

Research Article

Variations in T-Cell Response Among Cow's Milk Allergy Children With Or Without Sensitization to Casein Protein

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Abstract

Objective: To explore the potential differences in allergen-specific T immunological responses among cow's milk allergic(CMA) patients with or without casein sensitization.

Methods: A total of 25 children with IgE-dependent CMA were recruited from outpatient clinic in Beijing Children's Hospital from January to August of 2017, including 14 CMA patient with a positive sensitization to casein component, 11 CMA patient being sensitized to cow's milk, but a negative sIgE for casein component, and another 10 atopic control subjects without CMA were also included as negative control group in this study. The expression level of T cell related transcription factor in PBMCs or total CD4⁺ cells were detected by quantitative PCR, and the release of T cell related cytokines in donor's serum was determined by Luminex xMAP 200 System. Upon in-vitro stimulation with casein protein, the proliferation rates of PBMCs-derived T cells were assessed by flow cytometry, and T related cytokines production in culture supernatant was evaluated as well.

Results: A significantly higher level of GATA3 expression transcripts in PBMCs or total CD4⁺ cells were detected in casein-sensitized CMA patients than those without casein sensitization. An increase in IL-4 (74.1±16.5pg/ml vs 64.9±12.8 pg/ml, $P<0.05$) and a decrease in IL-10 (3.1±2.4 pg/ml vs 4.9±2.8 pg/ml, $P<0.05$) production were found in casein-sensitized CMA patient's plasma than patients without casein sensitization. Upon in-vitro stimulation with

cow's milk protein mixture, a significantly higher proliferation rate of casein-specific CD4⁺T cells (30.0±5.8% vs 20.7±5.5%, *P*<0.05) as well as increased productions of IL-4(4.1±13.6 vs 64.7±12.3pg/ml, *P*<0.05) and IL-13(48.7±10.3 vs 38.7±9.4pg/ml, *P*<0.05) were found in casein-sensitized CMA patients compared those without casein sensitization.

Conclusions: Our findings preliminarily revealed that a distinctive T cell mediated immune response among CMA patient with the different sensitized component.

Keywords: Cow's milk allergy; Casein; Food allergy; Sensitization; Specific IgE; Antigen-specific T-cells

Introduction

Cow's milk allergy (CMA) is a IgE-mediated immunologically adverse reactions, which affects about 2.5% of newborn infants in the first year of life [1-2]. Although double-blind, placebo-controlled food challenge(DBPCFC) is considered as gold standard for food allergy diagnosis, the application of DBPCFC in pediatric clinical practice is still limited due to its high-risk and time-consuming procedure [3-4]. Therefore, in-vitro measurement of cow's milk-specific IgE(sIgE), has been widely used in clinical practice as an alternative option for allergen laboratory diagnosis, to provide information on food allergen sensitization [5-6].

The composition of cow's milk is very complicated, there were much evidence to show being sensitized to some certain cow's milk components has proven to be more prone to a poor outcome of CMA [7-10]. Casein, as the most abundant protein, has been proven to be both more antigenic and allergenic than other cow's milk proteins [11]. Children with persistent CMA had an elevated levels of casein specific IgE antibodies (Cas-IgE) than those with transient CMA, and patients in this population were tend to have a higher risk of severe anaphylaxis and a delayed course of food reintroduce [12,13]. In addition, Machteld M et al. [14] study has proven distinctive properties of cow's milk specific-T cells responses existing among patients with persistent cow's milk, cow's milk-tolerant and nonallergic control subject, which suggesting that the potentially critical role of cow's milk allergen-specific T-cell-mediated reactivity of children with or without CMA. Accordingly, in order to clarify the detailed role of casein-specific T responses in cow's milk allergy, this study was designed to compare the casein-specific T cells response among CMA children with or without casein sensitization, basing on expression of T cell-related transcription factor, antigen-specific proliferation, and the production of T cell related-cytokine.

Material and Methods

Subjects

A total of 25 children with IgE-dependent CMA were recruited from outpatient clinic in Beijing Children's Hospital from January to August of 2017, including 14 CMA patient with a positive sensitization to casein component, 11 CMA patient being sensitized to cow's milk, but a negative sIgE for casein component, and another 10 atopic control subjects without CMA were also included as negative control group in this study. The diagnosis of CMA was based on the basis of a compatible and convincing clinical history relation with immediate cutaneous reaction

after intake of cow milk. All enrolled subjects were fully informed about the purpose of this study and nature of the study, which were approved by the medical ethics committee of Beijing Children's Hospital. Written informed consent was obtained from all parents/guardians of all the participants.

