Comparison of Patients with Eosinophilic Esophagitis with Extremely High and Low Esophageal Eosinophil Counts

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Introduction

Eosinophilic esophagitis (EoE) is an inflammatory disease characterized by eosinophil infiltration into the esophageal mucosa with a peak count of \geq 15 eosinophils per high-powered field (eos/hpf) following endoscopic biopsy. However, the range of esophageal eosinophilia can vary markedly from patient to patient. A key question in the field is to understand the relationship of eosinophil levels with disease features, especially since eosinophil-targeted therapies are now available. Patients with extremely high levels of esophageal eosinophilia have not previously been studied. It is unknown whether these patients exhibit different characteristics compared with EoE patients that have esophageal eosinophilia that is near the threshold of disease diagnosis. Given this, we aimed to establish whether any significant clinical, endoscopic, histologic, or transcriptomic features differ between patients with extremely high levels of esophageal eosinophilia and those with levels near the threshold of disease diagnosis.

Methods

Subject selection

Amongst a registry of patients with EoE, we identified a group of patients with the highest recorded levels of esophageal eosinophilia (eos/hpf >350), referred to as EoE-Hi. We subsequently identified a second group that had relatively low levels of esophageal eosinophilia (15-24 eos/hpf), referred to as EoE-Low. There were 74 patients in the registry with eosinophil counts of 15-24 eos/hpf on a distal esophageal biopsy. A random number generator was used to select the 14 patients that comprised the EoE-Low group.

