

Synergistic Effects of Interleukin-7 and Pre-T Cell Receptor Signaling in Human T Cell Development*

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Background: The role of IL-7 in human pre-T cell development is controversial.

Results: Using IL-7R α -modified Molt3, we demonstrate a synergistic effect of pre-TCR and IL-7 involving STAT5, Akt, and Erk1/2.

Conclusion: There is a stringent requirement for down-regulation of IL-7/IL-7R α signaling during human T cell development.

Significance: Understanding how IL-7 regulates human T cell development will help future development of T cell therapeutics.

The role of IL-7 in pre-T cell receptor (TCR) signaling during human T cell development is poorly understood. To study this, we engineered Molt3, a T cell progenitor T-acute lymphoblastic leukemia cell line, using lentiviral IL-7 receptor α (IL-7R α) to serve as a model system. IL-7 promoted pre-TCR activation in IL-7R α^{hi} Molt3 as illustrated by CD25 up-regulation after anti-CD3 stimulation. Anti-CD3 treatment activated Akt and Erk1/2 signaling pathways as proven using specific inhibitors, and IL-7 further enhanced both signaling pathways. The close association of IL-7R α with CD3 ζ in the pre-TCR complex was illustrated through live imaging confocal fluorescence microscopy. These results demonstrate a direct and cooperative role of IL-7 in pre-TCR signaling.

Maturation of T cell receptor (TCR)² $\alpha\beta$ or $\gamma\delta$ T cells from hematopoietic progenitor cells is a progressive and multistep process. T cell development has been characterized by the sequential expression of CD4 and CD8 coreceptors. The CD8⁻CD4⁻ double-negative (DN) thymocytes differentiate through the CD8 immature single-positive (SP) stage in mice and the CD4 immature SP stage in humans, followed by CD8⁺CD4⁺ double-positive (DP) cells and then CD4⁺ and CD8⁺ SP cells (1–4). IL-7, pre-TCR, and Notch signaling plays a crucial role in early T cell development. The transition from DN to DP is regulated by pre-TCR (5, 6) and Notch (7–11) signaling. IL-7 supports DN cell survival and proliferation; however, the role of IL-7 during DN-to-DP transition is not well characterized (12–17).

IL-7 is thought to play a cooperative role in pre-TCR signaling (18, 19). However, there are contradicting reports of both

inhibitory and supportive roles of IL-7 during DN-to-DP transition. An IL-7 dose-dependent inhibition of DN-to-DP transition *in vivo* has been reported (20), and IL-7 blocks differentiation at the DN3 stage in fetal thymus organ culture (21). In addition, IL-7 suppresses anti-CD3 antibody-mediated DN-to-DP transition in RAG-1 (recombination-activating gene-1)-deficient pre-T cells (22), and furthermore, IL-7 inhibits DN-to-DP transition in fetal thymus organ culture derived from IL-7 receptor α (IL-7R α) transgenic mice by inhibiting transcription of T cell factor, lymphoid enhancer factor, and retinoic acid-related orphan receptor γ (23). As pre-TCR signaling is central in DN-to-DP transition (24, 25), the above studies suggest that IL-7 signaling inhibits DN-to-DP transition by inhibiting pre-TCR signaling. On the other hand, IL-7R $\alpha^{-/-}$ mice display $\gamma\delta$ T cell deficiency and a reduced amount of $\alpha\beta$ T cells (26, 27). In humans, mutation in IL-7R α is associated with immunodeficiency (28). Anti-IL-7R α antibody inhibits DP transition and TCR $\alpha\beta$ expression in human CD34 cells in fetal thymus organ culture (29, 30). The latter studies suggest that IL-7/IL-7R α signaling plays a supportive role in DN-to-DP transition, which contradicts some other studies.

Insights into direct interaction of IL-7 and pre-TCR signaling pathways could help resolve the controversy. During T cell development, pre-TCR signals in an autonomous fashion independent of extracellular ligands (5, 31, 32); however, anti-CD3 antibody can activate pre-TCR signaling (33, 34). A TCR $\alpha\beta$ -deficient Jurkat cell line has been used to study pre-TCR signaling (35–37). However, Jurkat cells are considered to be mature T cells and do not recapitulate T cell precursor activities during pre-T cell development. Therefore, we evaluated different T cell progenitor T-acute lymphoblastic leukemia (T-ALL) cell lines and engineered an IL-7R α -expressing Molt3 cell line as a model pre-T cell system. Our results indicate that IL-7 enhances pre-TCR signaling through Erk1/2 and Akt pathways. This is consistent with the report that receptors with cytoplasmic domain-associated kinases aggregate with TCR and promote TCR signaling. On the other hand, the receptors with phosphatase-associated cytoplasmic domains move away from TCR (38). Further examination of the membrane compartmen-

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² The abbreviations used are: TCR, T cell receptor; DN, double-negative; SP, single-positive; DP, double-positive; IL-7R α , IL-7 receptor α ; T-ALL, T-acute lymphoblastic leukemia; eGFP, enhanced GFP.

