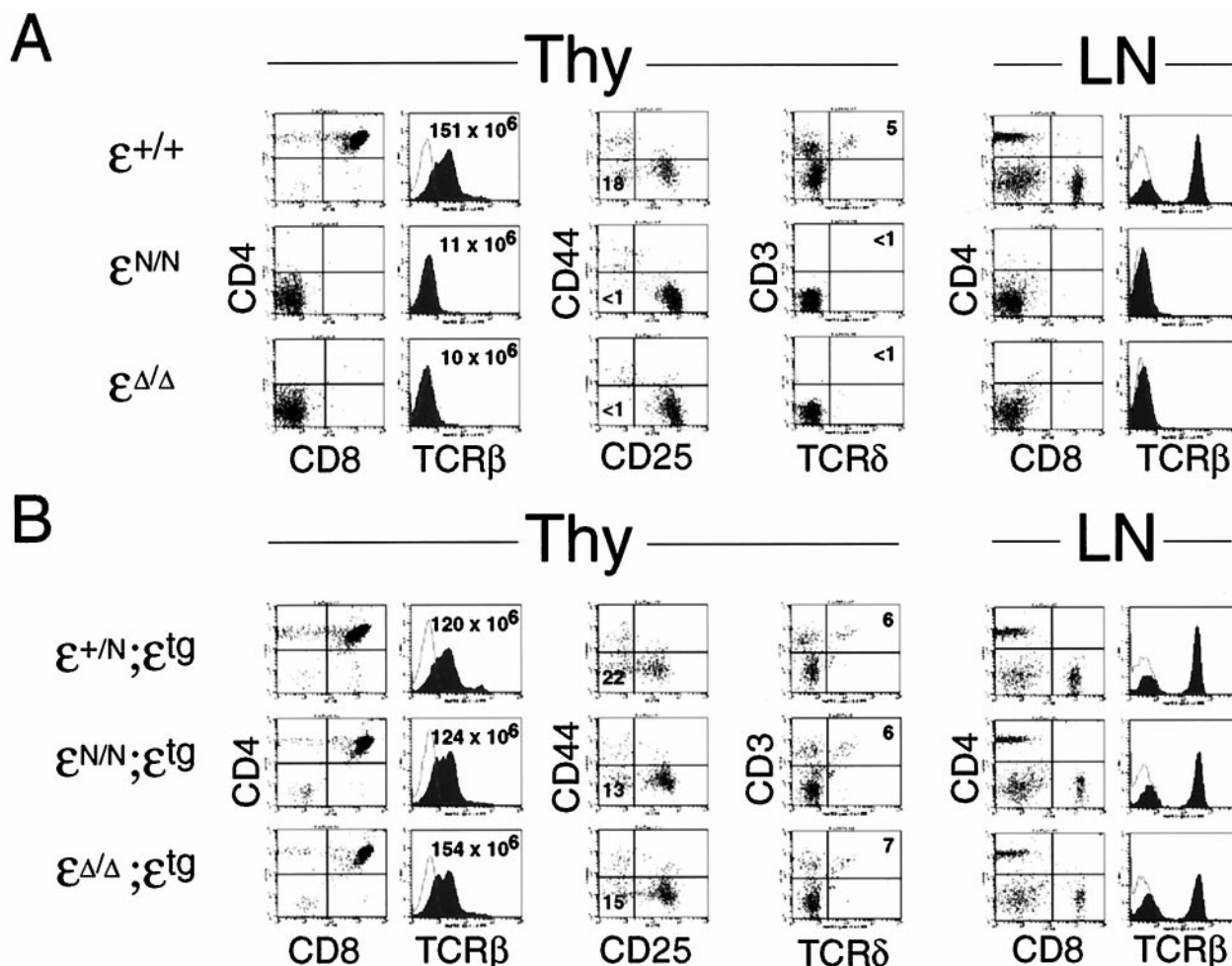


FIG. 3. (A) Phenotype of $CD3\epsilon^{N/N}$ and $CD3\epsilon^{\Delta/\Delta}$ mice. Thy, thymocytes; LN, lymph node. The left two panels show two-color (CD4 vs. CD8) and single-color (TCR β) analysis of thymocytes from ($CD3\epsilon^{+/+}$, ($CD3\epsilon^{N/N}$, and ($CD3\epsilon^{\Delta/\Delta}$) mice. Total thymocyte numbers from representative mice are provided in the two color plots. The central panels show two-color (CD44 vs. CD25) and two-color (CD3 vs. TCR δ) staining of gated (CD4⁻CD8⁻B220⁻) thymocytes. Numbers in quadrants indicate percentage of gated cells within the quadrant. The right two panels show two-color (CD4 vs. CD8) and single-color (TCR β) analysis of total lymph node cells. (B) Phenotype of $CD3\epsilon^{N/N};\epsilon^{tg}$ and $CD3\epsilon^{\Delta/\Delta};\epsilon^{tg}$ mice. The panels show representative data from ($CD3\epsilon^{+/N}$, ($CD3\epsilon^{N/N}$, and ($CD3\epsilon^{\Delta/\Delta}$) mice that contain a $CD3\epsilon$ transgene under the control of the human CD2 promoter and enhancer sequences (founder line C7530). Analysis is identical to that described in A.

nificantly, in contrast to $CD3\epsilon^{N/N}$ mice, thymocytes from $CD3\epsilon^{\Delta/\Delta}$ mice contained normal levels of $CD3\gamma$ and $CD3\delta$ transcripts, which were of the predicted size (Fig. 2B). Thus, the generation of $CD3\epsilon^{\Delta/\Delta}$ mice enabled a specific examination of the requirement for $CD3\epsilon$ in development.

