

---

# Specific Peanut Epitopes as a Biomarker for Desensitization During Epicutaneous Immunotherapy

**David M Fleischer, MD**

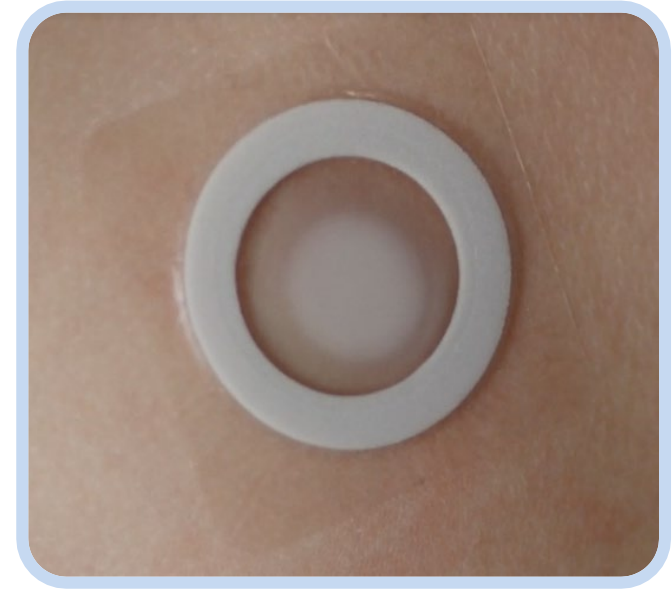
Professor of Pediatrics and Section Head  
Children's Hospital Colorado

University of Colorado School of Medicine, Aurora, CO, USA

**On behalf of:** Dianne E Campbell, Paul Kearney, Bob Getts, Todd D Green, Hugh A Sampson

# Background

- Epicutaneous immunotherapy (EPIT) is currently under investigation for the treatment of peanut allergy
  - **DBV712 250 µg** is a single, daily-dose patch applied to the back and is dosed at 250 µg (**~1/1000 of a peanut**)<sup>1-3</sup>
- Existing biomarkers such as total peanut-IgE and SPT used in the diagnosis of peanut allergy are not adequate for quantifying desensitization during immunotherapy
- Total peanut-IgG4 during EPIT is not highly correlated with treatment response during immunotherapy
- Measuring IgE and IgG4 reactivity to **epitopes** (smaller fragments of allergen to which an allergic individual can produce an antibody) from peanut protein allergens may help to quantify desensitization during EPIT



SPT=skin prick test.

1. Sampson HA, et al. *JAMA*. 2017;318:1798-1809. 2. Tilles SA, et al. *Ann Allergy Asthma Immunol*. 2018;121:145-149. 3. Parrish CP. *Am J Manag Care*. 2018;24:S419-S427.