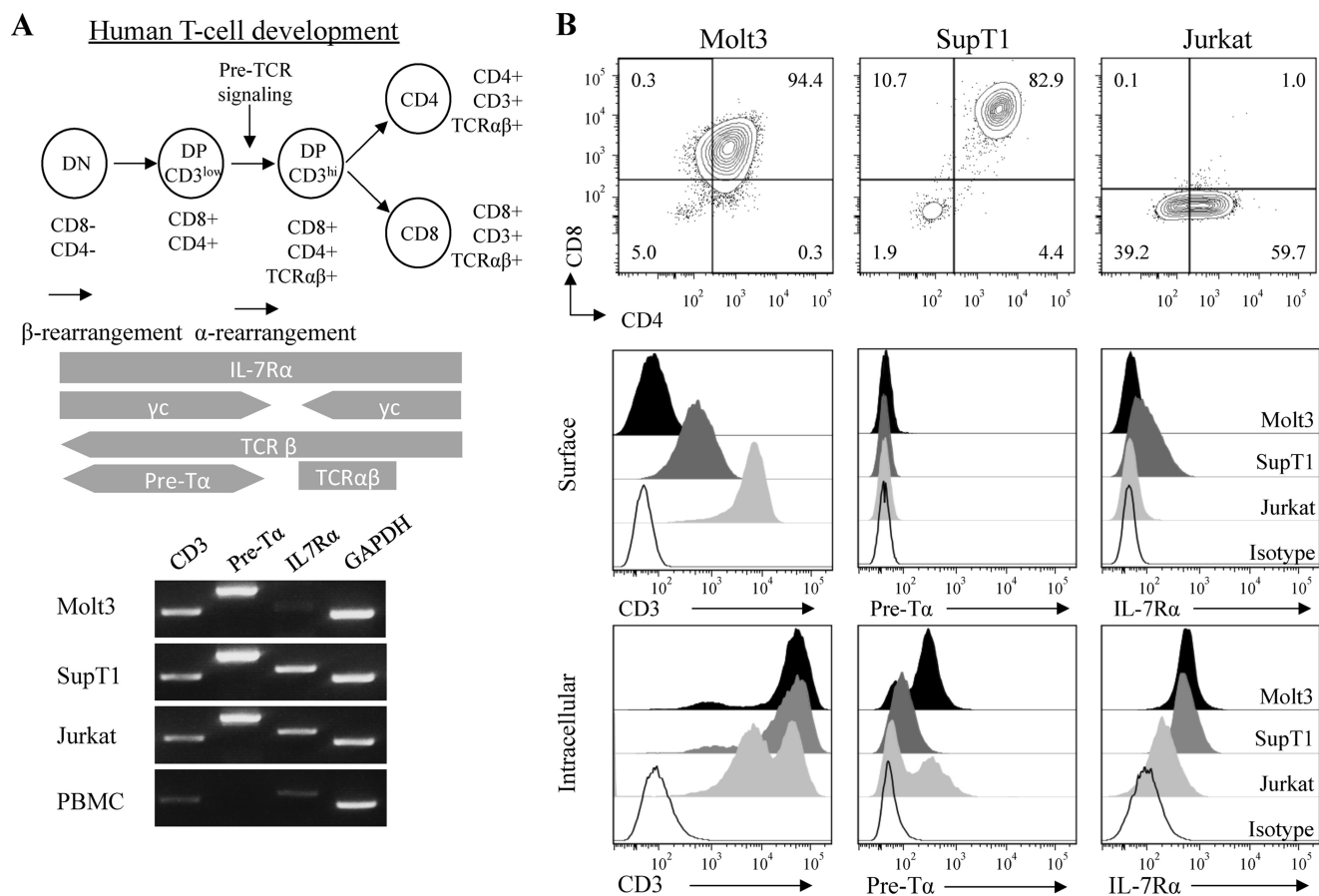


FIGURE 1. Analysis of pre-T signaling receptor molecules in Molt3, SupT1, and Jurkat T-ALL cell lines. A, schematic illustration of human T cell development. Results from the RT-PCR analysis of transcripts of CD3 ϵ , pre-T α , and IL-7R α using GAPDH as a control are also shown. PBMC, peripheral blood mononuclear cells. B, flow cytometry analysis of surface expression of CD8, CD4, CD3, pre-T α , and IL-7R α and intracellular expression of CD3, pre-T α , and IL-7R α .

talization of IL-7R α and CD3 ζ in live cells using confocal microscopy indicated that IL-7R α co-localizes with CD3 ζ , confirming a cooperative role of IL-7R α in pre-TCR signaling.

EXPERIMENTAL PROCEDURES

Flow Cytometry and Antibodies—The antibodies used for surface and intracellular staining were as follows: phycoerythrin-Cy7-conjugated anti-CD3 (clone SK7), Pacific Blue-conjugated anti-CD4 (clone RPA-T4), allophycocyanin-Cy7-conjugated anti-CD8 (clone SK1), FITC-conjugated anti-phospho-Tyr-694 STAT5 (clone 47), phycoerythrin-conjugated anti-phospho-Ser-473 Akt (clone M89-61), and Alexa Fluor 647-conjugated anti-phospho-Thr-202/phospho-Tyr-204 Erk1/2 (clone 20A) (BD Biosciences); anti-CD127 (clone 40131; R&D Systems, Minneapolis, MN); and anti-pre-T α (G-14; Santa Cruz Biotechnology). For flow cytometry staining, cells were first washed with PBS plus 2% FBS and blocked with mouse and human serum at 4 °C for 30 min. Cells were incubated with antibodies following the manufacturers' instructions. For each fluorochrome-labeled antibody used, an appropriate isotype control was included. After antibody staining, the cells were washed twice with FACS buffer and fixed with 2% paraformaldehyde. Intracellular staining was performed using a BD Cytotfix/Cytoperm kit (BD Biosciences) according to the manufacturer's protocol. For detection of phosphorylated proteins, cells

were stimulated using the indicated stimulus for 15 min. The cells were then fixed by adding an equal amount of pre-warmed 2% paraformaldehyde and permeabilized with 90% methanol for 30 min on ice or at -20 °C overnight. Next, the cells were washed and incubated with the indicated antibodies for 1 h at room temperature, followed by two washes with FACS buffer. Data were acquired using BD FACSDiva software (Version 5.0.1) on a BD LSR II flow cytometer and analyzed using FlowJo software (Version 7.1.3.0, Tree Star Inc., Pasadena, TX).

Lentiviral Vector Construction and Transduction—Lentiviral vectors were generated using the NHP/TYF lentiviral vector system as described previously (39, 40). IL-7R α cDNA was cloned into the pTYF transducing vector behind the human EF1 α promoter.

Inhibitors—Jak3 inhibitor V was purchased from Calbiochem. MEK1/2 inhibitor U0126 and PI3K inhibitor LY294002 were purchased from Cell Signaling Technology, Inc. (Danvers, MA).

RNA Extraction and Semiquantitative RT-PCR—RNA was harvested from cells using TRI Reagent (Sigma-Aldrich). 1 μ g of RNA was reverse-transcribed into cDNA using a two-step avian myeloblastosis virus RT-PCR kit (GeneChoice, Frederick, MD). The following primers were used for PCR: GAPDH,

