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Th2A and Th17 cell frequencies and regulatory markers as follow-up biomarker candidates for successful multifood oral immunotherapy

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To the Editor

In the context of food oral immunotherapy (OIT) the identification of biomarkers that can replace current standard burdensome and risky double-blind placebo-controlled food challenges (DBPCFC), which are used pre- and post-therapy¹, would be of great value in clinical settings. With this goal in mind, we undertook a comprehensive immune response analysis on blood cells obtained from a sub-group of multi-allergic participants in a clinical multi-OIT study (ClinicalTrials.gov number, NCT02626611)² to evaluate potential biomarker candidates. In the open-label phase of the multi-OIT study (n = 70, age 5– 22 years), participants received omalizumab (weeks 1–16) and multi-OIT (2–5 allergens; 1 g each; weeks 8-30), after which they were tested by food challenge (week 30). Subsequently, 60 eligible participants (excluding 10 drop-outs) were randomized 1:1:1 to receive in a blind manner either 0 mg, 300 mg, or 1 g of food allergens (weeks 30–36). These participants were then tested again by food challenge at week 36.

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Author contribution

KN, SCL and SC ran the trial and prepared samples. KN and LM designed the biomarker research study. SL performed the experiments. SL, KN and LM analysed the data. KN and LM supervised the research. KN and LM wrote the paper.

Conflict of interest

Co-authors (SL and LM) are employees at Stallergenes Greer.

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We analysed systemic changes in immune parameters, such as T helper (Th) cell and innate lymphoid cell (ILC) frequencies as well as dendritic cell (DC)-associated markers, occurring after 30-week multi-OIT, in a subset of 42 out of the 60 eligible patients. The demographics and baseline characteristics of patients are provided in Supplementary Table S1. Peripheral blood mononuclear cells (PBMCs) were stained with fluorescently labeled antibodies as described in Supplementary Methods according to the gating strategy shown in Supplementary Figure S1. We observed lower frequencies of both Th2A (p<0.05) and Th17 (p=0.0504) subsets in PBMCs at week 30 when compared to baseline (Figure 1.A and B, left panels). For the sake of simplicity, we divided ages in two groups instead of three groups since only 3 participants were over 15-year-old. Noticeably, the decrease of Th2A cell frequencies was statistically significant (p<0.05) only for patients aged 10 and over, although a positive trend was also detected in younger patients (Figure 1.A, **right panel**). Additionally, multi-OIT markedly (p<0.05) decreased Th17 cell frequencies in patient aged 10 and over, but no differences were seen for patients under 10-year-old. (Figure 1.B, right **panel**). To further explore cellular immune responses, we performed a detailed analysis of changes in innate immune cells by flow cytometry. However, no alterations were observed between groups in terms of cell frequencies for both type 1, 2 and 3 innate lymphoid as well as for DC subsets (data not shown). In addition, we monitored PBMCs for expression of markers associated with type 1 (*i.e. MX1*), type 2 (*i.e. GATA3* and *CD141*), or regulatory (i.e. C1q, STAB1 and FcyRIIIa) DCs previously identified to correlate with clinical efficacy of allergen immunotherapy (AIT) ^{3, 4} (See Supplementary Methods). We did not observe any changes in DC1 or DC2-associated markers. In contrast, 2 out of 3 DCreg markers exhibited statistical differences before and after multi-OIT. The expression of STAB1 was significantly (p<0.05) upregulated in PBMCs from multi-OIT patients at week 30 when compared to baseline (Figure 1.C, left panel) particularly for patients aged 10 and over (Figure 1.C, right **panel**). Multi-OIT did not alter the expression of $Fc\gamma RIIIa$ in patients' blood when considering the whole subset (Figure 1.D, left panel). However, $F_{C\gamma}RIII_a$ expression was significantly (p<0.05) increased in PBMCs from patients under 10 and correlated with a decrease of Th17 cell frequencies (Supplementary Figure S2; center panel) but was unchanged in patients older than 10 years of age (Figure 1.D, right panel).

At week 36, immune changes were also analysed in sub-groups of patients receiving either 0 mg (n=12), 300 mg (n=17), or 1 g (n=13) as aforementioned. No differences in terms of Th2A or Th17 cell frequencies (Supplementary Figure S3) or STAB1 expression (Figure 2.A, **upper panel**) were observed between baseline and week 36 within the 3 groups (0 mg, 300 mg, 1 g), likely due to a limited number of patients. It is thus difficult to conclude on the value of following such markers after cessation of OIT. Only on pooling data from the 300 mg and 1 g treated groups, *STAB1* was observed to be significantly (p<0.05) upregulated following a 36-week multi OIT, particularly in PBMCs from patients aged 10 and over (Figure 2.A, **lower panel**). Strikingly, *Fc* γ *RIIIa* expression was significantly (p<0.01) increased in PBMCs from patients receiving 1 g of food allergens up to week 36 (Figure 2.B, **upper panel**). On pooling data from the 300 mg and 1 g treated groups, we confirmed that this marker is significantly increased in PBMCs irrespective of age groups at W36 (but not at W30), suggesting that this DCreg marker is regulated in a time- and age-dependent fashion (Figure 2.B, **lower panel**).

