TNF receptor associated factor 3 plays a key role in development and function of invariant natural killer T cells

Zuoan Yi, Laura L. Stunz, and Gail A. Bishop

Abbreviations used: α-galactosylceramide, α-GalCer; iNKT, invariant NK T; LMC, littermate control; MFI, mean fluorescence intensity; TRAF3, TNF receptor associated factor 3.

TCR signaling is a prerequisite for early stage development of invariant natural killer T (iNKT) cells, whereas IL-15 signaling is required for expansion and maturation at later stages. In this study, we show that TNF receptor associated factor 3 (TRAF3) plays a critical role in the transition between these two distinct signaling pathways and developmental stages. TRAF3−deficient iNKT cells in CD4creTRAF3fl/fl (T-TRAF3−/−) mice exhibit defective up-regulation of T-bet and CD122, two critical molecules for IL-15 signaling, and as a consequence, IL-15−mediated iNKT cell proliferation and survival are impaired. Consistently, development of iNKT cells in T-TRAF3−/− mice shows a major defect at developmental stages 2 and 3, but not stages 0 and 1. We further demonstrated that defective T-bet up-regulation occurring during the stage 1 to stage 2 transition results from reduced TCR signaling in TRAF3−/− iNKT cells. In addition, mature TRAF3−/− iNKT cells displayed defective cytokine responses upon TCR stimulation. Collectively, our results reveal that by modulating the relative strength of TCR signaling, TRAF3 is an important regulator of iNKT cell development and functions.

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However, how the transition between signals from early to later stages is finely regulated to promote iNKT cell development remains ill defined. In this study, we identify TRAF3 as an essential regulator of iNKT cell development and function by modulating events dependent on the strength of TCR signaling.

stage (Borowski and Bendelac, 2005; Das et al., 2010; D’Cruz et al., 2010). TCR signaling in conjunction with the SLAM (signaling lymphocytic-activation molecule) is essential for early stage iNKT cell development, whereas IL-15 signaling is absolutely required for terminal expansion and maturation (Godfrey and Berzins, 2007; Godfrey et al., 2010; Engel and Kronenberg, 2012). However, how the transition between signals from early to later stages is finely regulated to promote iNKT cell development remains ill defined. In this study, we identify TRAF3 as an essential regulator of iNKT cell development and function by modulating events dependent on the strength of TCR signaling.
RESULTS AND DISCUSSION

TRAf3 is required for iNKT cell development

Studies of the T-TRAf3<sup>−/−</sup> mouse demonstrate that TRAF3 plays an essential role in T cell function and enhances TCR/CD28 signaling. Interestingly, although mature conventional T cells show striking functional defects in the absence of TRAF3, they develop normally and are present in normal numbers (Xie et al., 2011a). In contrast, we observed that T-TRAf3<sup>−/−</sup> mice exhibited an ~10-fold decrease in iNKT cells in the thymus and liver, and a twofold decrease in spleen in both percentage and total number (Fig. 1, A and B). Thus, in the current study, we investigated the role of TRAF3 in the development and function of iNKT cells. Together with reduced iNKT cell numbers, there was remarkably less IFN-γ and IL-4 produced by TRAF3-deficient iNKT cells upon in vivo stimulation with α-galactosylceramide (α-GalCer; Fig. 1 C). These observations led us to hypothesize that TRAF3 plays a distinct role in the development of iNKT cells, compared with conventional αβ T cells (Xie et al., 2011a).

Lck-Cre mediates depletion of floxed genes from as early as the DN2 T cell developmental stage in the thymus (Lee et al., 2001), earlier than that mediated by CD4-Cre. To examine the possibility that the presence of a few iNKT cells in T-TRAf3<sup>−/−</sup> mice might result from residual TRAF3 protein when iNKT cells start developing, we crossed TRAF3<sup>flx/flx</sup> mice with Lck-Cre mice. TRAF3 was completely depleted from the DN3 stage in Lck<sup>Cre</sup> TRAF3<sup>flx/flx</sup> mice, demonstrated by Western blot (Fig. 1 D). With this new mouse model, we found that iNKT cells were reduced to similar levels as in the CD4-Cre T-TRAf3<sup>−/−</sup> mouse (Fig. 1 E). This result implies that TRAF3’s role is important at very specific stages during iNKT cell development.