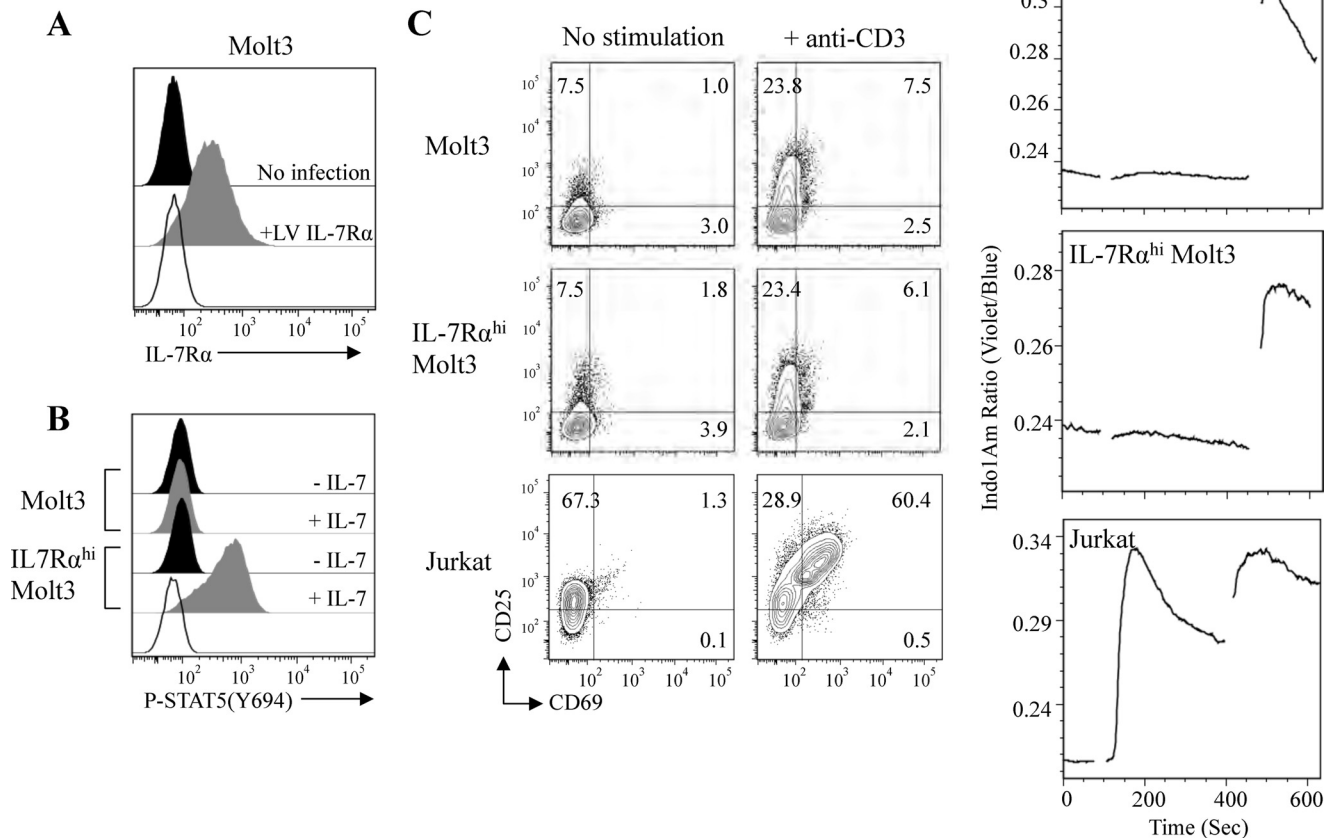


FIGURE 2. Analysis of IL-7 and anti-CD3 signaling in lentiviral vector-modified IL-7R α^{hi} Molt3. A, surface expression of IL-7R α on Molt3 (no infection) and lentiviral vector-modified IL-7R α^{hi} Molt3 (+LV IL-7R α). B, STAT5 activation in response to 100 ng/ml IL-7 by Phosflow antibody staining and flow cytometry analysis. C, surface expression of activation markers CD25 and CD69 in Molt3, IL-7R α^{hi} Molt3, and Jurkat cells after overnight incubation with anti-CD3/CD28 beads. D, intracellular free calcium measurement in Indo-1/AM-loaded Molt3, IL-7R α^{hi} Molt3, and Jurkat cells in response to anti-CD3 antibody (clone HIT3a) and ionomycin.

5'-CCG ATG GCA AAT TCG ATG GC-3' (forward) and 5'-GAT GAC CCT TTT GGC TCC CC-3' (reverse); pre-T α , 5' AGT ACA CAG CCC ATG CAT CTG TCA-3' (forward) and 5'-AAT GCT CCA AGA CTG GAG GAA GGA-3' (reverse); CD3 ϵ , 5'-TGA AGC ATC ATC AGT AGT CAC AC-3' (forward) and 5'-GGC CTC TGT CAA CAT TTA CC-3' (reverse); and IL-7R α , 5'-TCG CAG CAC TCA CTG ACC-3' (forward) and 5'-GTC ATT GGC TCC TTC CCG-3' (reverse). After 30 cycles of amplification (95 °C for 30 s, 55 °C for 30 s, and 72 °C for 60 s), PCR products were separated on a 2% agarose gel.

Intracellular Free Ca²⁺ Measurements—Molt3 and Jurkat cells were loaded with the Ca²⁺ indicator dye Indo-1/AM (Molecular Probes) and stimulated with HIT3a (5 $\mu\text{g/ml}$), anti-mouse IgG (0.15 mg/ml), or ionomycin (1 $\mu\text{g/ml}$). A 1-ml volume of cells at a density of 2×10^6 cells/ml was warmed to 37 °C and placed onto the BD LSR II flow cytometer. Base-line measurements were collected for 1 min, anti-CD3 antibody and anti-mouse IgG were added, and measurement was continued for an additional 5 min, followed by the addition of ionomycin and measurement for an additional 1–2 min. Stimulus-induced changes in the intracellular Ca²⁺ concentration were deter-

mined over time by monitoring the fluorescence emission ratio of the Ca²⁺-bound *versus* free form of Indo-1 at 405 and 495 nm, respectively; data were analyzed and plotted using FlowJo software.

Western Blotting—Antibodies for phospho-Erk1/2 (Thr-202/Tyr-204; D13.14.4E and 137F5) were obtained from Cell Signaling Technology, Inc. For Western blot analysis, cells were collected at specific time points post-treatment and washed with cold PBS before being lysed in Blue loading buffer with DTT (Cell Signaling Technology, Inc.) and supplemented with protease inhibitor mixture (Roche Diagnostics). Protein was separated on a 10% gel and then transferred to PVDF membrane (Millipore). The membrane was blocked and probed with antibodies as indicated and then incubated with peroxidase-conjugated secondary antibody (Cell Signaling Technology, Inc.). Bound proteins were visualized by SuperSignal West Pico chemiluminescent substrate (Thermo Scientific, Waltham, MA).

Statistical Analysis—Statistical analysis was performed by applying the Wilcoxon matched-pairs signed-rank test using GraphPad Prism 5 software.

