

ORIGINAL**Decreased expression of MHC class II and cathepsin E in dendritic cells might contribute to impaired induction of antigen-specific T cell response in NC/Nga mice**

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Abstract : NC/Nga (NC) mice are an animal model for human atopic dermatitis. We found that induction of antigen (Ag)-specific T cell response is diminished in ovalbumin (OVA)-immunized NC mice. Ability of Ag presentation in NC mouse dendritic cells (DCs) was significantly weaker than that in BALB/c and DBA/2 mouse DCs. Expression levels of MHC class II molecules and cathepsin E in NC mouse DCs were significantly lower than those in BALB/c and DBA/2 mouse DCs. These results indicate that low expression levels of MHC class II and cathepsin E might contribute to the defect in induction of Ag-specific T cells in NC mice. *J. Med. Invest.* 59 : 95-101, February, 2012

Keywords : *cathepsin E, dendritic cells, MHC class II, NC/Nga mice*

INTRODUCTION

NC/Nga (NC) mice have been shown to develop human atopic-like skin lesions with elevated serum IgE level when kept in conventional conditions (1, 2). NC mice have been used as model animals for evaluating the effects of anti-inflammatory drugs and food-derived components on the development of atopic dermatitis. In addition to allergic response, NC mice have been shown to possess unique immunological phenotypes, including a defect in toll-like receptor 9-mediated activation, deletion polymorphism of the FcγIIb gene promoter region, and a defect in oral tolerance (3-5). In this study, we found that NC mice have and impaired antigen (Ag)-specific T cell response after ovalbumin (OVA)

immunization. Therefore, we focused on the function of antigen (Ag)-presenting cells in NC mice

MATERIALS AND METHODS*Mice and immunization*

Specific pathogen-free female NC (H-2^d) (6), BALB/c (H-2^d) and DBA/2 (H-2^d) mice were purchased from SLC (Hamamatsu, Japan). DO11.10 mice were purchased from Jackson Lab. (Bar Harbor, ME, USA). All studies were performed in accordance with the ethical guidelines for animal experimentation by the Institution of Health Bioscience, the University of Tokushima and were approved by the institutional review board of the animal ethics committee.

Mice were intraperitoneally immunized with 10 μg of OVA (Sigma Chemical Co., MO, USA) absorbed in 1 mg of Aluminium Hydroxide Gel Adjuvant (HCl Biosector, Denmark) 2 times at a 2-week interval.

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Proliferation assay

Splenocytes from OVA-immunized mice were cultured in the medium supplemented with 10% fetal bovine serum, 50 $\mu\text{mol/L}$ 2-mercaptoethanol, 100 $\mu\text{g/mL}$ streptomycin, and 100 U/mL penicillin. Five $\times 10^5$ splenocytes were stimulated with 200 $\mu\text{g/mL}$ OVA, 1,000 $\mu\text{g/mL}$ OVA or with plate-bound anti-CD3 mAb (coated overnight at 1 $\mu\text{g/mL}$) in a 96-well flat-bottom plate at 37°C under 5% CO₂ for 72 h. For the last 8 h of culture, 1 μCi of [³H] thymidine deoxyribose (TdR) was added to the wells, and the amount of [³H]TdR incorporated was measured by a scintillation counter (Aloka, Tokyo, Japan).

Ag-presenting ability

CD11c⁺ cells were purified from 3 to 4 mice using CD11c microbeads and a MACS LS column (Miltenyi Biotec Inc., CA, USA) according to the manufacturer's instructions. CD4⁺ T cells were purified from DO11.10 transgenic mice, which express OVA-specific T cell receptor, using a CD4⁺ T cell isolation kit (Miltenyi Biotec). CD4⁺ T cells (2 $\times 10^5$ cells) from a DO11.10 mouse spleen were cultured in a 96-well plate with CD11c⁺ cells (10⁴ cells) in the presence of 200 $\mu\text{g/mL}$ OVA for 48 hr at 37°C/5% CO₂. Cultures were pulsed with 1 μCi of [³H]TdR for the last 8 hr of culture. Incorporated [³H]TdR was determined by a scintillation counter.