Determination of total and allergen-specific IgE antibodies

Serum levels of total IgE, specific IgE antibodies (sIgE) to cow's milk (CM-), casein (Cas), α -lactalbumin (α -La) and β -lactoglobulin (β -Lg) were measured using ImmunoCAP[®] (Phadia AB, Uppsala, Sweden). Any sensitization was regarded as positive when specific serum IgE levels were noted as being greater than 0.35 kU_A/L against the tested isolated cow's milk protein. The commercially available tests and reagents were used according to the instructions from the manufacturer.

Peripheral blood mononuclear cells (PBMCs) and total CD4⁺ T lymphocytes isolation

PBMCs were isolated from heparinized venous blood by density-gradient separation (Ficoll-Paque PLUS; GE Healthcare, Barcelona, Spain). Blood was diluted 1:2 with phosphate buffered saline (HAOYANG, Tianjin, China), then centrifuged using Ficoll density gradient solution (HAOYANG, Tianjin, China) at 2000 rpm for 20 min at 4 °C to isolate the PBMCs. Resting CD4⁺ T cells were positively selected from PBMCs using magnetic CD4 MicroBeads (Miltenyi Biotec) or negatively selected using a human CD4⁺ T cell enrichment kit (StemCell Technologies). The purity of CD4⁺ T cells was confirmed by using the FACScan system (BD Biosciences).

Cell proliferation rate evaluation upon in-vitro allergen-specific stimulation

PBMCs (2×10^6 cells/mL) were cultured in vitro for 7 days at 37 °C in 5% CO₂ with α -MEM medium alone as negative controls, or with 200 μ g/mL mixture of purified α -, β -, and κ -caseins (Sigma-Aldrich, USA), or with 4 μ g/mL phytohemagglutinin (PHA) (Sigma-Aldrich, USA) was used as positive control. Cow's milk allergen-specific T cells proliferation status was identified using ex vivo carboxyfluorescein succinimidyl ester (CFSE) dilution before antigen stimulation with modifications to the method described by Turcanu et al. [16]. Briefly, PBMC samples were labeled with 5 mmol/L CFSE for 10 min at 37 °C.

Cytokine profile analysis

A custom designed MILLIPLEX MAP kit (Millipore Corporation, Bilerica, MA, USA) and the Luminex xMAP 200 System (Luminex, Austin, Texas, USA) were used to quantify Th1 (IL-12, IFN- γ), Th2 (IL-4, IL-13), Th17 (IL-17, IL-23) and Treg (IL-10, TGF- β) in patients' serum and in cell culture supernatant following manufacturer's instructions. Briefly, the samples were thawed and incubated with cytokine-specific antibody-coupled beads, followed by incubation with indicator antibodies. After incubation, samples were subjected to the Luminex 200 System. All standards and samples were run in duplicates. Mean Fluorescent Intensity (MFI) data was collected and analyzed using a weighted 5-parameter logistic or spline curve-fitting method for calculating cytokine/chemokine concentrations (pg/ml). Results are expressed as amount of each cytokine detected after the stimulation with casein minus the amount detected after stimulation with the negative control.

RNA preparation and quantitative reverse transcription-polymerase chain reaction

Total RNA from PBMCs or isolated CD4⁺ cells were extracted using RNeasy plus mini kit (Qiagen) according to manufacturer instruction. RNA template was quantified using a NanoDrop ND1000 instrument (Thermo Fisher Scientific). RNA was converted to cDNA using PrimeScriptTM RT reagent Kit (TAKARA, Dalian, China) according to the manufacture's instruction. Quantitative reverse-transcription PCR (qPCR) was performed using a SYBR Premix Ex Taq kit (TaKaRa, Dalian, China), following the manufacturer's instruction. Primer specific for tested gene were listed in Table 1. The amplification program used was: 1 cycle of 10 minutes at 95 °C, 40 cycles of 15 seconds at 95 °C, and finally 1 cycle of 1 minutes at 60°C. All reactions were performed in triplicate. The mean value of the replicates for each sample was expressed as the quantification cycle (Ct). The Ct data were normalized using GAPDH and the relative gene expression level were calculated with the $2^{-\Delta\Delta C_t}$ method.