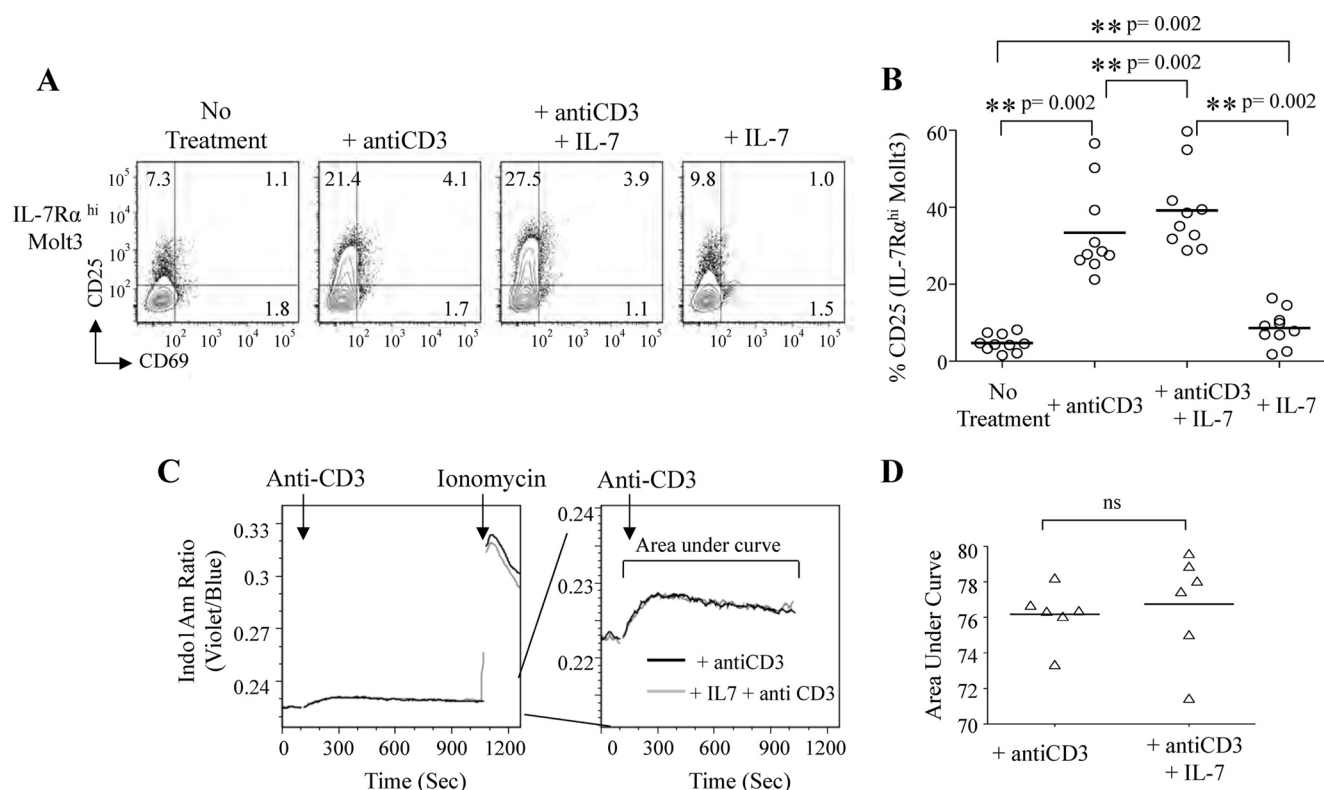


FIGURE 3. IL-7 enhances anti-CD3 antibody-induced pre-TCR signaling but not calcium flux induction in IL-7R α^{hi} Molt3. *A*, flow cytometry analysis of activation markers CD25 and CD69 in IL-7R α^{hi} Molt3 after overnight treatment as indicated. *B*, statistical analysis using the Wilcoxon matched-pairs signed-rank test ($n = 10$). *C*, intracellular free calcium measurement in Indo-1/AM-loaded IL-7R α^{hi} Molt3 in response to anti-CD3 antibody (clone HIT3a) and IL-7 using ionomycin as a control. *D*, the area under curve was quantified using FlowJo software, and data were analyzed using GraphPad Prism 5 ($n = 6$). *ns*, not significant.

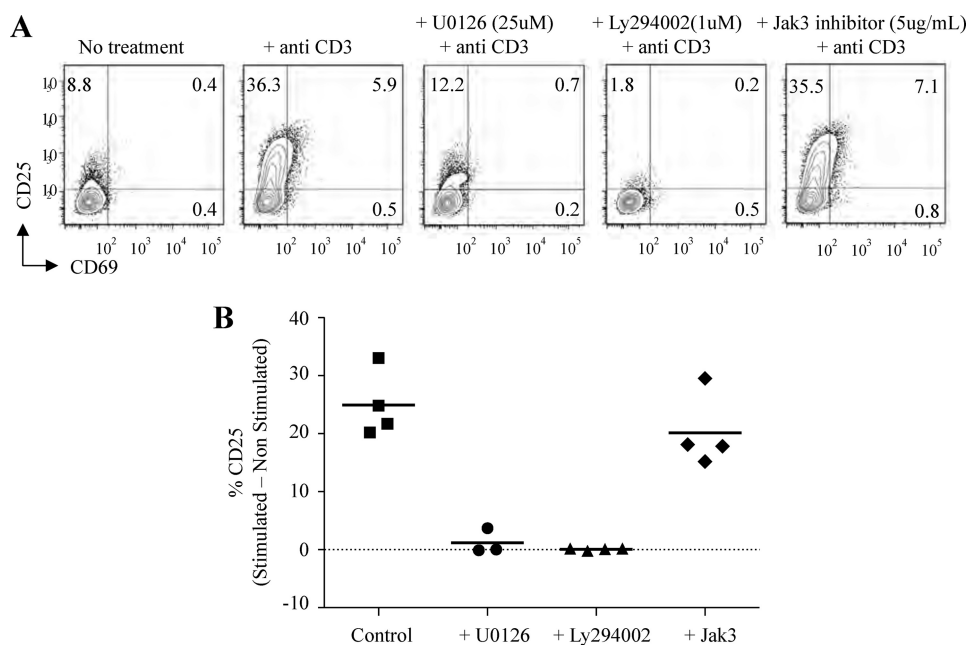


FIGURE 4. Analysis of pre-TCR signaling pathways in IL-7R α^{hi} Molt3 using specific kinase inhibitors. *A*, analysis of pre-TCR activation markers in IL-7R α^{hi} Molt3 after overnight activation and inhibitor treatment as indicated. *B*, percentage of CD25 expression in inhibitor-treated versus control anti-CD3 antibody-stimulated cells after subtraction of the non-stimulated control. Data represent at least three independent experiments.