Flow cytometry

Expression of MHC class I, MHC class II and CD80 molecules was analyzed by flow cytometric analysis. Purified CD11c⁺ cells were incubated with FITC-conjugated anti-mouse H-2K^d mAb (clone SF1-1.1), PE-conjugated anti-mouse I-A mAb (clone M5/114/15.2) or FITC-conjugated anti-CD80 mAb (clone 16-10A1) or 30 min at 4°C. After washing the cells, stained cells were analyzed with a FACSCalibur and CellQuest software (Becton Dickinson, CA, USA).

Quantitative PCR analysis

To assess gene expressions, mRNA levels of genes involved in MHC class II expression were determined by reverse transcription of total RNA followed by PCR analysis. Total RNA was isolated from CD11c⁺ cells using a RNeasy Micro kit (Qiagen Science, MD, USA). First-strand cDNA was reverse-transcribed at 42°C for 60 min and at 95°C for 5 min from 2 μg of the extracted total RNA with M-MLV

reverse transcriptase (Invitrogen, CA, USA) and a random primer. We performed real-time PCR using specific primers and SYBR green dye (Takara Bio, Japan) in a Light Cycler real-time PCR system (Roche Diagnostics, Germany) according to the manufacturer's instructions. The primers used were 5'-CAGATACCATCAACTGCGACCA-3' (sense) and 5'-TCTGCTCCAATGTGCTCTATGAAG-3' (antisense) for CTIIIT, 5'-AGTCAGTGGAGGCCAGCTTA-3' (sense) and 5'-CCAGGAGGCACCAACTGTAT-3' (antisense) for NF-Y α , 5'-GCTGACGCTGAGAACCTAC-3' (sense) and 5'-CGATGGGAGCTGAAGGTAGA-3' (antisense) for RFX5, 5'-CACAGACGGCGTTTATGAGA-3' (sense) and 5'-TGC-TCCACCTTGACATCATA-3' (antisense) for H-2^d, 5'-CTGACCCTGAAGTACCCCATTTGAACA-3' (sense) and 5'-CTGGGGTGTGAAGGTCTCAAACATG-3' (antisense) for β -actin'. The relative abundance of target transcripts was normalized to the constitutive expression of β -actin.

Western blot analysis

CD11c⁺ cells were lysed in lysis buffer containing 1% Noidet-P40, 150 mM NaCl, 10 mM Tris-Cl (pH 7.5), and 1 mM EDTA. The cell lysates were dissolved by SDS-PAGE and transferred onto a PVDF membrane. The membrane was blotted with an Ab to cathepsin S (Santa Cruz, CA, USA), cathepsin E (Novas Biologicals, CO, USA) or β -actin (Epitomics, Inc., CA, USA) and visualized using an enhanced chemiluminescence system (Amersham Biosciences, UK). Quantification of protein expression was performed by densitometry analysis of the autoradiograms.

Statistical analysis

Data were analyzed using one-way analysis of variance followed by the Scheffe post hoc test for multiple comparisons. Values in the text are means \pm SD. Differences were considered significant at $P < 0.05$.

RESULTS

Decreased Ag-specific T cell response in OVA-immunized NC mice

OVA-specific proliferation response after OVA immunization in NC mice was significantly decreased compared to that in BALB/c and DBA/2 mice. However, proliferation response to anti-CD3 mAb in NC mice was comparable to that in BALB/c and

DBA/2 mice (Fig. 1), suggesting that T cell receptor-mediated signal transduction is normal in NC mice. This finding indicates the possibility that weak induction of Ag-specific T cells in NC mice contributes to the process of Ag-specific T cell induction.

Characterization of CD11c⁺ dendritic cells (DCs) from NC mice

We next examined CD11c⁺ cell function because DCs are known to be a crucial cell population for initiation of acquired immune responses. Ag-presenting ability of DCs was evaluated by using DO11.10 CD4⁺ cells that carried the OVA-specific T cell receptor gene. In the Ag presentation process toward

to MHC class II, at first Ag-presenting cells capture Ag, digest the captured Ag in the lysosome, and then present Ag-derived peptide/MHC class II complex on the cell surface. In the case of intact OVA stimulation, proliferation response of DO11.10 CD4⁺ cells when co-cultured with NC mouse CD11c⁺ cells was significantly decreased compared to that when co-cultured with BALB/c or DBA/2 mouse CD11c⁺ cells (Fig. 2). We evaluated the ability of Ag-internalization in NC mouse CD11c⁺ cells using FITC-labeled OVA, but a significant difference was not observed compared to that in BALB/c mouse CD11c⁺ cells (data not shown). Next, cell surface expressions of MHC class I, MHC class II, and CD80 were