| Gene name | Forward primers (5'-3') | Reverse primers (5'-3') |
|----------------|-------------------------|-------------------------|
| T-bet | GGGAAACTAAAGCTCACAAAC | CTCTCCGTCGTTACCTCAAC |
| FOXP3 | CAGCACATTCCCAGAGTTCCTC | GCGTGTGAACCAGTGGTAGATC |
| ROR γ t | CTCAAAGCAGGAGCAATGGAA- | AGGGAGTGAGAAGTCAAAGA |
| GATA3 | CGGCTTCGGATGCAAGTC | GTCGAGGTTGCCCCACAG |
| GAPDH | AGCCGAGCCACATCGCT | CAGCCCTGGTGACCAGGC |

Table 1: Primers used for real-time quantitative PCR

Statistical analysis

Statistical analysis was performed using SPSS version 19.0 statistical software (IBM, Armonk, NY, USA) and graphs were generated using the prism software (GraphPad, La Jolla, CA, USA). Qualitative variables were compared between the groups by chi-square tests and continuous variables were analyzed by the nonparametric Mann-Whitney U test for unpaired comparisons. *P* values of less than 0.05 were considered significant.

Results

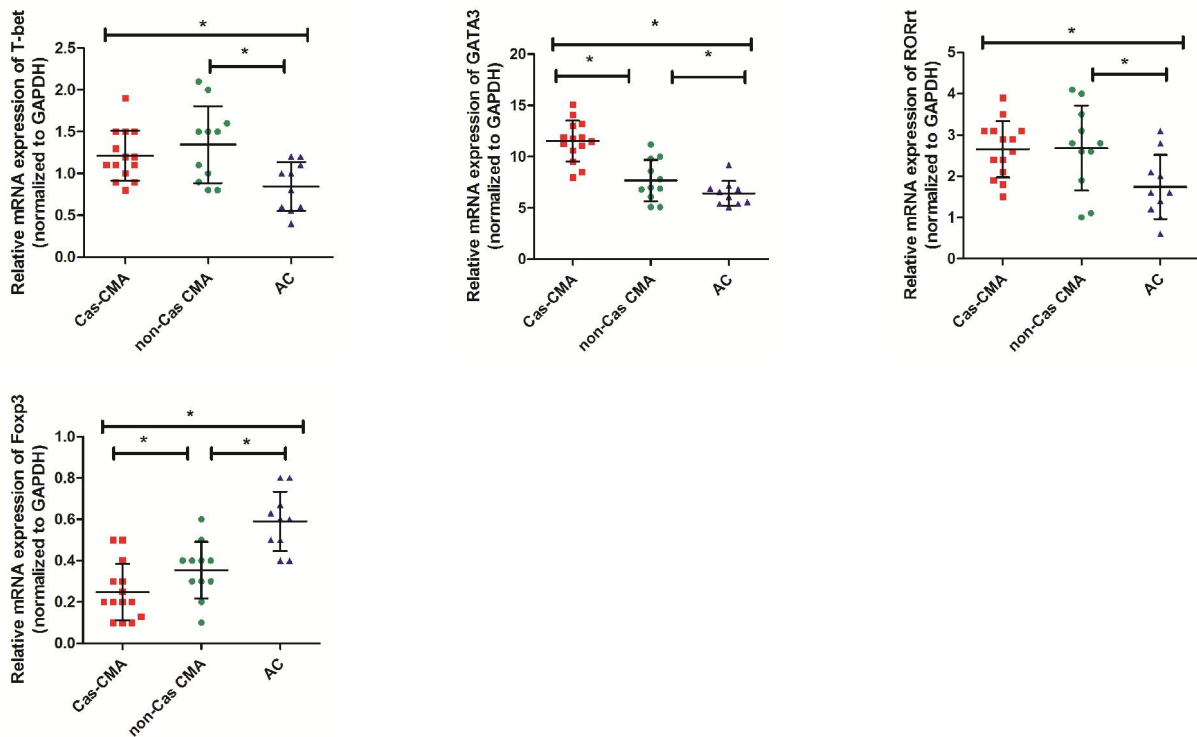
Study subjects

A total of 25 children with IgE-dependent CMA were recruited in the present study, including 11 CMA patients without casein sensitization (non-Cas CMA group, age range: 0.3-6.0 years, mean 3.2 years) and 14 CMA patients with casein sensitization (Cas-CMA group, age range: 0.1-9.1 years, mean 4.5 years). The diagnosis of CMA was based on convincing clinical history, usually occurring immediately or starting several hours after the intake of moderate to large amounts of milk or its products or infant formula. Another 10 atopic controls (AC group, age range: 1.0-8.6 years, mean 4.8 years) were enrolled in the present study, among of them there were 5 children with a physician-diagnosed food allergy but no history of CMA.