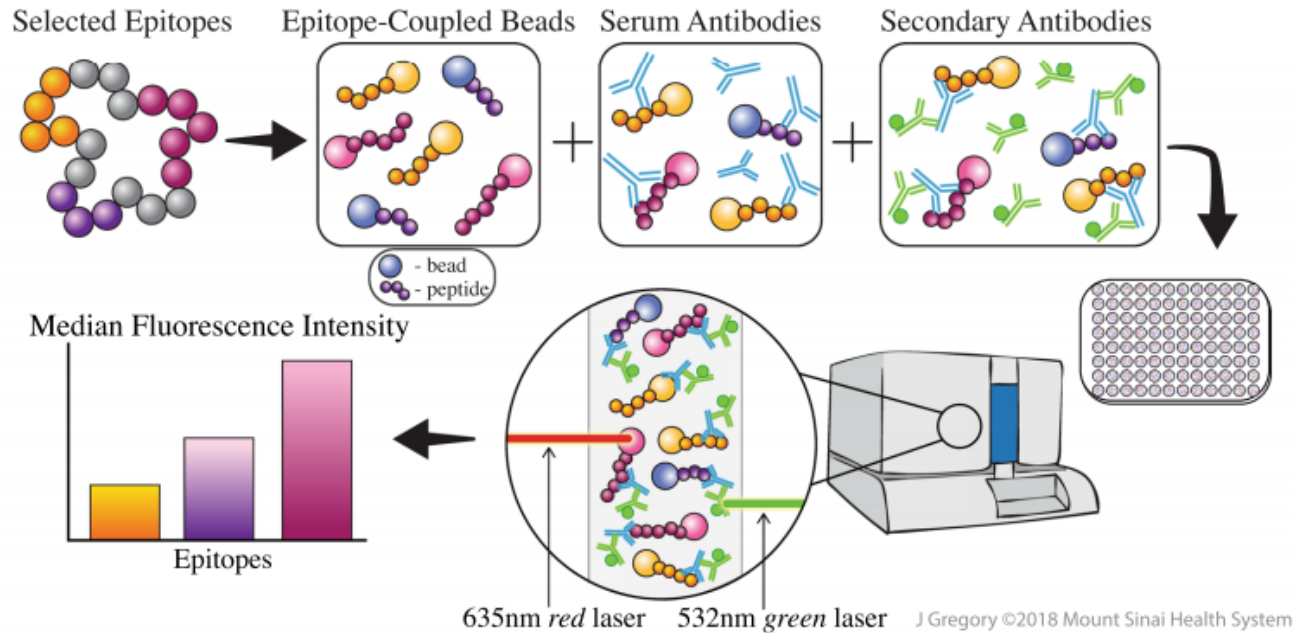
# Aims and Objectives

---

- Assess performance of the **Bead-Based Epitope Assay** (BBEA), which enables simultaneous quantification of antibodies recognizing sequential linear protein epitopes, in predicting desensitization during EPIT
- The specific **objectives** of the study were to:
  - Analyze the samples from subjects who participated in PEPITES (Phase 3 DBRCT of DBV712 250 µg in peanut allergic children) for IgE and IgG4 reactivity to Ara h 1, Ara h 2, and Ara h 3 peanut allergen epitopes using the BBEA
  - Derive predictors of desensitization progress over time as well as predictors of desensitization above a peanut protein eliciting dose threshold (>300 mg) following 12 months of treatment with DBV712 250 µg

# Methods: BBEA Technology<sup>1</sup>

- A **Bead-Based Epitope Assay** (BBEA) platform was used to monitor the reactivity of **IgE and IgG4** in subjects' serum to 64 linear **epitopes** from Ara h 1, Ara h 2, and Ara h 3



- The BBEA methodology enables simultaneous quantification of antibodies binding to sequential epitopes
- Epitopes are covalently coupled to unique fluorescent microspheres (Luminex)
- Epitope-labeled beads are mixed to form a master library
- Patient plasma and a secondary fluorophore-labeled antibody are then incubated with the beads
- The Luminex instrument uses dual-lasers for quantification (red for beads, green for secondary antibodies)
- For each epitope, the signal is quantified as a median fluorescence intensity (MFI)

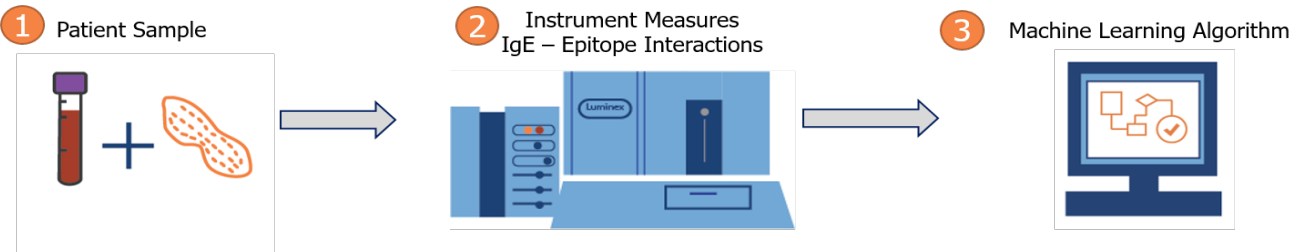


Figure adapted from Suprun et al.<sup>1</sup> (<http://creativecommons.org/licenses/by/4.0/>)  
1. Suprun M, et al. *Sci Rep.* 2019;9:18425. doi.org/10.1038/s41598-019-54868-7.