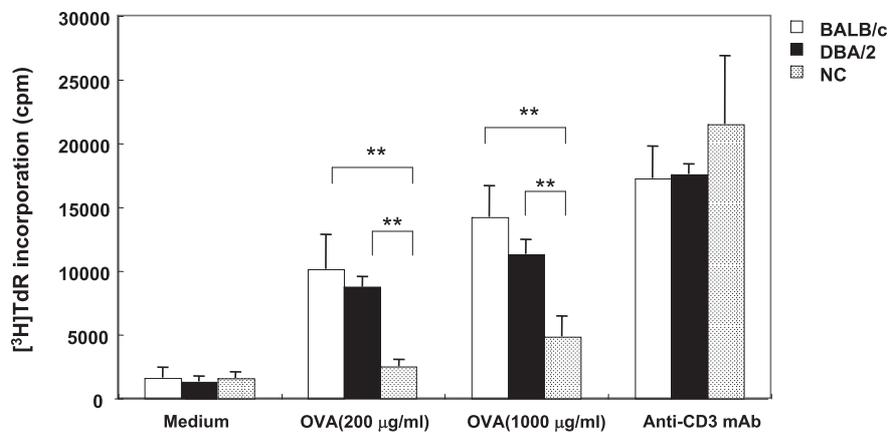


Fig. 1. OVA-specific T cell responses in BALB/c, DBA/2 and NC mice. Mice were immunized with OVA 2 times at a 2-week-interval. Two weeks after the second immunization, spleen cells were stimulated with OVA or anti-CD3 mAb for 72 hr *in vitro*. Proliferation response was determined by counting incorporated [³H]TdR activity. Results are expressed as means ± SD. White, black and hatched bars indicate BALB/c, DBA/2 and NC mice, respectively. Data are presented as means ± SD. ***P* < 0.01.

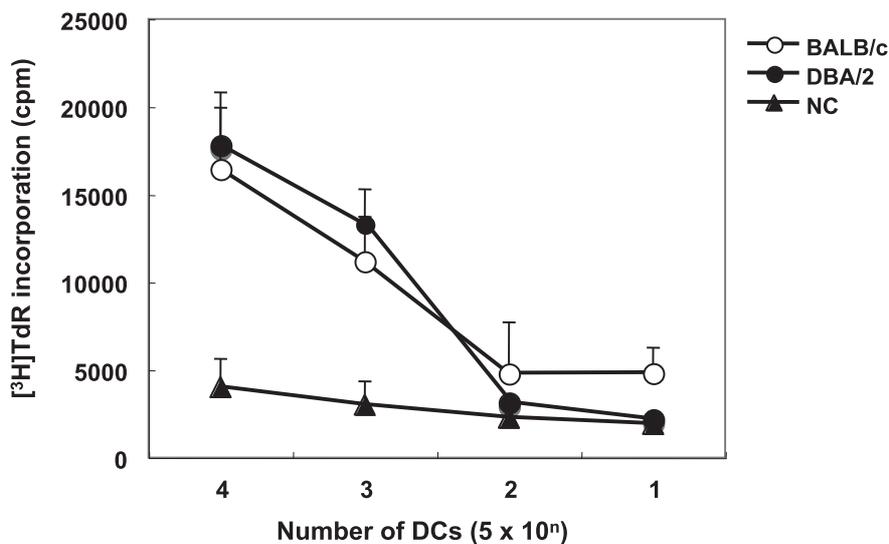


Fig. 2. Characterization of NC mouse DCs. DO11.10 CD4⁺ cells were stimulated with OVA in the presence of indicated number of DCs from BALB/c, DBA/2 or NC mice. Ag-presenting ability of DCs was evaluated by incorporated [³H]TdR activity. Levels of incorporated [³H]TdR were less than 1,500 cpm when DO11.10 CD4⁺ cells or DCs alone were cultured with OVA. Levels of incorporated [³H]TdR were less than 500 cpm when DO11.10 CD4⁺ cells or DCs alone were cultured with medium only. Results are expressed as means ± SD. The data are representative of two independent experiments of pooled samples from 3 or 4 mice.