Gene expression of T-related transcription factors

Using quantitative real-time assay, expression level of genes encoding transcription factors of Th1(T-bet), Th2(GATA3), Th17(ROR γ t) and Treg(Foxp3) subsets were compared among CMA children with casein sensitization (Cas-CMA), CMA children without casein sensitization (non-Cas CMA), and atopic controls (AC). Our results showed that there was an intra-group difference existing in GATA3 or Foxp3 expression respectively, showing that a higher level of GATA3 expression (11.53 ± 2.00 vs 7.67 ± 2.10 , $P<0.05$) and a decreased level of Foxp3 (0.25 ± 0.14 vs 0.35 ± 0.14 , $P<0.05$) were detected in casein-sensitized CMA patients in comparison with CMA patients without casein sensitization (Figure 1a). Next, in order to describe the changing trend of T-related transcription factor among different subgroups more precisely, we isolated the total CD4⁺ cell population from peripheral blood and examine the expression profiles of T-related transcription factors. However, our results showed that the intra-group differences were only found in GATA3 expression (10.09 ± 2.49 vs 7.16 ± 1.97 , $P<0.05$), and ROR γ t (2.24 ± 0.42 vs 1.68 ± 0.76 , $P<0.05$), respectively. There is a slightly decreased level of FoxP3 expression (0.29 ± 0.10 vs 0.33 ± 0.12 , $P>0.05$) in CD4⁺T cells isolated from casein-sensitized CMA patients than those without it, this difference was not statistically significant (Figure 1b).

(a)



(b)