Phenotype of $CD3\epsilon^{\Delta/\Delta}$ Mice. Analysis of T cell development in $CD3\epsilon^{\Delta/\Delta}$ mice revealed a phenotype virtually indistinguishable from that of $CD3\epsilon^{N/N}$ mice (ref. 12 and Fig. 3). Total thymocyte numbers in both $CD3\epsilon^{\Delta/\Delta}$ and $CD3\epsilon^{N/N}$ mice were $<10\%$ of control ($CD3\epsilon^{+/+}$), and the thymocytes that were present consisted entirely of the DN cells (i.e., both DP and $CD4^+CD8^-$ or $CD4^-CD8^+$ thymocytes were absent) (Fig. 3A). DN thymocytes progress through four stages of maturation before their transition to the DP stage: $CD44^+CD25^- \rightarrow CD44^-CD25^+ \rightarrow CD44^{-/lo}CD25^+ \rightarrow CD44^-CD25^-$ (22). Rearrangement of the $TCR\beta$ locus is first detected in $CD44^{-/lo}CD25^+$ thymocytes, and progression of these cells to the $CD44^-CD25^-$ stage is thought to require expression of the pre-TCR (or TCR) complex (2, 22). Consistent with this idea, thymocyte development is arrested at the $CD44^{-/lo}CD25^+$ stage in both $Rag^{-/-}$ and $CD3\epsilon^{N/N}$ mice (refs. 7, 8, and 12 and Fig. 3A). Significantly, thymocyte development was arrested at the identical ($CD44^{-/lo}CD25^+$) stage in $CD3\epsilon^{\Delta/\Delta}$ mice (Fig. 3A). As observed previously in $CD3\epsilon^{N/N}$ mice (12), $TCR\beta$, $TCR\gamma$, and $TCR\delta$ gene rearrangements were readily detectable in thymocytes from $CD3\epsilon^{\Delta/\Delta}$ mice (data not shown). Thus, these results demonstrate that $CD3\epsilon$ specifically is required for maturation of $CD44^{-/lo}CD25^+$ thymocytes to the $CD44^-CD25^-$ stage. In addition, the fact that T cell development is blocked at the identical stage in $CD3\epsilon^{N/N}$ and $CD3\epsilon^{\Delta/\Delta}$ mice indicates that $CD3\gamma$ and $CD3\delta$ are incapable of compensating for the loss of $CD3\epsilon$.