TRAf3 affects iNKT development in a cell-intrinsic manner

To explore whether defective iNKT cell development in T-TRAf3<sup>−/−</sup> mice is a cell-autonomous effect, we generated BM chimeric mice by transferring a 1:1 ratio of BM from C57BL/6 mice (WT, CD45.1<sup>+</sup>) and T-TRAf3<sup>−/−</sup> mice (CD45.2<sup>+</sup>), into lethally irradiated WT mice (CD45.1<sup>+</sup> CD45.2<sup>+</sup>). 8 wk later, analysis of recipients showed that the percentage of iNKT cells developing from T-TRAf3<sup>−/−</sup> BM was ~10-fold less than from WT BM in all lymphoid organs examined, whereas the percentage of CD4<sup>+</sup> and CD8<sup>+</sup> T cells was not different (Fig. 1, F and G). Similar results were obtained with 1:20 or 20:1 ratios of mixed BM (unpublished data). In addition, the expression levels of CD1d, CD150, and Ly108 on double-positive thymocytes were not different between T-TRAf3<sup>−/−</sup> and littermate control (LMC) mice (unpublished data). To determine whether this is a TRAF3-specific cell-autonomous effect, T-TRAf3<sup>−/−</sup> BM was transduced with a virus expressing the traf3 gene and transplanted into sublethally irradiated Rag1-deficient (Rag1<sup>−/−</sup>) mice. Analysis of these mice 8 wk after reconstitution showed that forced expression of TRAF3 significantly increased the percentage of iNKT cells, but empty viral vector had no effect (Fig. 1 H). These results demonstrate that TRAF3 affects iNKT cell development in a cell-intrinsic manner.

TRAf3 affects iNKT cell development after positive selection

It is well established that iNKT cell development in the thymus undergoes four distinct stages (Godfrey, et al., 2010). The presence of a few thymic iNKT cells in T-TRAf3<sup>−/−</sup> mice allowed us to estimate at which stages TRAF3 exerts its effect. With enriched thymic iNKT cells, we observed 10-fold fewer absolute numbers of iNKT cells in T-TRAf3<sup>−/−</sup> than in LMC mice. However, the percentage of stage 0 and stage 1 iNKT cells was more than sevenfold higher, and for stage 2 was twofold higher; in contrast, the percentage of iNKT cells in stage 3 was lower in T-TRAf3<sup>−/−</sup> mice than in LMC mice (Fig. 2 A). Therefore, the absolute number of iNKT cells in stage 2 and, in particular, stage 3 was drastically reduced.
whereas the numbers in stage 0 and 1 were relatively normal or just slightly reduced in T-TRAF3−/− mice (Fig. 2 B). This result indicates that TRAF3 is not essential for precursor selection (stage 0) and transition from stage 0 to 1 but is critical for the burst proliferation from stage 1 to stages 2 and 3. Consistent with this observation, there was reduced proliferation of iNKT cells at stages 1 and 2 in T-TRAF3−/− mice (Fig. 2 C). These results, in conjunction with findings in LckCreTRAF3flox/flox mice that TRAF3 deficiency did not completely block iNKT cell development, indicate that the involvement of TRAF3 in iNKT cell development is stage specific. We next explored whether the expression level of traf3 differs in distinct stages. Results indicate that traf3 gene expression increased during iNKT cell development, being highest in stage 3 and lowest in stages 0 and 1 (Fig. 2 D). The expression level of traf3 suggests that TRAF3 might play a more important role in mature iNKT cells.