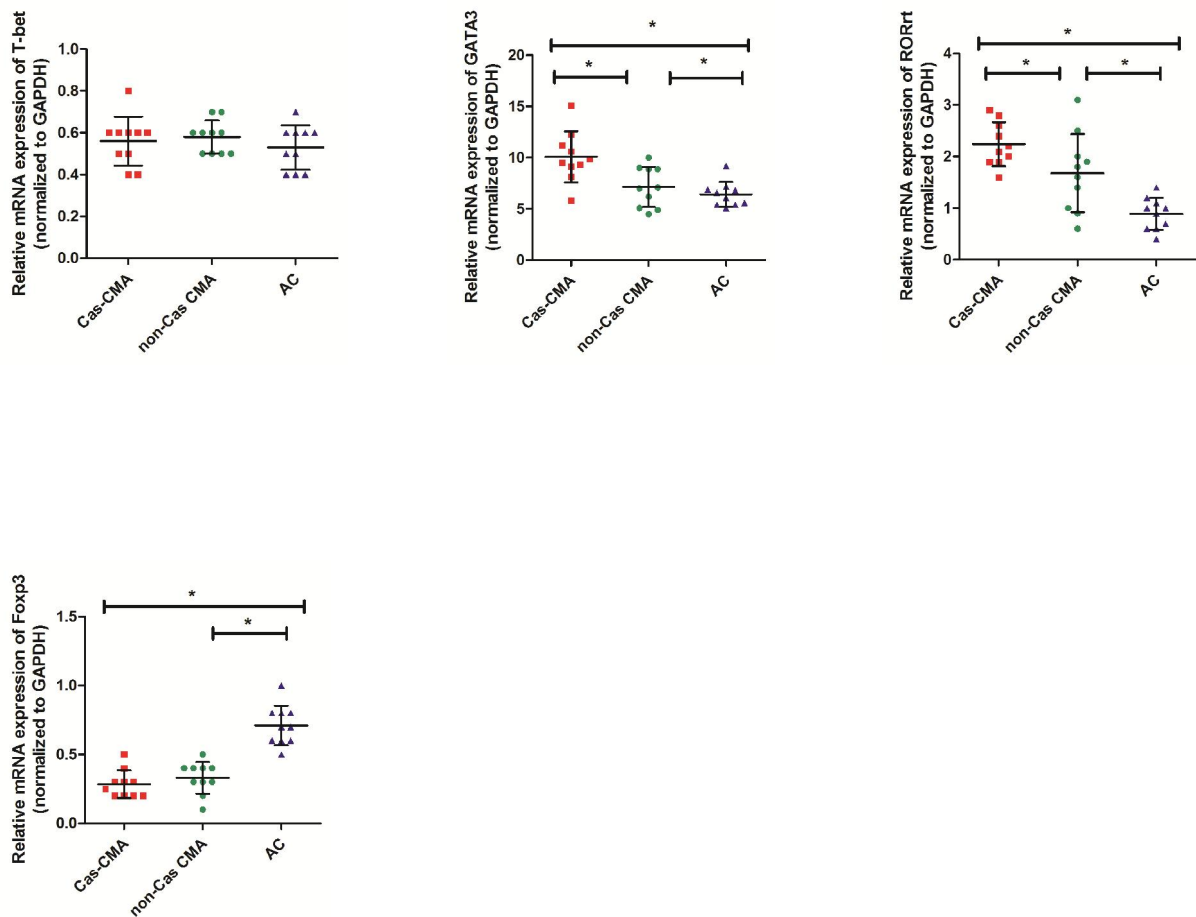


Figure 1: Analysis of T subpopulation related transcription factors expression among study subjects

Th1/Th2/Th17/Treg cytokines measurement

T-cell subpopulation related cytokines in patient's serum were measured using a multiplex Luminex[®] detection system, enabling to detect subtle variations in a small-volume sample with a higher sensitivity. Our results showed that the average concentrations of Th1 cytokine (IL-12:13.3±3.3 pg/ml vs 11.0±2.3 pg/ml, $P<0.05$), Th2-related cytokines (IL-4:69.5±14.7pg/ml vs 56.5±11.2pg/ml, $P<0.05$; IL-13: 37.5±15.9pg/ml vs 33.6±10.2pg/ml, $P<0.05$), and Th17-related cytokines (IL-17: 28.6±12.3pg/ml vs 23.4±5.0pg/ml, $P<0.05$) were significantly increased in CMA patients than that in atopic controls. However, an obviously decreased expression of Treg-related cytokines were noted in CMA children than atopic control subjects (IL-10:4.0±2.6pg/ml vs 3.5±1.8 pg/ml, $P<0.05$; TGF-β: 3.6±1.3pg/ml vs 5.6±1.3pg/ml, $P<0.05$). Furthermore, through an intra-group comparison among CMA patients with or without casein sensitization, there showed an increase in IL-4 (74.1±16.5pg/ml vs 64.9±12.8 pg/ml, $P<0.05$) with a decrease in IL-10 (3.1±2.4 pg/ml vs 4.9±2.8 pg/ml, $P<0.05$) production in casein-sensitized CMA patient's plasma than those patients without casein sensitization (Figure 2).