# Methods: Subjects and Samples

---

## Subjects

- **89 peanut-allergic subjects** who participated in the 12-month, randomized, double-blind, placebo-controlled **PEPITES** study<sup>1</sup>
  - Subjects received active treatment with DBV712 250 µg (n=61) or placebo (n=28)

## Samples

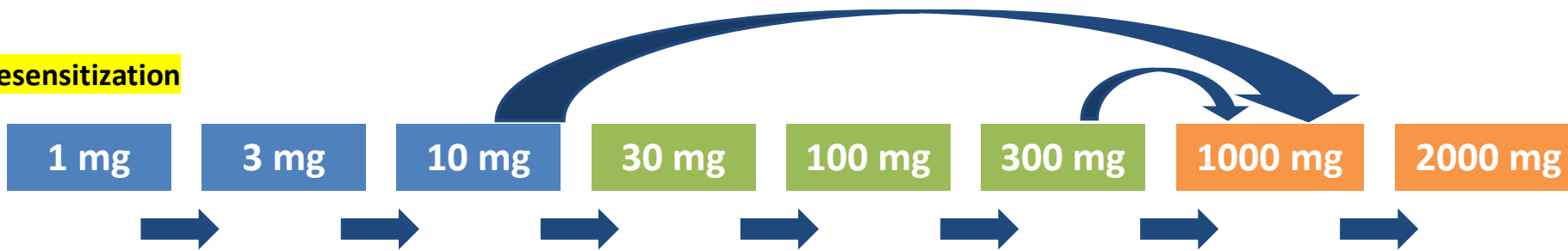
- Serum samples were analyzed using the BBEA at 0, 3, 6, and 12 months for IgE and IgG4 reactivity to 64 linear epitopes from Ara h 1, Ara h 2, and Ara h 3 (esIgE and esIgG4)
  - The BBEA method was applied under SOPs to all subjects in triplicate and randomized across plates
  - Raw data was processed: noise removal, log normalized, triplicates merged
  - Analysis was performed using linear regression models
  - Accuracy, sensitivity, specificity, and AUC were generated
- Serum samples were analyzed for total peanut-specific IgE and IgG4

# Methods: Endpoints

## Endpoints

- **Threshold desensitization** was defined as an eliciting dose (ED) >300 mg peanut protein at Month 12 (ie,  $\geq 1000$  mg) irrespective of baseline ED (all  $\leq 300$  mg)
- **Progressive desensitization** was defined as ED threshold increased by at least 1 dose from baseline to Month 12