**Defective response of TRAF3−deficient iNKT cells to IL-15 stimulation**

IL-15 is important in terminal stage expansion, maturation, and homeostasis of iNKT cells (Matsuda et al., 2002; Castillo et al., 2010; Gordy et al., 2011). Our observations with T-TRAF3−/− mice share similarities with the phenotypes reported for IL-15–deficient mice, which encouraged us to explore whether TRAF3 deficiency affects IL-15–mediated iNKT cell expansion and homeostasis. Because the expression of CD122 on iNKT cells can be regulated by T-bet (Townsend et al., 2004; Matsuda et al., 2007), we first examined the expression level of T-bet and CD122. Flow cytometry data showed that expression of T-bet is remarkably reduced (Fig. 3 A), as is CD122 (Fig. 3 B) in TRAF3−/− thymic iNKT cells. Notably, T-bet or CD122 exhibited only basal level expression at stages 0 and 1 but were drastically up-regulated in stages 2 and 3 in LMC mice, consistent with the distinct roles of TCR and IL-15 signaling in different stages of iNKT development and maturation. To investigate whether the reduction of T-bet and CD122 affected IL-15–mediated proliferation of iNKT cells, thymocytes depleted of CD8+ T cells were CFSE labeled and incubated with IL-15 for 4 d. Assessment of CFSE dilution by flow cytometry showed much lower cell proliferation in TRAF3−/− than in LMC stages 2 and 3 iNKT cells (Fig. 3 C). IL-15–induced phosphorylation of STAT5 was also impaired in T-TRAF3−/− iNKT cells (Fig. 3 D). These results indicate that TRAF3 deficiency impairs IL-15–mediated proliferation of iNKT cells by reducing T-bet and CD122 expression.

Bcl-xL is a major target of IL-15 signaling and essential for maintaining iNKT cell survival (Gordy et al., 2011). Examination of Bcl-xL expression in T-TRAF3−/− mice shows that stage 3 thymic iNKT cells expressed much less than LMC (Fig. 3 E). To explore whether this results in impaired survival, we estimated survival status of iNKT cells in vitro. Upon tetramer stimulation, stage 3 T-TRAF3−/− iNKT cells underwent much faster cell death at all time points examined, but

![Figure 3. Defective response of TRAF3-deficient iNKT cells to IL-15 stimulation.](image-url)

Enriched thymic iNKT cells were surface stained as in Fig 2 and further stained for T-bet (A) or Bcl-xL (E). (A, left) Mean fluorescence intensity (MFI) of T-bet in a single representative mouse from each group. (A, right) Bar graph representing a summary of relative T-bet expression in TRAF3−/− versus LMC stage 2 iNKT cells (the MFI for the LMC was set at 100%). (B) CD122 expression on stage 2 iNKT cells. Error bars in A and B represent mean values ± SD of four mice. (C) Proliferation of stage 2 and 3 thymic iNKT cells was evaluated by the percentage of CFSE dilution. Error bars are mean values ± SD of three separate experiments. (D) IL-15–induced phosphorylation of STAT5 in thymic iNKT cells (data represent three independent experiments, LMC = 85 ± 4.7%, T-TRAF3−/− = 51 ± 5.1%, P < 0.01). (E) Bcl-xL expression in stage 3 iNKT cells. Data are from one of three mice with similar results (LMC vs. T-TRAF3−/−: MFI = 3,636 ± 606 vs. 1,809 ± 23, P = 0.027). (F) Tetramer-enriched thymic iNKT cells were incubated for indicated time. Surface staining to distinguish different developmental stages was performed and followed by staining for Annexin V and PI. Error bars are mean values ± SD of three separate experiments.
this was confined to stage 3 of maturation (Fig. 3 F). Similarly, more cell death was found in TRAF3\(^{-/-}\) iNKT cells without stimulation (unpublished data). These results suggest that reduced expression of Bcl-xL could contribute to decreased iNKT cell number in T-TRAf3\(^{-/-}\) mice. Collectively, TRAF3 deficiency led to a failure to up-regulate T-bet and CD122 and further impaired both IL-15 mediated terminal expansion and survival of iNKT cells.