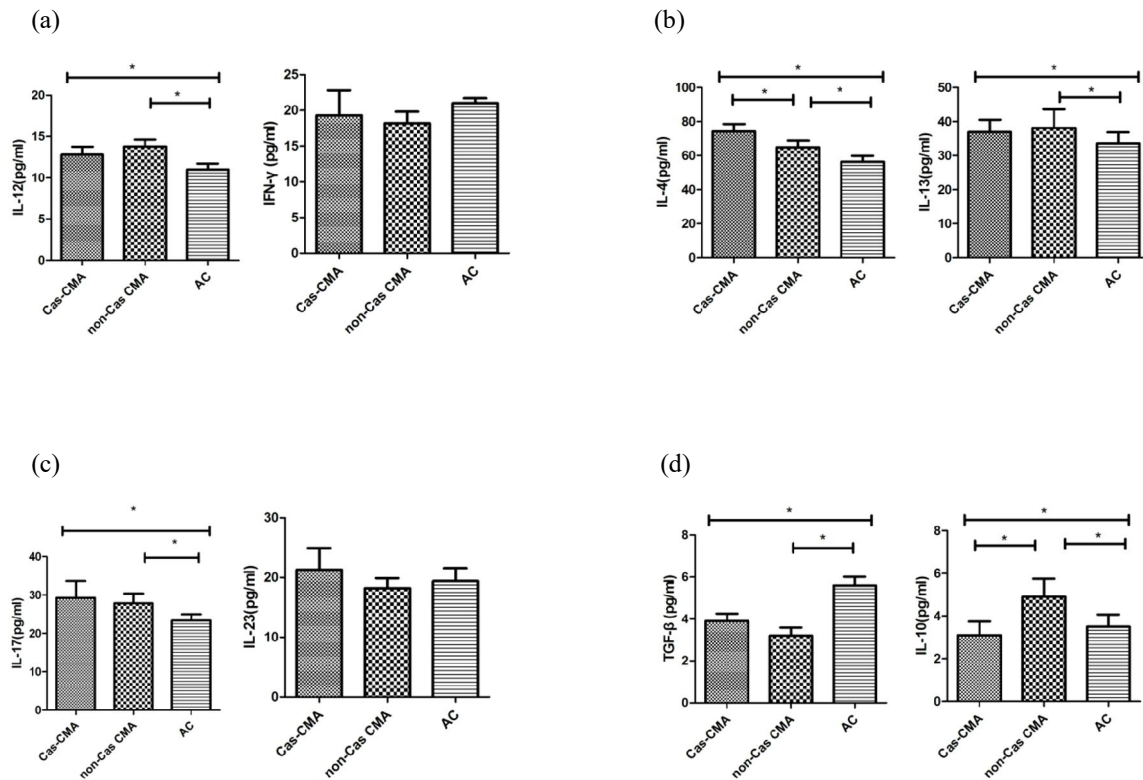


Figure 2: Comparison of circulating Th1/Th2/Th17/Treg related cytokine levels in peripheral blood among included subjects.

Cytokine levels of (a)Th1 (IL-12, IFN- γ), (b)Th2 (IL-4, IL-13), (c)Th17 (IL-17, IL-23) and (d)Treg (IL-10 and TGF- β) in peripheral blood from enrolled patient's blood were detected using a custom designed MILLIPLEX MAP kit (Millipore Corporation, Bilerica, MA, USA) and the Luminex xMAP 200 System (Luminex, Austin, Texas, USA). Each symbol represents individual samples and horizontal lines show the median values. * P value < 0.05 considered significant.

T cells proliferation and cytokines release after in-vitro stimulation

In order to explore the potential differences in cow's milk-specific T cell reactivity, PBMCs were isolated from study subjects and stimulated with a commercial cow's milk protein extract for 7 days in vitro. The proliferation status of cow milk-specific CD4⁺ T-cell frequency was assayed by CFSE dilution, showing that a significantly higher proliferation rate of cow's milk-specific CD4⁺T cells in casein-sensitized CMA patients (30.0 \pm 5.8%) than that of non-Cas CMA group(20.7 \pm 5.5%) and atopic controls(7.7 \pm 1.9%) (Figure3a). In addition, upon in-vitro stimulation with cow's milk protein, although there were significant differences in Th2 (IL-4, IL-13), Th17 (IL-17, IL-23) and Treg (IL-10 and TGF- β) among these 3 subgroups, a intra-group comparison revealed that the variations in levels of IL-4 (4.1 \pm 13.6 vs 64.7 \pm 12.3 pg/ml, $P < 0.05$), IL-13(48.7 \pm 10.3 vs 38.7 \pm 9.4 pg/ml, $P < 0.05$) and IL-10

(22.1 ± 5.3 vs 26.8 ± 4.8 pg/ml, $P < 0.05$) between casein-sensitized CMA patients and those without casein sensitization remained statistically different (Figure 3b).