As predicted from the early block in thymocyte development, mature $CD4^+CD8^-$ or $CD4^-CD8^+$ T cells were absent from the periphery of $CD3\epsilon^{\Delta/\Delta}$ mice (Fig. 3A). Moreover, surface $\alpha\beta$ TCR and $\gamma\delta$ TCR complexes were undetectable in both the thymus and peripheral lymphoid organs as assessed by staining for $TCR\beta$ and $TCR\delta$ chains (Fig. 3A). As previously reported for $CD3\epsilon^{N/N}$ mice (12), the development of intestinal intraepithelial T cells also was arrested in $CD3\epsilon^{\Delta/\Delta}$ mice whereas B and NK cell development appeared unaffected (data not shown).

Expression of a $CD3\epsilon$ Transgene Rescues T Cell Development in $CD3\epsilon^{\Delta/\Delta}$ Mice. To confirm that the phenotype exhibited by $CD3\epsilon^{\Delta/\Delta}$ mice could be attributed entirely to the absence of $CD3\epsilon$, we reconstituted $CD3\epsilon$ expression in $CD3\epsilon^{\Delta/\Delta}$ mice by the introduction of a $CD3\epsilon$ transgene. Our own investigations have shown that transgenes placed under the control of the T-lineage-specific human $CD2$ promoter and enhancer begin to be expressed in thymocytes at the DN, $CD44^{-/lo}CD25^+$ stage and continue to be expressed at all subsequent stages of development (ref. 23 and data not shown). Thus, the $CD3\epsilon$ transgene construct should provide expression of $CD3\epsilon$ at the stage in development at which an arrest is first observed in $CD3\epsilon^{\Delta/\Delta}$ mice (Fig. 3A). Because overexpression of $CD3\epsilon$ has been shown to block T cell development (24), we used transgenic founder lines that express relatively low levels of $CD3\epsilon$ for reconstitution experiments (0.25- and $2\times$ wild-type). These transgenes did not inhibit thymocyte development when introduced into $CD3^{+/+}$ or $CD3^{N/N}$ mice (Fig. 3B and data not shown).

As depicted in Fig. 3B, reexpression of $CD3\epsilon$ in $CD3\epsilon^{\Delta/\Delta}$ mice effectively reversed the developmental defects as assessed by the presence of DN, $CD44^-CD25^-$ thymocytes, as well as DP and $CD4^+CD8^-$ and $CD4^-CD8^+$ thymocytes in $CD3\epsilon^{\Delta/\Delta}CD3\epsilon^{tg}$ mice. In addition, total thymocyte cellularity was increased markedly (to normal levels) in $CD3\epsilon^{\Delta/\Delta}CD3\epsilon^{tg}$ mice, and mature T cells were present in the periphery (Fig. 3). Finally, reexpression of $CD3\epsilon$ also restored surface expression of $\alpha\beta$ - and $\gamma\delta$ -TCR complexes on both immature and mature

T cells (Fig. 3A and B). Thus, these results demonstrate that the defects in T cell development observed in $CD3\epsilon^{\Delta/\Delta}$ mice are caused specifically by $CD3\epsilon$ deficiency. Moreover, because the $CD3\epsilon$ transgene would not be expected to restore expression of $CD3\epsilon$ at stages in development that precede the DN, $CD44^{-/lo}CD25^+$ stage, these findings lend support to the idea that $CD3\epsilon$ first is required at the point at which expression of the pre-TCR begins (12).