RESULTS

IL-7R Expression and Pre-TCR Signaling in T-ALL Cell Lines— Key stages of human T cell development have been characterized as DN, CD3^{low} DP, CD3^{hi} TCR $\alpha\beta^+$ DP, CD8 SP, or CD4 SP

as shown at top of Fig. 1. Full-length transcripts of TCR β and pre-T α have been detected in the CD3^{low} DP stage, and pre-TCR signaling drives progression to CD3^{hi} DP and induces allelic exclusion at the TCR β locus, followed by TCR α locus

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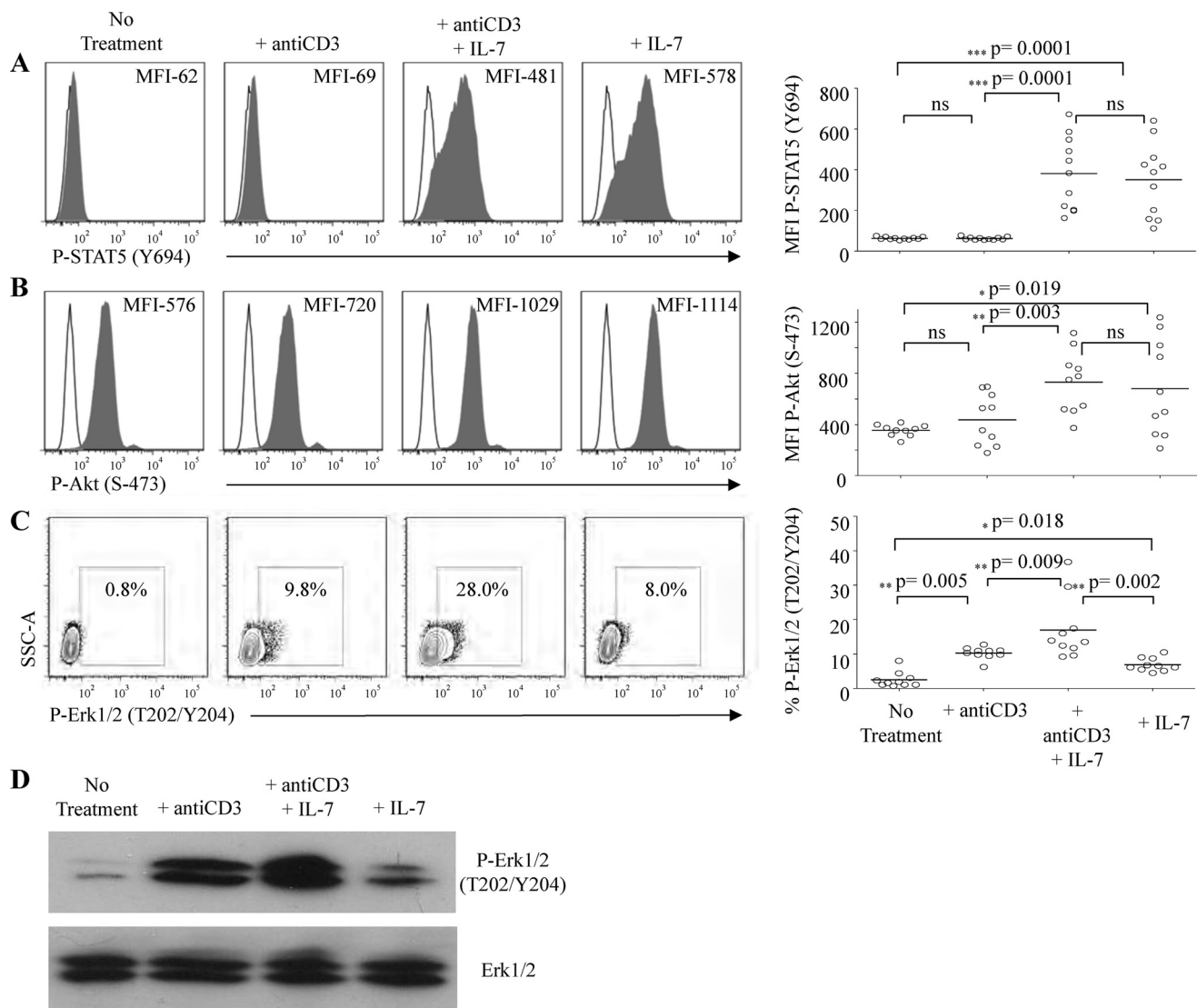


FIGURE 5. Synergy of IL-7 with pre-TCR signaling through activating Akt and Erk1/2. IL-7 α^{hi} Molt3 cells treated with 100 ng/ml IL-7 and anti-CD3/CD28 beads for 15 min. Cells were fixed; permeabilized; stained with antibodies against activated STAT5 (A), Akt (B), and Erk1/2 (C); and subjected to flow cytometry analysis. The mean fluorescence index (MFI) for STAT5 and Akt and the percentage for Erk1/2 are shown and are summarized on the right via the Wilcoxon matched-pairs signed-rank test ($n = 10$). D, Western analysis of Erk1/2 activation in response to the indicated treatment. ns, not significant.

rearrangement (41). We have recently established that concomitant IL-7/IL-7 α signaling and pre-TCR activation play a central role in promoting DN-to-DP transition as well as T cell maturation during human T cell development *in vitro* (73). To investigate the direct effect of IL-7 on pre-TCR signaling, we examined several T-ALL cell lines, including Jurkat, SupT1, and Molt3.

In mice, pre-TCR signaling is initiated at the DN3 stage in the absence of coreceptors. In humans, pre-TCR signaling is initiated at the CD4 immature SP and early DP stages of development. Molt3 resembles an early stage of DP T cells characterized by low CD3 and low CD4/CD8 coreceptor expression; SupT1 resembles a later stage of DP T cell development with high CD3 and coreceptor expression. As Molt3 has a rearranged TCR β chain and is positive for intracellular TCR β expression (42, 43), we further evaluated the intracellular and extracellular expression of CD3, pre-T α , and IL-7 α .

RT-PCR analysis of RNA showed comparable levels of CD3 and pre-T α but reduced IL-7 α transcripts in Molt3 compared

with SupT1 and Jurkat cells (Fig. 1A). Analysis of surface markers showed that Molt3 is CD4 $^{\text{low}}$ CD8 $^{\text{low}}$ DP with low surface CD3 expression (Fig. 1B, upper panels), but intracellular staining revealed that cytoplasmic CD3 expression in Molt3 was similar to that in SupT1 and Jurkat cells (Fig. 1B, lower panels). Intracellular staining of IL-7 α and pre-T α showed comparable expression of IL-7 α but higher pre-T α in Molt3 compared with SupT1 and Jurkat cells (Fig. 1B). In addition, Molt3 showed CD7 $^+$ CD1a $^{\text{low}}$ expression (data not shown). Together, these results suggest that Molt3 is arrested in the CD3 $^{\text{low}}$ DP pre-T cell development stage, during which pre-TCR signaling occurs (44). However, it lacks surface IL-7 α expression, a critical requirement for IL-7 signaling (45–47).