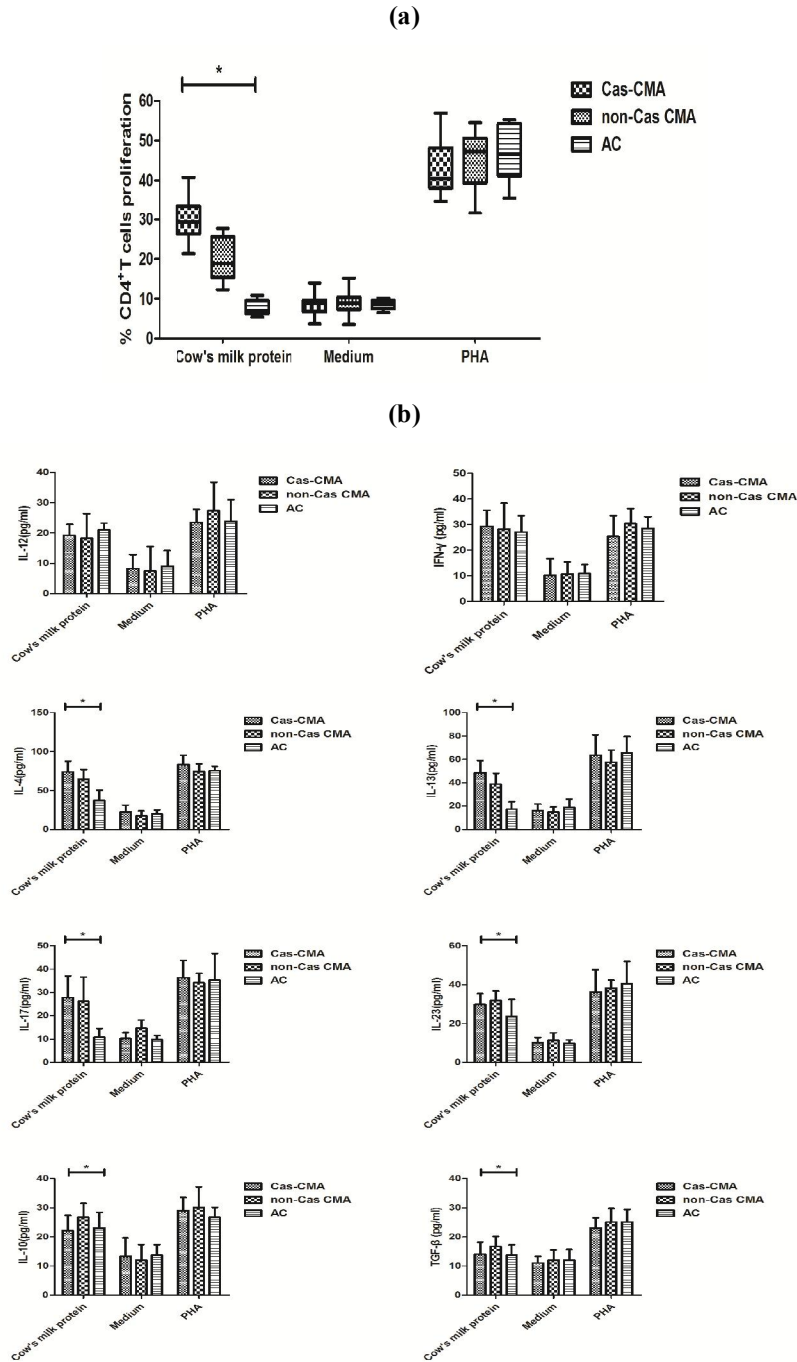


Figure 3: Analysis of T cell proliferating status and cytokines production among CMA patients with casein sensitization(Cas-CMA), CMA patients without casein sensitization(non-Cas CMA), and atopic controls(AC) in vitro stimulation with cow milk protein.

Note: PBMCs were isolated from heparinized venous blood by density-gradient separation. PBMCs (2×10^6 cells/mL) were cultured in vitro for 7 days at 37 °C in 5% CO₂ with medium alone as negative controls, or 200 µg/mL mixture of purified alpha-, beta-, and kappa-caseins, or with 4 µg/mL phytohemagglutinin (PHA) as positive control.

(a) T cell proliferation rate were measured by a CFSE-based flow cytometry method.

(b) The release of T-related cytokines were determined by a custom designed MILLIPLEX MAP kit (Millipore Corporation, Bilerica, MA, USA). **P* value <0.05 considered significant.

Discussion

Cow's milk is the first source of antigens encountered in infancy. The overall prognosis of cow's milk allergy is favorable showing that the majority of infants with CMA could outgrow their allergy, however, there was still a small percentage of children retaining their persistent sensitivity to cow's milk into the second decade [16]. Casein is the most most abundant protein components in cow's milk, approximately accounting for 80% of total milk proteins. A cross-sectional observation study in Taiwanese population revealed that the frequency of sensitization to casein was found up to 40% of their study population [17], while were similar to those obtained from western studies [18,19]. Moreover, due to its heat stability and unique epitope structure casein protein is thought to be the most potent allergenic proteins in cow's milk, with an evidence showing that heating milk at 120 °C for 15 min did not get rid of the antigenicity of bovine casein absolutely [20]. In real-world clinical practice, children with a persistent high casein-specific IgE are prone to exhibit a more severe phenotype of CMA and a delay of intolerance to cow's milk [21]. Comparing with sIgE to whole cow's milk, monitoring the level of casein-sIgE can serve as a bio-marker to predict clinical reactivity to cow's milk with a higher sensitivity and specificity [22].