Of interest, introduction of the $CD3\epsilon$ transgene also fully corrected the developmental defect in $CD3\epsilon^{N/N}$ mice (Fig. 3B). This result was unexpected because DN thymocytes from $CD3\epsilon^{N/N}$ mice contain abnormally low levels of $CD3\gamma$ and $CD3\delta$ transcripts (ref. 12 and Fig. 2B), and thymocyte development is compromised in the absence of either $CD3\gamma$ or $CD3\delta$ (13, 14). Analysis of $CD3\gamma$ and $CD3\delta$ gene expression in total thymocytes from $CD3\epsilon^{N/N}CD3\epsilon^{tg}$ mice (which consisted mostly of DP cells) revealed normal levels of $CD3\gamma$ or $CD3\delta$ transcripts (Fig. 4). Thus, the restoration of thymocyte development in $CD3\epsilon^{N/N}CD3\epsilon^{tg}$ mice could be attributed to normalization of $CD3\gamma$ and $CD3\delta$ expression as well as reconstitution of $CD3\epsilon$. There are several possible explanations for these results. For example, the inhibitory effect of PGK-NEO mice may be variable between individual cells, such that some DN thymocytes express relatively normal levels of $CD3\gamma$ and $CD3\delta$ and these cells are expanded selectively when $CD3\epsilon$ is reexpressed. Alternatively, the transcriptional suppression of $CD3\gamma$ and $CD3\delta$ by PGK-NEO may be cell cycle- or stage-dependent. Regardless of the mechanism, these results demonstrate that the inhibitory effect of PGK-NEO on $CD3\gamma$ and $CD3\delta$ gene expression is not developmentally limiting if $CD3\epsilon$ expression is restored.

In summary, these results reveal a specific requirement for $CD3\epsilon$ in T cell development not exhibited by any of the other invariant subunits of the TCR complex (i.e., $CD3\gamma$, $CD3\delta$, and ζ) (11, 13, 14). The present data are consistent with previous results obtained in T cell lines demonstrating that $\alpha\beta$ TCR complexes are still expressed, though at reduced levels, in the absence of $CD3\gamma$, $CD3\delta$, or ζ but that $CD3\epsilon$ is absolutely required for surface expression of $TCR\alpha/\beta$ heterodimers (25–27). The absence of DP thymocytes and $\gamma\delta$ TCR⁺ cells in

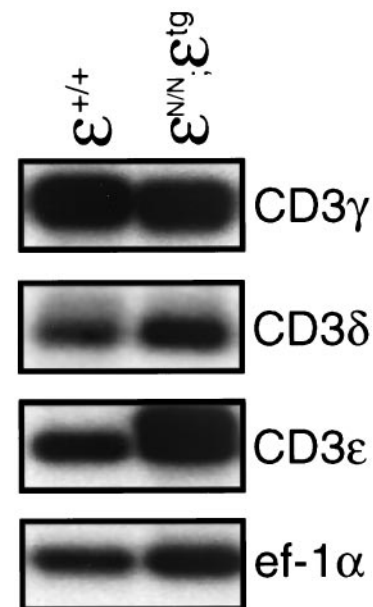


Fig. 4. Northern blot analysis of total thymocyte RNA from $CD3\epsilon^{+/+}$ and $CD3\epsilon^{N/N};\epsilon^{tg}$ mice. Blots were hybridized with probes corresponding to $CD3\epsilon$, $CD3\gamma$, $CD3\delta$, and ζ and then were hybridized with the probe corresponding to the ubiquitously expressed $ef-1\alpha$ to assess the integrity and quantity of RNA in each lane.

CD3 $\epsilon^{\Delta/\Delta}$ mice demonstrates that CD3 ϵ is also required for the expression and/or function of the pre-TCR and $\gamma\delta$ TCR complexes. That this need not necessarily have been the case is suggested by the discovery that the pre-TCR and $\gamma\delta$ TCR differ from the $\alpha\beta$ TCR in their requirement for CD3 δ (13, 28). Taken together, current data indicate that a minimal requirement for pre-TCR/TCR expression and/or function is the expression of either a CD3 $\gamma\epsilon$ or CD3 $\delta\epsilon$ heterodimer, although the former are much more efficient at promoting the assembly and surface expression of pre-TCR and $\gamma\delta$ TCR complexes. Finally, although these experiments demonstrate an obligate role for CD3 ϵ in T cell development, it is important to note that the defect in CD3 $\epsilon^{\Delta/\Delta}$ mice results in loss of both the structural and signal transducing functions of CD3 ϵ . Therefore, additional studies will be required to determine whether CD3 ϵ -mediated signal transduction is required at specific stages of development.

We thank B. J. Fowlkes for critical review of the manuscript. K.J.W. is a Howard Hughes Medical Institute–National Institutes of Health Research Scholar.

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