Ectopic Up-regulation of IL-7 α in Molt3 and Response to IL-7 and Anti-CD3 Stimulation—SupT1 expresses high levels of CD3 and IL-7 α and would be an ideal choice to study the effect of IL-7 on pre-TCR signaling. However, we found that SupT1 failed to respond to both IL-7 and anti-CD3 stimulation

as assessed by the lack of STAT5 activation, lacked up-regulation of T cell activation markers CD25 and CD69, and did not show induction of calcium flux. Hence, we constructed a lentiviral vector encoding IL-7R α and established an IL-7R α^{hi} Molt3 cell line. Flow cytometry analysis confirmed surface IL-7R α expression in the lentiviral IL-7R α -transduced Molt3 cell line (Fig. 2A, +LV IL-7R α). IL-7 responsiveness was then assessed by the induction of STAT5 phosphorylation; the result illustrated that IL-7 induced STAT5 (Tyr-694) activation in IL-7R α^{hi} Molt3 but not in unmodified Molt3 (Fig. 2B). We further evaluated the response to anti-CD3 stimulation by the analysis of T cell activation markers CD25 and CD69 (48). Whereas anti-CD3 antibody activated CD25 and CD69 in Jurkat cells, both the unmodified and IL-7R α^{hi} Molt3 cell lines up-regulated CD25, but only induced low levels of CD69 in response to anti-CD3 stimulation (Fig. 2C). TCR stimulation directly corresponds to calcium flux, which can be measured by Indo-1/AM violet/blue ratio. Thus, we evaluated induction of calcium flux, and the results showed a low level of response to anti-CD3 stimulation of both unmodified and IL-7R α^{hi} Molt3 compared with control ionomycin treatment or anti-CD3 antibody-stimulated Jurkat cells (Fig. 2D). Therefore, the anti-CD3 signaling consequence differed between the early T cells (Molt3) and the mature T cells (Jurkat).

Effect of IL-7 on Anti-CD3 Antibody-induced Pre-TCR Signaling—To more precisely assess the effect of IL-7 on anti-CD3 antibody-mediated pre-TCR activation, we tested four conditions (no treatment, anti-CD3 Ab, anti-CD3 Ab plus IL-7, and IL-7) using IL-7R α^{hi} Molt3 cells (Fig. 3A). The results showed a consistent increase ($n = 10$, $p = 0.002$) in the expression of CD25 after anti-CD3 Ab plus IL-7 treatment (Fig. 3B). Interestingly, IL-7 alone also induced a slight increase in CD25 ($p = 0.002$) (Fig. 3, A and B). The intensity and duration of TCR stimulation directly correspond to calcium flux, which can be measured by Indo-1/AM violet/blue ratio over time. Evaluation of the effect of IL-7 on anti-CD3 antibody-induced calcium flux showed no change in both the intensity and duration of calcium flux (Fig. 3, C and D).

Identification of Multiple Pre-TCR Signaling Pathways—Pre-TCR signaling drives DN-to-DP transition, but the signaling partners of the cytoplasmic tail of the human pre-TCR complex have not yet been identified (6). Pre-TCR signaling occurs through pathways reminiscent of signaling after TCR ligation in mature T cells based on knock-out and transgenic mouse studies (6, 31, 49–53). However, critical differences exist between the mouse and human cytoplasmic domains of the pre-T α chain (5, 25, 54, 55).

There are three potential pathways for human pre-TCR signaling: Erk1/2-, PI3K-, and γ_c -mediated signaling pathways. To understand how IL-7 enhances pre-TCR signaling, we first examined the candidate pathways involved in pre-TCR signaling in IL-7R α^{hi} Molt3. For the PI3K pathway, we tested the inhibitor LY294002. Erk1/2 signaling pathway activation is important for DN-to-DP transition, so we tested the inhibitor U0126 (56). To evaluate the γ_c signaling pathway, we utilized a Jak3 inhibitor, as γ_c chain signal transduction begins with the binding of Jak3 (57–59). IL-7R α^{hi} Molt3 was pretreated with the indicated inhibitors for 2 h, followed by anti-CD3 stimula-

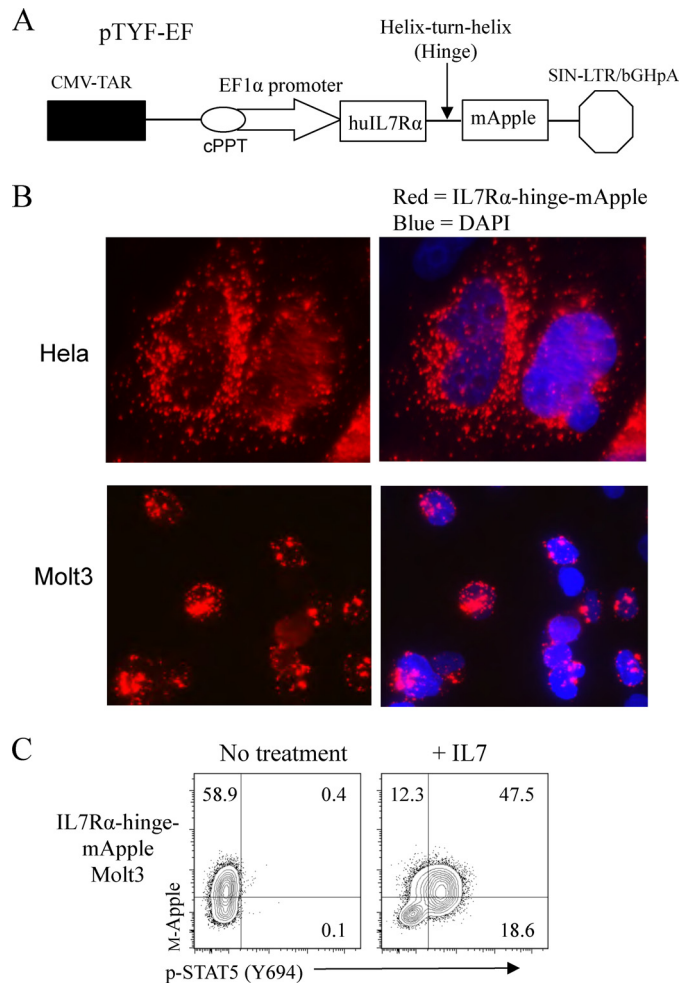


FIGURE 6. Live imaging and functional analysis of an IL-7R α -red fluorescent protein fusion. A, functional IL-7R α -mApple lentiviral vector construct with a helix-turn-helix hinge domain. CMV-TAR, cytomegalovirus-TAT transactivation response element; cPPT, central polyuridine tract; hu, human; SIN-LTR/bGHpA, self-inactivating long terminal repeat/bovine growth hormone polyadenylation signal. B, fluorescence images of HeLa and Molt3 cells infected with IL-7R α -hinge-mApple. C, STAT5 activation in response to IL-7 in Molt3 cells infected with lentiviral IL-7R α -hinge-mApple.

tion and overnight incubation. Analysis of CD25 and CD69 by surface staining and flow cytometry showed that the Erk1/2 and PI3K-Akt pathways were required for pre-TCR signaling, yet the γ_c signaling pathway was dispensable (Fig. 4, A and B). Similar results were obtained using unmodified Molt3 (data not shown).