The aim of the present study was to compare the difference in T cells-mediated responses among CMA patients with or without casein sensitization. Our results showed that T cell isolated from casein-sensitized CMA patients polarized a Th2-type cytokines profiles, which featured by higher serum levels of IL-4 and IL-13 in comparison with those CMA patient without casein sensitization or atopic controls. Our results is also in line with prior studies demonstrated that casein-specific T cell clones derived from persistent CMA patients present a Th2-high signature with a higher release of IL-4 and IL-5, however, T cell responses isolated from non-allergic subjects polarized a Th1-type response [23]. Accordingly, our finding suggested that an altered T cell-mediated immunologic responses towards different protein components in cow's milk, especially for casein allergen component, might be one of the determinant factors of the outcome for cow's milk allergy.

Another Foxp3⁺regulatory T subsets (Tregs) with suppressive function were analyzed in this study. We firstly examined whether there were differences in expression level of Foxp3 among study subjects, a key transcription factor for Tregs differentiation. Our results showed that the expression levels of Foxp3 mRNA in peripheral blood mononuclear cells (PBMCs) from patients with cow's milk allergy were significantly decreased than that in atopic controls, moreover, the variation in Foxp3 gene expression in casein-sensitized CMA patients were also found statistically different from those CMA patients without casein sensitization. Although we observed a slightly

decreased level of FoxP3 expression in CD4⁺T cells isolated from CMA patients with casein sensitization when comparing with those without casein sensitization, the intra-group difference between these 2 groups was not statistically significant. However, the opposite results were also obtained showing an increased number of CD4⁺Foxp3⁺Tregs was found in both cow's milk allergy group and cow's milk tolerant group after in-vitro stimulation with casein [24]. We assumed one possible reason for this discrepancy might be explained by the different clinical features of enrolled CMA subjects. Instead of basing on OFC as enrollment criteria in other previous studies, the diagnosis of CMA in present study was mainly based on the convincing clinical history after cow's milk consumption, the presentation of clinical symptom might include not only gastrointestinal symptoms, but also any of following urticaria, angioedema, wheezing, or a combination of above. Therefore, the differences in clinical characteristic of enrolled subjects might be one of possible reasons for the variations of Treg-mediated responses.

Next, the level of Treg-related cytokines in serum, such as IL-10 and TGF- β , were compared among different subgroups as well. Our data showed that when comparing with CMA patients without casein sensitization there was an obviously reduced serum level of IL-10 in casein-sensitized CMA patients. In addition, after in-vitro stimulation with commercial cow's milk protein mixture, a similar result was obtained showing that the level of IL-10 was statistically lower in CMA patients with casein sensitization than those without it. IL-10 is a potent anti-inflammatory cytokine produced mainly by monocytes, lymphocytes and other cells, including macrophages, B cells, mast cells, and eosinophils [25]. IL-10 are thought to have played an important roles in maintaining intestinal homeostasis and oral tolerance induction, based on the mechanism of oral tolerance involves the deletion or anergy of T cells reactive with specific antigens associated with the expanding population of IL-10- producing regulatory T cells (Tr1) [26]. Moreover, there were much evidence reporting some single nucleotide polymorphisms (SNPs) located in IL-10 gene promoter were associated with the food allergy in Taiwanese or Japanese population, indicating that the potentially critical role of IL-10 in the pathogenesis of food allergy [27]. Moreover, Scott-Taylor et al. [28] reported that children with FA also had reduced numbers of IL-10-producing T cells in the circulation, moreover, the proportions of T cells of food sensitized patients spontaneously secreting IL-10 without antigen stimulation were lower than non-atopic controls, which was consistent with our present study.

In conclusion, the current study implicates a distinctive T cell mediated immune response among CMA patient with different sensitized component. However, inevitably, there are still some limitations in present study design. First, we should conduct standardized OFC as the enrollment criteria of CMA patients. Second, a longitudinal monitoring of casein-specific T cell response with growth status of CMA children should be further conducted, which could be helpful to provide more evidence to understand the molecular mechanism of casein-specific T cells during the CMA process.

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Conflict of Interest Statement

We declare that we have no conflict of interest.

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