determined by flow cytometric analysis. Expression level of MHC class II in NC mouse CD11c⁺ cells was significantly lower than that in BALB/c mouse CD11c⁺ cells. Reduced MHC class II expression was observed not only in DCs but also in B cells in NC mice (data not shown). In the case of MHC class I and CD80, expression of these molecules in NC

mouse DCs was slightly lower than that in BALB/c and DBA/2 mouse DCs (Fig. 3). Although we also evaluated transcription factors that contribute to MHC class II expression, differences in the expression of CIITA, NF-Y α and RFX5 were not observed between NC, BALB/c and DBA/2 mice (Fig. 4 B-D).

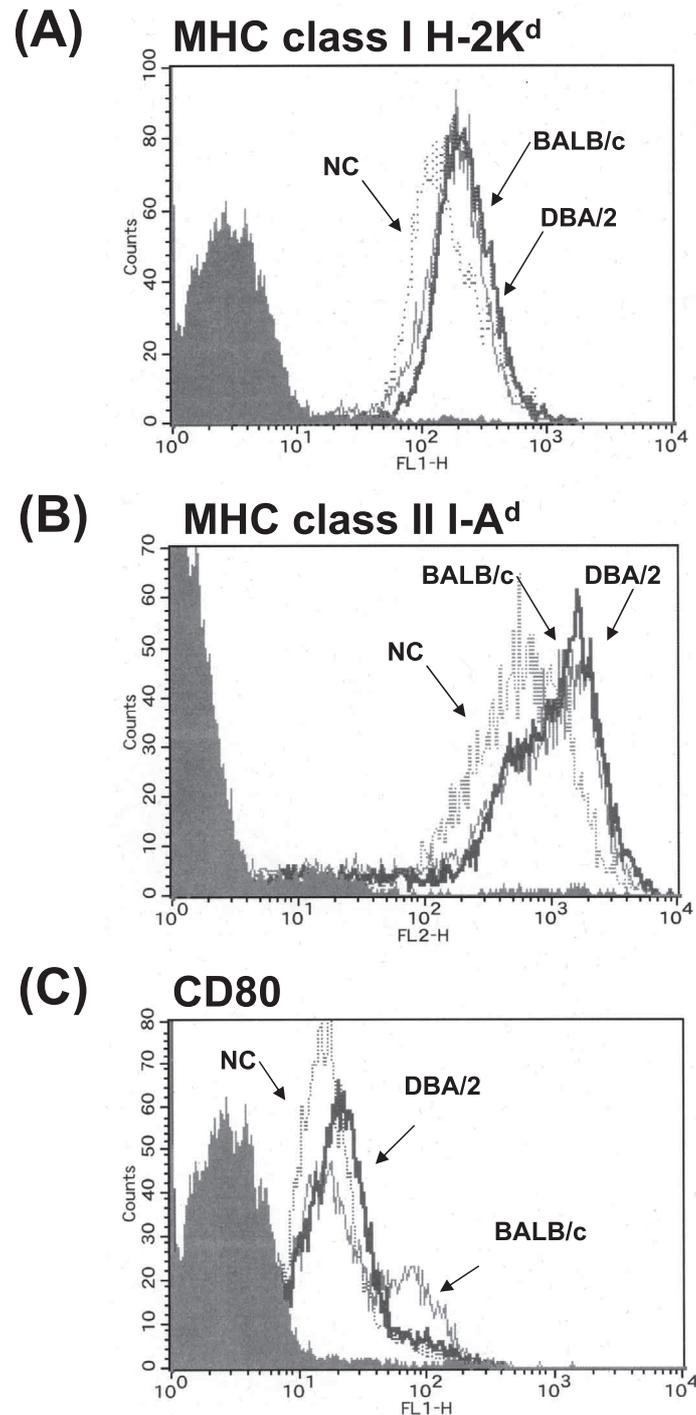


Fig. 3. Expression of MHC class I, MHC class II and CD80 in DCs from BALB/c, DBA/2 and NC mice. Expression of MHC class II was determined as described in the "Materials and methods" section. Shaded histograms represent cells stained with an isotype-matched control Ab.