Effect of IL-7 on Pre-TCR-mediated Erk1/2 and Akt Activation—To further characterize the effect of IL-7/IL-7R α signaling on the Erk1/2 and PI3K-Akt pathways in pre-TCR signaling, we treated IL-7R α^{hi} Molt3 with IL-7 and anti-CD3 Ab. Activation of STAT5 served as a control for IL-7 signaling. We used phosphorylation-specific antibodies and flow cytometry analysis for quantitative assessment of STAT5, Akt, and Erk1/2 activation. Preliminary flow cytometry analyses of these target proteins at 5, 15, 30, and 60 min post-stimulation established that all three proteins were optimally phosphorylated at 15 min, and thus, we chose this time point for further experiments. The results showed no STAT5 activation in response to anti-CD3 stimulation, but a large population of cells exhibited phospho-

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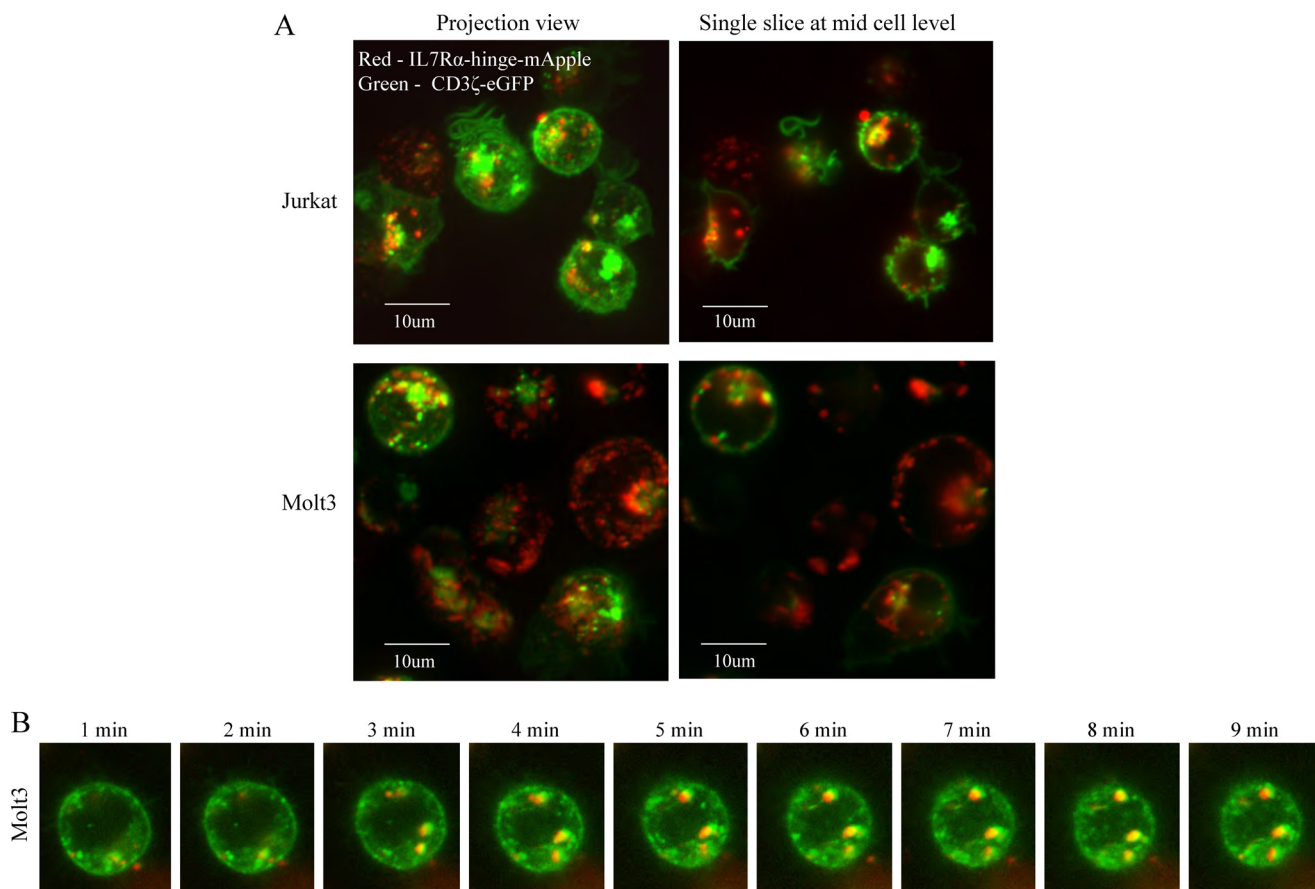


FIGURE 7. **Dynamic intracellular interaction of IL-7R α and CD3 ζ .** *A*, confocal images of Molt3 and Jurkat cells infected with lentiviral IL-7R α -hinge-mApple and CD3 ζ -eGFP. *B*, time-lapse confocal fluorescence images of Molt3 cells co-infected with lentiviral IL-7R α -hinge-mApple and CD3 ζ -eGFP.

STAT5 in response to IL-7 treatment (Fig. 5A). Molt3 is a γ -secretase inhibitor-resistant cell line and has PTEN deficiency and therefore has constitutively active Akt (60). We found that pre-TCR signaling enhanced Akt activation on some occasions, but the effect lacked statistical significance. On the other hand, IL-7 alone significantly increased Akt activation based on mean fluorescence index (Fig. 5B). Anti-CD3 stimulation activated Erk1/2 (from 0.8 to 9.8%) (Fig. 5C); interestingly, IL-7 alone was able to activate Erk1/2 (8%) and showed a synergistic effect upon anti-CD3 stimulation (28%, $n = 10$) (Fig. 5C), which was further confirmed by Western blotting (Fig. 5D).