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These results establish for the first time that omalizumab-facilitated multi-OIT induces changes in immune polarization-based readouts. Multi-OIT promoted a decrease in Th2A and Th17 cell frequencies while increasing regulatory markers in blood, particularly evidenced for patients aged 10 and over for which desensitization has been successful. Such results will need to be confirmed with a larger cohort of patients in DBPC clinical trials since we acknowledge the limits of our study due to a low number of participants. Additionally, long-term therapy should be considered as it may enable sustained effect and allow cessation of immunotherapy². Our findings are fully aligned with previous reports monitoring follow-up candidate biomarkers in AIT studies for respiratory allergies^{3, 4}. Although a surrogate biomarker of AIT efficacy has not been validated so far, immune changes documented in peripheral blood or mucosal tissues of patients include the downregulation of allergen-specific Th2 cells ^{5, 6}. On the other hand, a limited number of studies have analyzed a change in Th17 polarization with mixed conclusions ^{7, 8}. We also confirmed the interest of both STAB1 and Fc yRIIIa as hallmark DCreg markers in the mechanism of action of immunotherapy ^{3, 4}. This is consistent with observations that *STAB1* is expressed by tolerogenic DCs/macrophages and that it contributes to fetal implantation in the human decidua ^{4, 9}. In contrast, the identification of FcyRIIIA as a DCreg marker was less expected since this receptor is more involved in inflammatory mechanisms ⁴.

Altogether, our findings pave the way for further understanding of the mechanisms involved in food immunotherapy. Positive outcomes on systemic immune parameters, as observed in this study, confirm the value of monitoring (particularly in PBMCs from pediatric patients), Th2/Th17 cell frequencies and DCreg markers as potential replacements for oral food challenges. We also suggest that successful multi-OIT involve different immune pathways according to the maturation level of the immune systems during infancy.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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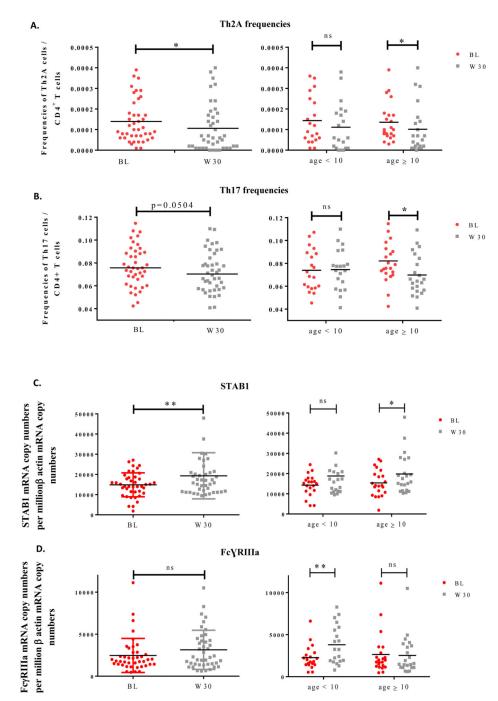


Figure 1: Th2A and Th17 cell frequencies decreased while the regulatory marker STAB1 and FcyRIIIa increased after multi OIT.

PBMCs from patients receiving multi OIT (n=42) were analysed at baseline (BL) and week 30 (W30) for Th2A (A) and Th17 (B) cell frequencies by flow cytometry and for STAB1 (C) and Fc γ RIIIa (D) gene expression by quantitative PCR. Age-related factors were analysed for patients under 10 and aged 10 and over. Wilcoxon tests were used for statistical analysis. *P<0.05; **p<0.01; ns=non-significant.

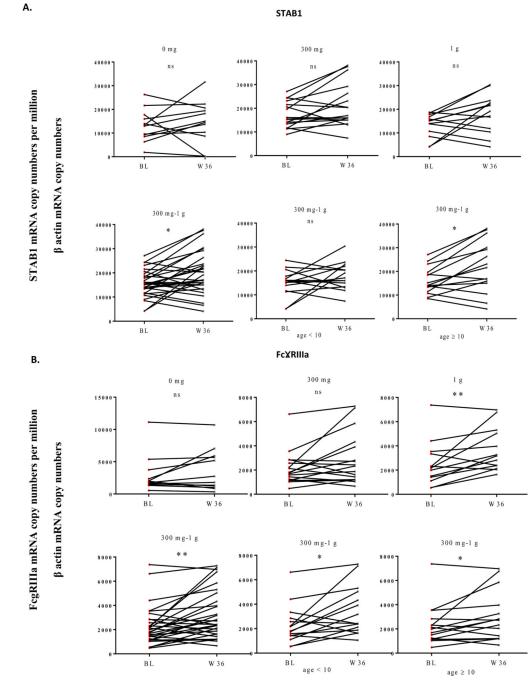


Figure 2: Regulatory markers STAB1 and FcγRIIIa are confirmed to be upregulated in PBMCs from patients after extension of multi OIT.

At week 30 (W30), patients receiving OIT were randomised in 3 groups: patients interrupting the treatment (0 g; n=12), patients continuing OIT with 300 mg (n=17) or 1 g (n=13) until week 36 (W36). PBMCs from patients were analysed at baseline (BL) and W36 for STAB1 (**A**) and FcyRIIIa (**B**) gene expression by quantitative PCR. Age related factor were analysed for patients under 10 and aged 10 and over. Wilcoxon tests were used for statistical analysis. *P<0.05; **p<0.01; ns=non-significant.

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