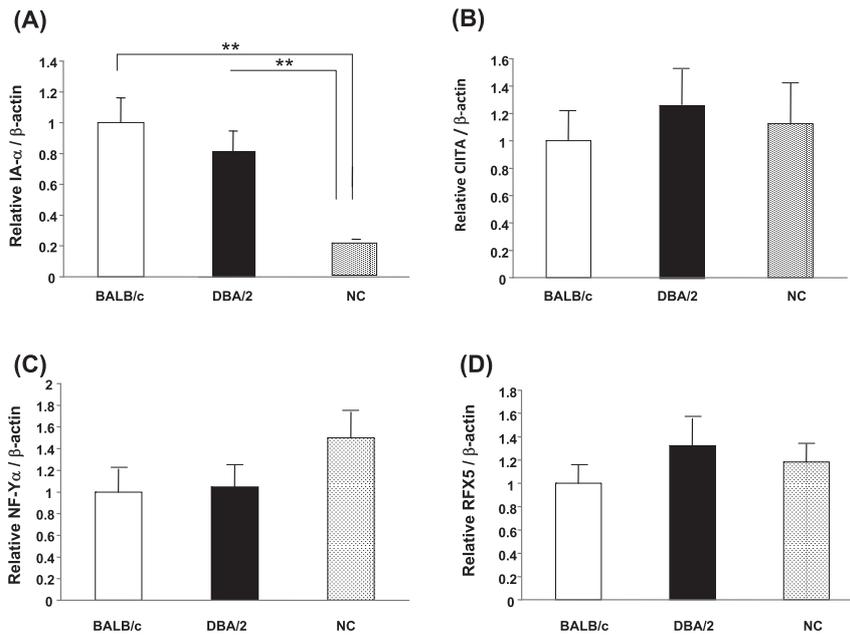


Fig. 4. CIITA, NF- γ and RFX5 mRNA expression levels in DCs. cDNA was prepared from purified CD11c⁺ cells and analyzed for expression of CIITA (A), NF- γ (B), RFX5 (C) and β -actin message by quantitative RCR as described in the “Materials and methods” section. Data are presented as means \pm SD. ** $P < 0.01$.

Decreased expression of cathepsin E in NC mouse CD11c⁺ DCs

Cathepsin S is a member of the papain family of cysteine proteases (7), and cathepsin E is an intracellular aspartic protease of the pepsin superfamily (8). Both molecules have been shown to be crucial for Ag processing and presentation in Ag-presenting

cells (9). Expression of cathepsin S and cathepsin E in DCs was determined by Western blot analysis. A difference in cathepsin S expression was not observed between BALB/c, DBA/2 and NC mouse CD11c⁺ cells. In contrast, it was found that expression of cathepsin E in NC mouse DCs was significantly decreased compared to that in BALB/c and DBA/2 mouse DCs (Fig. 5). Densitometry analysis