### Threshold desensitization



### Progressive desensitization

# Results: Epitope Mapping – Progressive and Threshold Desensitization Predictors

## Threshold Desensitization (ED $\geq$ 1000 mg)

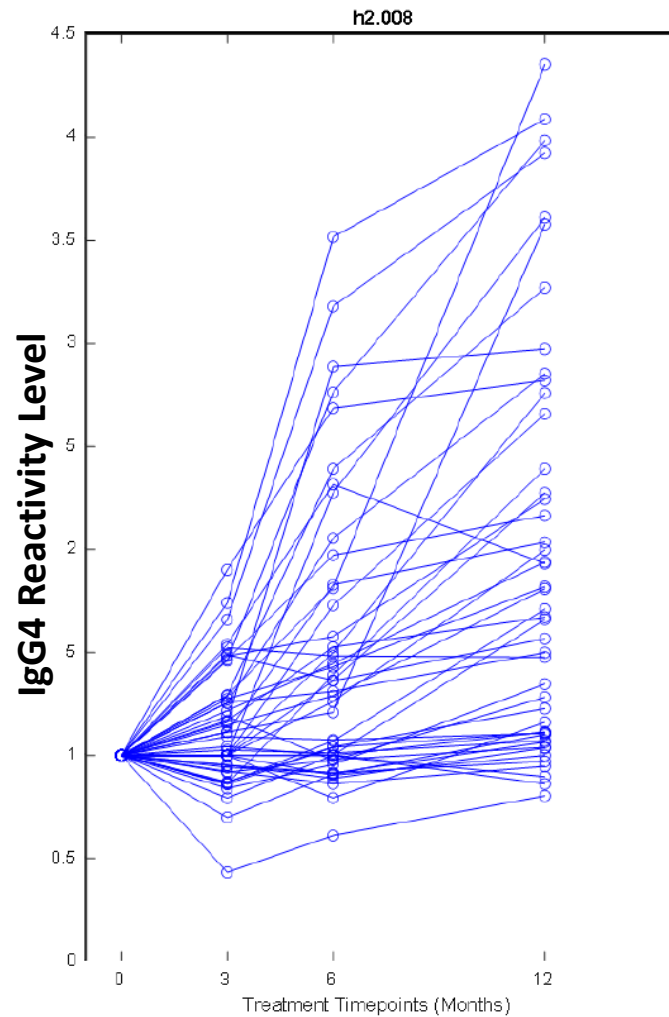
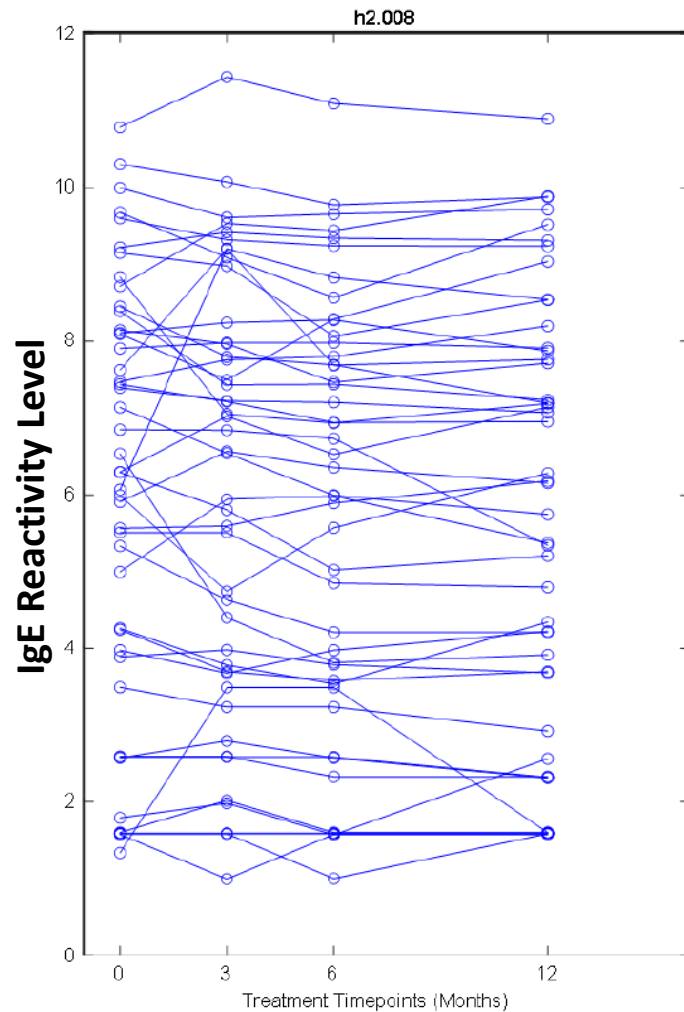
	12 Months
<b>Best epitopes (IgG4)</b>	<i>h2.010</i> <i>h3.102</i> <i>h1.029</i> <i>h1.090</i>
<b>AUC</b>	<b>0.97</b>
<b>% accuracy (95% CI)</b>	93% (90, 98)

## Progressive Desensitization (Improvement in ED)

	12 Months
<b>Best epitopes (IgE)</b>	h1.041 h1.022 h2.008
<b>AUC</b>	<b>0.92</b>
<b>% accuracy (95% CI)</b>	91% (83, 93)

- Addition of baseline subject characteristics or AE rate during treatment did not improve performance in either analysis

# Results: esIgE and esIgG4 to h2.008 Reactivity for All Subjects



- For epitopes such as h2.008, treatment increased IgG4 reactivity to epitopes over time
- Similar behavior was seen for h1.029



# Results: Threshold Desensitization

---

## Threshold Desensitization at Month 12 – 4 Epitope Linear Regression Model (IgG4)

	Coefficient	SE	tStats	P value
<b>(Intercept)</b>	-0.535	1.593	-0.336	0.737
<b><i>h2.010</i></b>	6.357	3.312	1.919	0.055
<b><i>h3.102</i></b>	-5.910	2.970	-2.033	0.042
<b><i>h1.029</i></b>	-7.037	2.667	-2.639	0.008
<b><i>h1.090</i></b>	6.339	2.579	2.458	0.014
Chi <sup>2</sup> -statistic vs constant model: 41.9, P value =1.77e-08, <b>AUC=0.97</b>				