**TRAF3 is required for T-bet up-regulation by impacting TCR signaling during iNKT cell development**

Up-regulation of T-bet can be initiated and maintained by TCR signaling (Bajénoff, et al., 2002; Matsuoka, et al., 2004) in synergy with IFN-γ/STAT-1 signaling (Lighvani, et al., 2001; Afkarian, et al., 2002). Our previous report that TRAF3 is required for normal levels of TCR signaling in conventional T cells (Xie et al., 2011a) prompted us to hypothesize that defective up-regulation of T-bet during the transition from stage 1 to stage 2 results from reduced strength of TCR signaling. With optimized stimulation by PBS57-CD1d tetramer and staining conditions, we found that phosphorylation of the MAP kinases ERK and p38 was detectable in LMC iNKT cells within 2–5 min of tetramer stimulation. However, ERK phosphorylation at all stages of T-TRAf3\(^{-/-}\) iNKT cell development was defective compared with that seen in LMC iNKT cells (Fig. 4 A). Similar results were obtained for p-p38 (unpublished data). To verify that weakened TCR signaling was responsible for reduced T-bet expression, mice were treated with anti-CD3 Ab. 18 h later, thymic iNKT cells were enriched and stained for T-bet. Results show that T-bet expression was markedly increased in stage 1 of LMC iNKT cells, which was equal to the level in stages 2 and 3 in untreated control iNKT cells. However, there was only minimal up-regulation of T-bet in T-TRAf3\(^{-/-}\) iNKT cells after anti-CD3 Ab treatment (Fig. 4 B). Notably, T-bet expression in stages 2 and 3 was not further up-regulated upon stimulation in LMC iNKT cells, indicating that it reaches its peak levels at these stages. These results support our hypothesis that TRAF3 is required for optimal TCR signaling to up-regulate T-bet expression during the transition from stages 1 to 2. To assess whether defective T-bet expression is responsible for impaired iNKT cell development, T-bet overexpression by retroviral transduction in BM chimeric mice showed that the percentage of iNKT cells was dramatically increased in T-bet-overexpressing cells, compared with virus-negative cells or those transduced with viral vector only. Consistently, CD122 expression was also increased in T-bet-overexpressing cells (Fig. 4, C and D). This result indicates that T-bet overexpression can rescue defective iNKT cell development induced by TRAF3 deficiency.

**TRAF3-deficient iNKT cells are functionally impaired**

During iNKT cell development, stage 2 immature cells migrate into the periphery, further mature into functional iNKT cells, and acquire NK cell surface markers. Analysis of cell surface markers of splenic iNKT cells showed that they are either slightly to moderately decreased or unchanged (Fig. 5 A), indicating that a few iNKT cells can still mature in the absence of TRAF3. To investigate whether the function of these mature iNKT cells is impacted by TRAF3 deficiency, cytokine production by peripheral iNKT cells stimulated with α-GalCer was examined. Intracellular staining revealed that the percentage of TRAF3\(^{-/-}\) iNKT cells producing IFN-γ and IL-4 was markedly reduced compared with LMC iNKT cells (Fig. 5 B). Results from BM chimera mice showed that defective IFN-γ and IL-4 production by TRAF3\(^{-/-}\) iNKT cells was a cell-intrinsic defect (unpublished data). This result is comparable with the phenotype of TRAF3\(^{-/-}\) conventional CD4\(^{+}\) and CD8\(^{+}\) T cells, which show defective cytokine production in response to TCR stimulation (Xie, et al., 2011a). Examination of TCR signaling showed that TRAF3\(^{-/-}\) splenic iNKT cells exhibited much less p-ERK and p-p38 after tetramer stimulation than LMC iNKT cells (Fig. 5 C). These results indicate that TRAF3 is also required for TCR signaling during mature iNKT cell function.
The stage-specific effect of TRAF3 on iNKT cell development suggests that TRAF3 is a key connector between TCR and IL-15 signaling. TRAF3 deficiency didn’t completely block the development of stages 2 and 3, indicating the involvement of additional regulators in the transition stages. Defective iNKT cell development is consistent with our findings that the percentage of memory (CD44hiCD122hi) CD8+ T cells is decreased in T-TRAF3−/− mice (unpublished data). These defects in both iNKT cells and CD8+ T cells indicate that they might share similar mechanisms. One possibility is that when they become mature T cells with down-regulation of CD24, TRAF3 plays more important roles in optimizing TCR signaling. Our findings also raise future questions: why is TRAF3 required for iNKT cell but not conventional T cell development? Does TRAF3 play distinct roles in different T cell subsets? Precisely how does TRAF3 participate in proximal TCR signaling? All are the focus of our ongoing studies. The findings of this study reveal TRAF3 to have rheostat function in modulating TCR signal strength. Our results highlight the diverse roles played by TRAF3 in different immune cells and the importance of TRAF3 in multiple aspects of T cell biology.