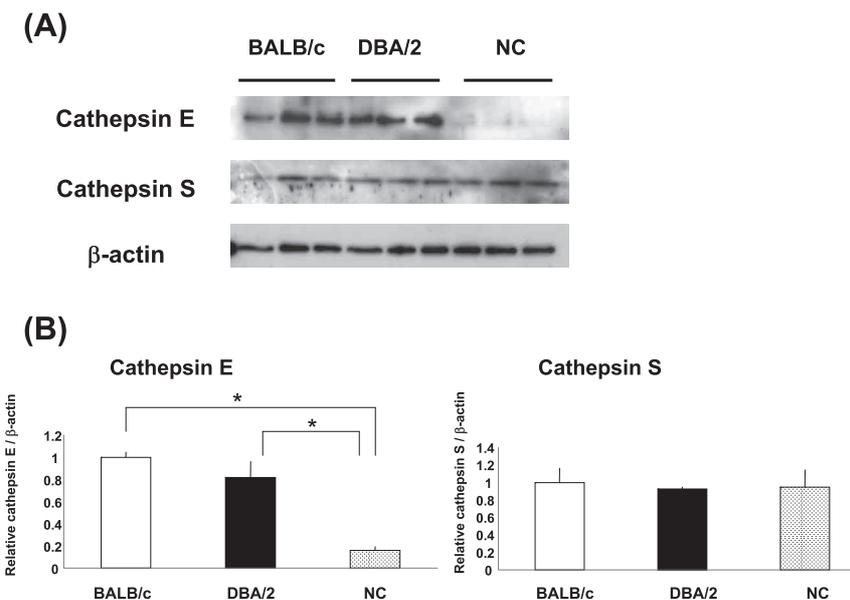


Fig. 5. Expression of cathepsin E and S in DCs. (A) Expression of cathepsin E and cathepsin S was analyzed by Western blotting with anti-cathepsin E and anti-cathepsin S mAb. To ensure the presence of equal amounts of protein, the membranes were re-probed with an anti- β -actin mAb. (B) Histograms show the cathepsin E/ β -actin and cathepsin S/ β -actin ratio. Data are presented as means \pm SD. * $P < 0.05$.

of autoradiograms showed that the relative expression level of cathepsin E in NC mice is one-tenth of that in BALB/c and DBA/2 mice.

DISCUSSION

NC mice provide a new animal model for the study of atopic-like dermatitis and allergy disease. The contribution of possible immunological factors to the development of allergic response in NC mice has been revealed (6, 10-12). In addition to the development of allergic response, it has been shown that NC mice possess unique immunologic phenotypes [3-5]. In the course of investigating the mechanism for the onset of allergic response in NC mice, we found that Ag-specific T cell response following Ag immunization was severely impaired (Fig. 1). To elucidate the mechanism for this phenotype, we focused on DC function in NC mice. Consistent with the impaired induction of Ag-specific T cells, Ag-presenting ability in NC mice was significantly weaker than that in BALB/c and DBA/2 mice (Fig. 2). Furthermore, expression level of MHC class II in NC mouse DCs was about half of that in BALB/c mouse DCs (Fig. 3). The main control element regulating transcription is a highly conserved 150-bp promoter proximal region situated upstream of the transcription initiation site. Additional promoter distal regulatory elements have been identified (13). At first, we examined the nucleotide sequence in conserved regulatory elements, named distal Y'-S' motif and proximal S-Y motifs, between NC and BALB/c mice, but a difference was not found. In addition, expression of DNA-binding proteins, CIITA, RFX and NF-Y, were examined by real-time PCR assay and no differences in expression levels were found (Fig. 4 B-D).

Several lines of evidence suggest that cathepsin E plays a role in Ag processing via the MHC class II pathway (14). Treatment with a cathepsin E-specific inhibitor inhibits Ag processing, resulting in an impaired Ag-specific T cell response. Similar to the results obtained by using a specific inhibitor, cathepsin E-deficient mouse DCs showed reduced ability of antigen presentation. Cathepsin E is mainly present in the endosome compartment in DCs. In this study, we found that expression of cathepsin E, but not that of cathepsin S, in NC mouse DCs was weak. Reduced Ag presenting ability of exogenous Ag in NC mouse DCs was similar to that in cathepsin E-deficient mouse DCs or DCs treated with a cathepsin

E-specific inhibitor. From these findings, it seems likely that a low level of cathepsin E expression contributes to defect in Ag-presenting ability in NC mice. However, it is not known whether a low level of cathepsin E expression directly affects the expression of MHC class II because expression of MHC class II in cathepsin E-deficient mouse DCs was comparable to that in wild-type mice (15). We speculate that decreased expression of MHC class II and cathepsin E synergistically affect Ag-presenting ability in DCs, resulting in impaired induction of Ag-specific T cell response in NC mice.

NC mice have been shown to develop human atopic-like skin (1, 2). Interestingly, cathepsin E-deficient mice develop atopic-like dermatitis. Tsukuba *et al.* showed that development of atopic dermatitis in cathepsin E-deficient mice contributes to the systemic accumulation of IL-18 and IL-1 β . They also showed cathepsin E activity in erythrocyte ghosts in NC mice and found that cathepsin E activity in NC mice was lower than that in C57BL/6 mice (16).

In this study, we found a new immunologic phenotype in NC mice. Low MHC class II and cathepsin E expression levels in an experimental animal have not previously been reported. Therefore, NC mice will provide a new useful animal model for investigation of induction of Ag-specific immune response and acquired immunity.

CONFLICT OF INTEREST

None of the authors have any conflicts of interest to declare.

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