# Results: Progressive Desensitization at Month 12

## Progressive Desensitization at Month 12 – 3 Epitope Linear Regression Model (IgE)

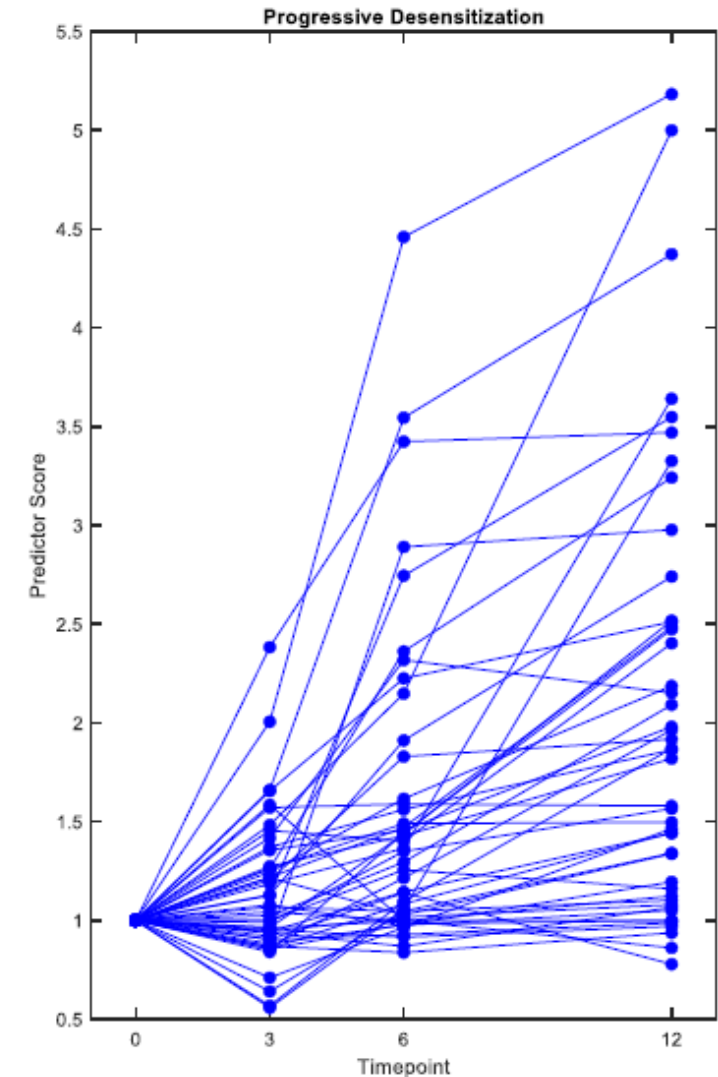
	Coefficient	SE	tStats	P value
<b>(Intercept)</b>	6.932	2.361	2.936	0.0033
<b>h1.041</b>	-0.634	0.279	-2.271	0.0232
<b>h1.022</b>	0.585	0.237	2.463	0.0138
<b>h2.008</b>	-0.423	0.226	-1.872	0.0612
Chi <sup>2</sup> -statistic vs constant model: 23.2, <i>P</i> =3.71e-05, <b>AUC=0.92</b>				