**MATERIALS AND METHODS**

**Animals.** TRAF3fl/fl mice were previously described (Xie et al., 2007) and backcrossed with C57BL/6 mice for 10 generations. TRAF3fl/fl mice were bred with CD45.1+ mice (Taconic) and Lck{	extsuperscript{cre}} mice (The Jackson Laboratory). CD45.2+ C57BL/6 and congenic CD45.1+ C57BL/6 mice (The Jackson Laboratory) were bred to generate CD45.2+ and CD45.1+ double-positive mice. Rag1−/− mice were provided by F. Sutterwala (University of Iowa, Iowa City, IA). Mice of 6–12 wk of age were used for all experiments. All mice were maintained in facilities under specific pathogen-free conditions at The University of Iowa and were used in accordance with National Institutes of Health guidelines under an animal protocol approved by the Animal Care and Use Committee of the University of Iowa.

**Cytokine detection.** For serum cytokine detection, 5 μg α-GalCer (Avanti Polar Lipids, Alabaster, AL) in 100 μl PBS was injected intraperitoneally into mice. Serum was collected at 2 and 5 h after injection. IFN-γ and IL-4 were measured by ELISA according to the manufacturer’s instructions (ELISA Ready-SET-Go; eBioscience). For secondary activation of B cells, splenocytes were collected 5 h after injection and stained with antibodies against CD69 and B220. To detect cytokine production by single cells, splenocytes were isolated at 1.5 h after α-GalCer injection and incubated with Brefeldin A in vitro for another 2 h. Intracellular staining for IL-4 and IFN-γ was performed using Cytofix/Cytoperm reagents (BD).

**Retrovirus transduction and bone marrow chimeras.** Recipient CD45.1+CD45.2+ congenic C57BL/6 mice were given 1,100 rad γ-irradiation 16 h before transfer. BM cells harvested from the tibiae and femurs of T-TRAF3−/− (CD45.2+) and WT (CD45.1+) mice were depleted of B220+ and CD3+ cells by magnetic bead separation (Miltenyi Biotec). 5 × 10⁶ cells from each mouse strain were mixed and injected intravenously into recipient mice.

For virus packaging, the mouse traf3 or Tbx21 gene was cloned and inserted into the retrovirus backbone pMIG. pMIG:pMIG-traf3 or pMIG-Tbx21 and helper vector pCLECO were cotransfected into 293T epithelial cells using lipofectamine (Invitrogen). Supernatant was harvested after 48 h. Lineage-negative BM cells were purified using a kit (Miltenyi Biotec), stimulated overnight with a cytokine combination (IL-6, IL-3, and SCF [Peprotech]), and transduced with viral supernatant. Rag1−/− mice were sublethally irradiated with 500 rad γ-irradiation and rested overnight. 0.5 × 10⁶ transduced BM cells were transferred by i.v. injection. The resulting chimeras were analyzed 8 wk later.

**Flow cytometry.** Single-cell suspensions were prepared from thymi and spleens, and erythrocytes were lysed. Liver mononuclear cells were isolated (Xie et al., 2011a). For flow cytometry staining, nonspecific Ab binding was blocked with anti-mouse CD16/32 mAb and cells stained with fluorescently labeled antibodies against TCR-β (H57-597), CD4 (L3T4), CD8α (53–67), B220 (RA3-6B2), CD24 (M1/69), CD44 (IM7), NK1.1 (PK136), CD69 (H1.2F3), CD25 (eBio7D4), NK2D2 (A10), CD49b (DSX), CD1d (1B1), CD45.2 (104), CD45.1 (A20), CD122 (TM-b1), IL-4 (BVD6-24G2), IFN-γ (XM1G1.2), T-bet (eBioB10), SLAM6 (13G3-19D), and CD150 (9D1). For Bcl-xL staining, cells were stained with anti-Bcl-xL Ab (5H4) followed by anti-mouse–APC secondary Ab. All antibodies were purchased from eBioscience, BD, or Cell Signaling Technology. In intracellular staining of T-bet and Bcl-xL, cells were first stained for surface markers, fixed and permeabilized with the Foxp3 Staining Buffer Set (eBioscience), and stained