Dynamic Interaction of IL-7R α and CD3 ζ —As IL-7 enhanced pre-TCR signaling, we further evaluated a possible direct interaction between IL-7R α and CD3 ζ , the latter being a central component of the pre-TCR signaling complex. We engineered an IL-7R α -mApple (monomeric red fluorescent protein) fusion to track IL-7R α in live cells. Lentiviral IL-7R α -mApple gene-transduced cells expressed a membrane form of the IL-7R α fusion but lacked STAT5 activation function in response to IL-7 (data not shown). To overcome this defect, we inserted a helix-turn-helix spacer between IL-7R α and mApple as illustrated in Fig. 6A. HeLa and Molt3 cells were transduced with the lentiviral IL-7R α -hinge-mApple fusion gene. Immunofluorescence detection showed abundant expression of the fusion protein in the perinuclear compartment (Fig. 6B). We treated IL-7R α -hinge-mApple Molt3 cells with IL-7 and examined

STAT5 activation through phospho-STAT5 intracellular Ab staining and flow cytometry. The result showed that IL-7 treatment markedly induced phospho-STAT5 in IL-7R α -hinge-mApple Molt3 cells (Fig. 6C). To evaluate the localization of IL-7R α with respect to CD3 ζ , a CD3 ζ -eGFP fusion gene (kindly provided by Dr. M. Krummel, University of California, San Francisco) was cloned into a lentiviral expression vector. The functional analysis of the CD3 ζ -eGFP fusion protein has been reported previously (61). Jurkat and Molt3 cells were cotransduced with lentiviral vectors encoding IL-7R α -hinge-mApple and CD3 ζ -eGFP, and the transduced cells were imaged live under a confocal microscope. We found that IL-7R α (red) and CD3 ζ (green) tended to co-localize (yellow) in microclusters formed intracellularly or on the cell surface (Fig. 7A). Time-lapse images illustrated that IL-7R α was moving in close proximity with CD3 ζ in Molt3 (Fig. 7B), suggesting a cooperative role of IL-7R α in pre-TCR signaling.

DISCUSSION

IL-7 inhibits pre-TCR signaling both *in vitro* and in transgenic mice (20–23). However, studies of *in vitro* human T cell development and patients with mutations in IL-7R α suggest an important role of IL-7 signaling in early T cell development (28, 29). Because of the complexity of these systems, the involvement of IL-7 in pre-TCR signaling has not been characterized. The goal of this study was to decipher the role of IL-7 in pre-TCR signaling. After extensive analysis of existing human

T-ALL cell lines, we established an IL-7 α -modified Molt3 cell line as a DP pre-T model system and demonstrated a synergistic role of IL-7 in pre-TCR signaling. Additionally, we reported that, in addition to STAT5 and Akt, IL-7 activated the Erk1/2 pathway in IL-7 α^{hi} Molt3. It has been shown that the cytoplasmic tail of IL-7 α is associated with Lck, the kinase normally associated with T cell coreceptors CD4 and CD8. Thus, the observation that IL-7 suppresses DP transition could be a result of indirect transcriptional regulation of the coreceptor CD4 gene rather than a direct effect on pre-TCR signaling; in fact, IL-7 is known to enhance the expression and activity of CD4 silencer-binding factors such as c-Myb (62).

There is discordance in the RNA and intracellular and surface protein expression of the CD3, pre-T α , and IL-7 α genes in the three T-ALL cell lines analyzed here. The levels of CD3 RNA and intracellular protein expression in Molt3 were comparable with those in SupT1 and Jurkat cells, but surface CD3 expression was markedly different (highest in Jurkat and lowest in Molt3 cells), suggesting possible post-translational regulation of surface CD3 expression. Molt3 was previously characterized as CD3-negative; however, this could be an artifact of the anti-CD3 antibody clones used (63). Many studies use antibody clones specifically recognizing CD3 ϵ paired with CD3 δ or CD3 γ ; the anti-CD3 antibody clone used in our study does not require such pairing. Moreover, pre-TCR does not require CD3 δ for its function (64). Unlike murine pre-T α , human pre-T α has an endoplasmic reticulum retention signal in its cytoplasmic domain and is strongly associated with the CD3 ζ chain (54, 55). We detected abundant pre-T α RNA expression in all three T-ALL cell lines, yet surface pre-T α was barely detected, and only Molt3 showed increased intracellular pre-T α expression. Interestingly, IL-7 α transcripts were present at extremely low levels in Molt3 compared with SupT1 and Jurkat cells, but the levels of intracellular protein expression were comparable. However, surface IL-7 α expression was detected only in SupT1. Thus, the modified Molt3 cell line expressing high surface IL-7 α (IL-7 α^{hi} Molt3) represents a rational model system for the study of pre-TCR and IL-7/IL-7 α signaling in pre-T cells.

IL-7 has been shown to activate the Erk1/2 pathway in pre-B cells but not in mature T cells (65, 66). Activation of the PI3K and Erk1/2 pathways in pre-T cells in response to IL-7 has not been previously evaluated (67). We found that IL-7 activated Akt and STAT5, as well as Erk1/2, which is in line with the finding that IL-7 activates Erk1/2 in an IL-7-dependent immature T cell line, TAIL-7 (68, 69). Unlike STAT5 and Akt activation, both of which showed substantial increases upon IL-7 treatment, IL-7 activated Erk1/2 only in a small percentage of cells (~8%), which correlated well with the marginal increase in CD25 expression in response to IL-7. Our findings suggest that IL-7 activates Erk1/2 in pre-T cells through a separate pathway in parallel to the canonical γ_c -dependent Jak-STAT activation pathway.

During mouse T cell development, IL-7 α is expressed in DN cells, shut down in DP cells, and then re-expressed in mature T cells (70, 71). Different from mouse thymocytes, IL-7 α is expressed in all stages of developing T cells in humans, but its binding partner γ_c is down-regulated during

the CD3 $^{\text{low}}$ DP and CD3 $^{\text{hi}}$ DP stages, resulting in a loss of STAT5 activation (72). During human T cell development, we found that there is a stringent requirement for down-regulation of IL-7/IL-7 α signaling in pre-TCR signaling (73). Here, we reported that IL-7 delivers additional signals similar to pre-TCR signals in a T-ALL cell line. The lack of down-regulation of IL-7 may aberrantly allow the precursors to bypass TCR β selection and progress toward the next stage with an improperly rearranged TCR β chain. Our finding provides a plausible explanation for the necessary down-regulation of IL-7 signaling during the pre-TCR stage of T cell development.

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