# Results: Progressive Desensitization Over Time

## Progressive Desensitization

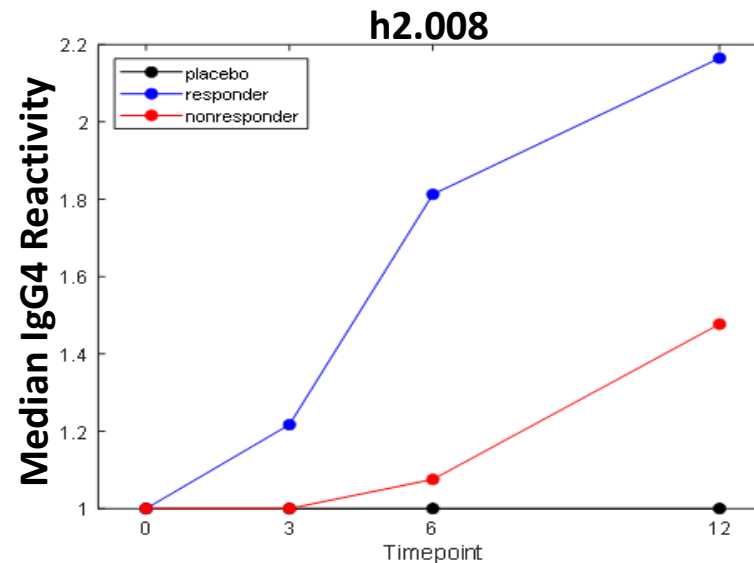
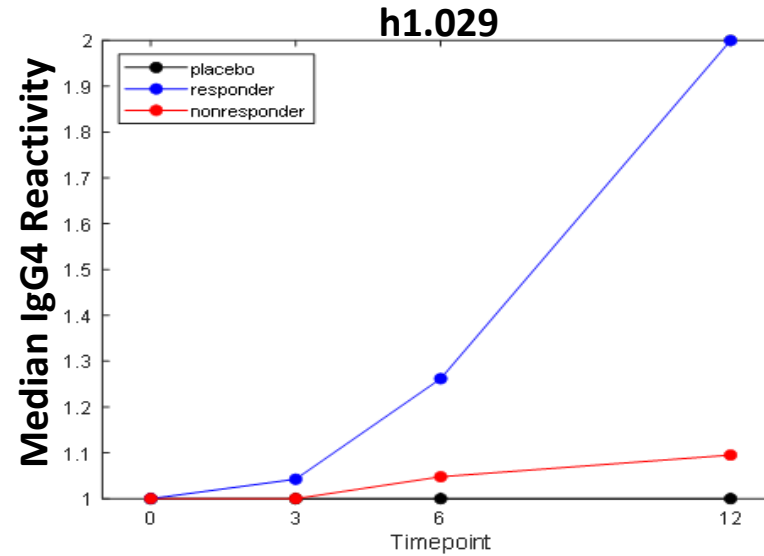
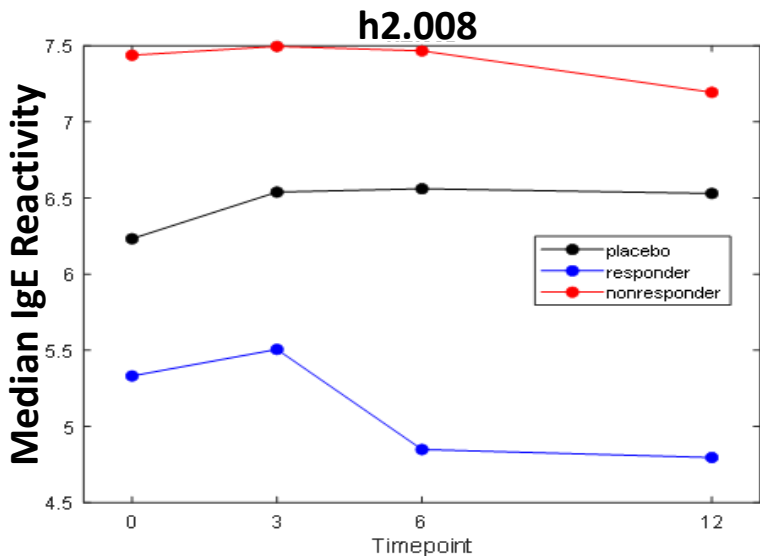
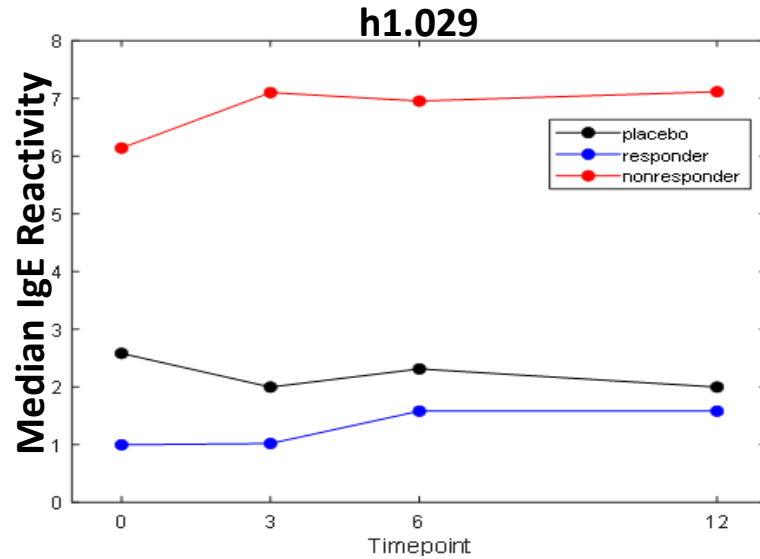
	3 Months	6 Months	12 Months
<b>Best epitopes (IgE)</b>	h1.041 h1.022 h2.008	h1.041 h1.022 h2.008	h1.041 h1.022 h2.008
<b>AUC</b>	0.90	0.91	0.92
<b>% accuracy</b>	89%	89%	91%