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**Figure 5. TRAF3-deficient iNKT cells display defective response to stimulation.** (A) Splenocytes from LMC or T-TRAF3−/− mice were stained for indicated surface markers. iNKT cells were gated and shown in histograms. Data shown are representative of one of six mice (for MFI of TCR-β, NK2G2, and NK1.1, P > 0.05; for MFI of CD44, CD49b, and CD69, P < 0.05). (B) Intracellular IFN-γ and IL-4 in splenic iNKT cells were evaluated by flow cytometry after α-GalCer stimulation. Data shown are representative of one of four mice (for IFN-γ, P = 0.002 [LMC vs. T-TRAF3−/−]; for IL-4, P = 0.0027). (C) Splenocytes were stained with PE-labeled PBS57-CD1d tetramer at 4°C for 45 min and then incubated at 37°C for 2 min. Surface staining for TCR-β and intracellular staining for p-ERK and p-p38 were performed after fixation and permeabilization. Right: p-p38 staining. Left: relative MFI comparing Tr versus Ctl, Ctl set at 100%. Error bars are mean values ± SD of three independent experiments.
with relevant Abs. For thymic iNKT cell purification, thymocytes were stained with PE-labeled PBS57-CD1d tetramer, followed by anti-PE beads and purified with a magnetic column (Mátyás Biotech). Alternatively, thymic iNKT cells were enriched by depleting CD8+ cells, using PE-labeled anti-CD8 Ab and anti-PE beads. BrDU in vivo incorporation was measured according to the manufacturer’s instructions (BrDU Flow kit; BD). Flow cytometric analysis and cell sorting were performed using a FACS LSRII or Aria (BD) at The University of Iowa Flow Cytometry Facility. Results were analyzed with FlowJo software (Tree Star).

**In vitro survival assay.** Enriched thymic iNKT cells were stained with or without PBS57-CD1d tetramer and incubated in complete RPMI medium. Samples were taken at various time points and stained for cell surface markers and then stained with FITC-labeled Annexin V (eBioscience) in Annexin V binding buffer for 15 min. Finally, propidium iodide (PI) was added and cells were analyzed by flow cytometry without fixation.

**In vitro proliferation assay.** Thymocytes depleted of CD8+ cells were labeled for 10 min with 10 µM CFSE in accordance with manufacturer’s instructions (Invitrogen). IL-15 was added at indicated concentrations and cells were incubated in complete medium for 4 d. CFSE dilution was analyzed by flow cytometry.

**TCR signaling induced by PBS57-CD1d tetramer.** Thymic iNKT cells enriched at 4°C with PE-labeled PBS57-CD1d tetramer or splenocytes stained with tetramer were incubated in a 37°C water bath for 2 or 5 min. Cells were fixed immediately with 4% paraformaldehyde for 10 min at room temperature, washed, and permeabilized with 0.1% Triton X-100 for 5 min. Surface staining was performed after washing, followed by intracellular staining for phospho-ERK or phospho-p38 (Cell Signaling Technology). Cells were analyzed by flow cytometry.

**In vivo stimulation with anti-CD3 Ab.** For T-bet up-regulation experiments, LMC or T-TRAF3−/− mice were injected intravenously with 25 µg anti-CD3 Ab (2C11; eBioscience). Thymic iNKT cells were enriched after 18 h and surface-stained for CD24, NK1.1, and CD44, followed by intracellular staining for T-bet. Cells were analyzed by flow cytometry.

**Cell sorting and Western blots.** Thymocytes from LckCre;TRAF3flox/flox mice were stained and sorted for DN1, DN2, and DN3. Cell lysates were separated by 10% SDS-PAGE and transferred to PVDF membranes. Anti-TRAF3 (M20) and anti-actin (C-2; Santa Cruz Biotechnology, Inc.) Abs were used for TRAF3 and actin detection.

**Real-time PCR.** Different stages of thymic iNKT cells were sorted. RNA was extracted with RNeasy Protocol (QiAGEN) and cDNA synthesized using SuperScript II (Invitrogen). RT-PCR was performed on an ABI PRISM 7900 Sequence Detection System (Applied Biosystems) using SYBR Green using SuperScript II (Invitrogen). RT-PCR was performed on an ABI PRISM 2720 real-time PCR system (Applied Biosystems) using 10 µM of each primer. Real-time PCR was performed on a 96-well plate using SYBR Green and followed by intracellular staining for CD4. Cells were analyzed by flow cytometry.

**Statistics.** Data represent mean ± SEM. Statistical comparisons of differences between sample means used Student’s t test.

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