AUC=area under the curve.



Scores above 1 indicate progressive desensitization over time.

# Results: Responders vs Non-Responders vs Placebo – Epitopes h1.029 and h2.008



- IgG4 reactivity to each epitope differentiated between placebo, responders, and non-responders with a markedly different trajectory over the 12 months by group
- Faster increases in IgG4 reactivity to h1.029 and h2.008 was associated with treatment response
- IgE reactivity to h1.029 and h2.008 remained stable or slightly declined over 12 months of treatment

# Results: Comparison of esIgE/IgG4 to Total Peanut-sIgE/sIgG4

Total peanut-sIgE and sIgG4/sIgE were demonstrated to be poorer predictors of progressive desensitization, with AUCs of 68% and 64%, respectively, at 12 months

**sIgE Predictor of Progression Desensitization**

	Coefficient	SE	tStats	P value
<b>(Intercept)</b>	1.4447	0.45808	3.1538	0.0016113
<b>x1 [h1.041]</b>	-0.0012408	0.0010524	-1.1789	0.23842
Chi <sup>2</sup> -statistic vs constant model: 1.36, <i>P</i> =0.244, <b>AUC=0.68382</b>				

**sIgE and sIgG4 Predictor of Progression Desensitization**

	Coefficient	SE	tStats	P value
<b>(Intercept)</b>	1.2014	0.53651	2.2392	0.025142
<b>x1 [h1.041]</b>	-0.0016422	0.0011868	-1.3837	0.16644
<b>x2 [h1.022]</b>	0.061838	0.076652	0.80674	0.41981
Chi <sup>2</sup> -statistic vs constant model: 2.14, <i>P</i> =0.344, <b>AUC=0.64216</b>				

# Summary and Future Directions

---

## **BBEA may be a highly accurate tool for monitoring desensitization during EPIT**

- Highly accurate models for predicting both **threshold** and **progressive desensitization** were able to be built using 3, 4, and 5 epitope (eslgE and eslgG4) algorithms
- Overall peanut-eslgE reactivity did not change significantly with treatment over time, while peanut-eslgG4 reactivity increased
- Faster increases in eslgG4 reactivity to h1.029 and h2.008 were associated with treatment response
- Lower eslgE reactivity to h1.029 and h2.008 was associated with response to treatment
- Total peanut-slgE and/or slgG4 were shown to be relatively poor predictors of progressive desensitization in this population
- Analyses are ongoing to validate these findings in a larger sample and to examine predictors for desensitization beyond 12 months

# Acknowledgements

---

**Dianne E Campbell<sup>1,2</sup>, Paul Kearney<sup>3</sup>, Bob Getts<sup>3</sup>, Todd D Green<sup>2,4</sup>,  
David M Fleischer<sup>5</sup>, Hugh A Sampson<sup>2,6</sup>**

1. Children's Hospital at Westmead, Sydney
2. DBV Technologies SA, Montrouge, France
3. AllerGenis, Hatfield, PA, USA
4. UPMC Children's Hospital of Pittsburgh and University of Pittsburgh School of Medicine, Pittsburgh, PA, USA
5. Children's Hospital Colorado, University of Colorado School of Medicine, Aurora, CO, USA
6. Department of Pediatrics, Allergy and Immunology, Icahn School of Medicine at Mount Sinai, New York, NY, USA