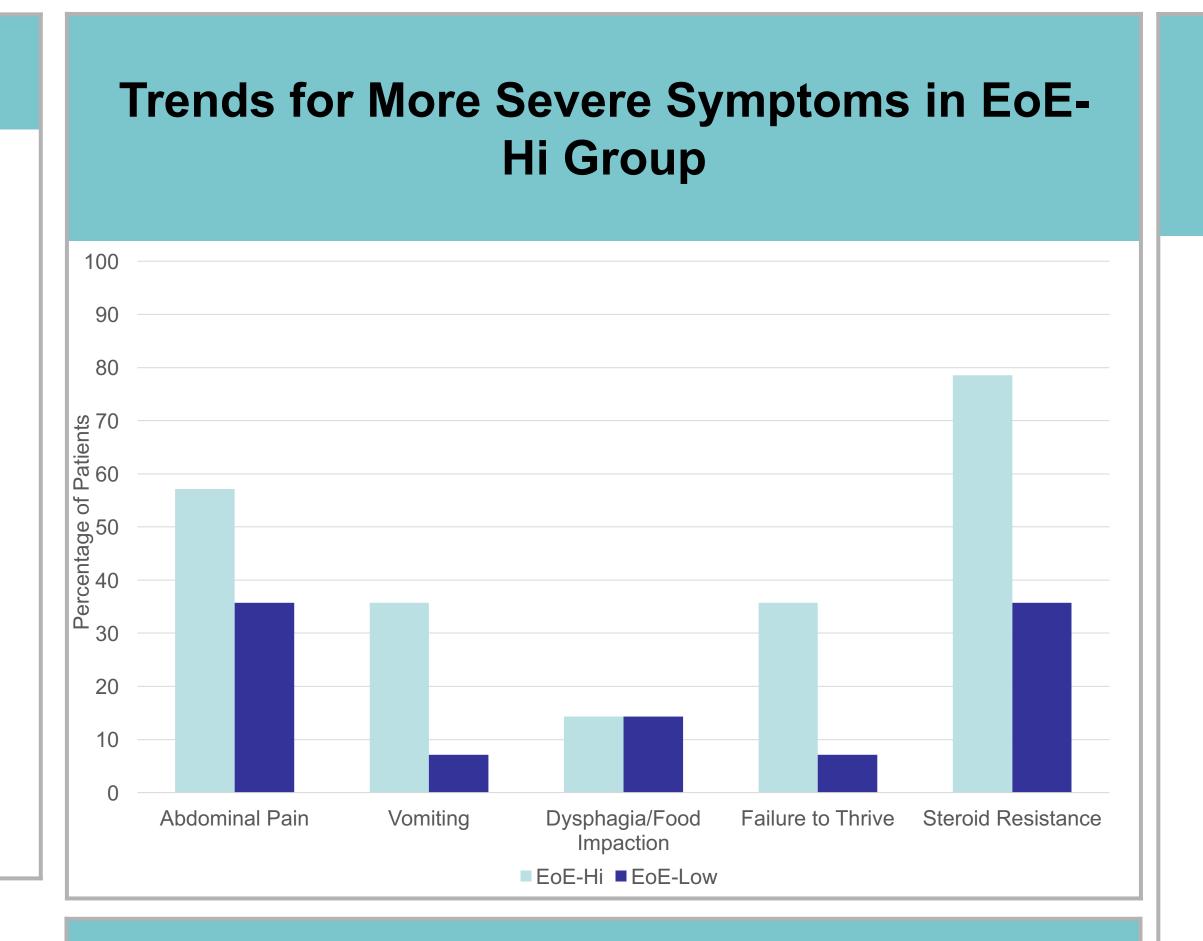
Analysis

Phenotypic and clinical characteristics were gathered on the basis of electronic medical records and detailed questionnaires as part of a research registry. Endoscopic characteristics were assessed on the basis of findings from EGD operative reports. Histologic characteristics were classified on the basis of the histologic scoring system. Molecular analysis was performed using the 96-gene EoE Diagnostic Panel (EDP).

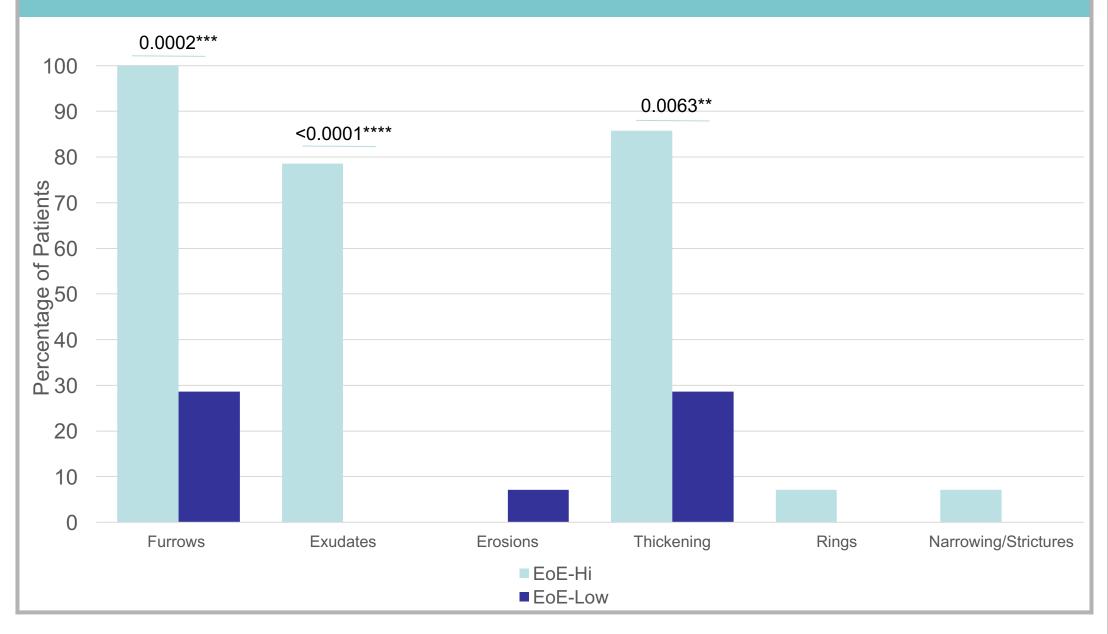
Patients with Extremely High Eosinophil **Counts Have a Unique Phenotype**

	Eos/hpf >350 (n=14)	Eos/hpf 15-24 (n=14)	P-value
Male sex, no (%)	78.6	78.6	>0.9999
Age at Biopsy (y), mean ± SD	13.4 ± 2.7	6.3 ± 0.9	0.0205
Age at Diagnosis (y), mean \pm SD	$7.3 \pm 2.9 \text{ (n=11)}$	4.4 ± 0.8 (n=12)	0.3313
Duration of Disease*(y), mean ±			
SD	$10.9 \pm 0.9 (n=12)$	$6.4 \pm 0.6 (n=12)$	0.0008
Eosinophils/hpf (mean ± SD)	433.3 ± 96.6	18.6 ± 2.9	<0.0001
Absolute Eosinophilia (mean \pm SD)	$686.2 \pm 97.8(n=13)$	481.5 ± 74.7 (n=13)	0.1094
IgE (mean \pm SD)	426.3 ± 150.4 (n=9)	338.8 ± 245.1 (n=5)	0.7521
History of other EGIDs ^{+,} no. (%)	3 (23.1) (n=13)	0	0.2276
Atopic Co-morbidities, no. (%)	14 (100)	9 (64.3)	0.0407
Atopic Dermatitis	10 (71.4)	6 (42.8)	0.2519
Allergic Rhinitis	13 (92.8)	7 (50)	0.0329
Asthma	9 (64.3)	8 (57.1)	>0.9999
Food Allergy	9 (64.3)	6 (42.8)	0.4495
History of Anaphylaxis	5 (35.7)	4 (28.6)	>0.9999

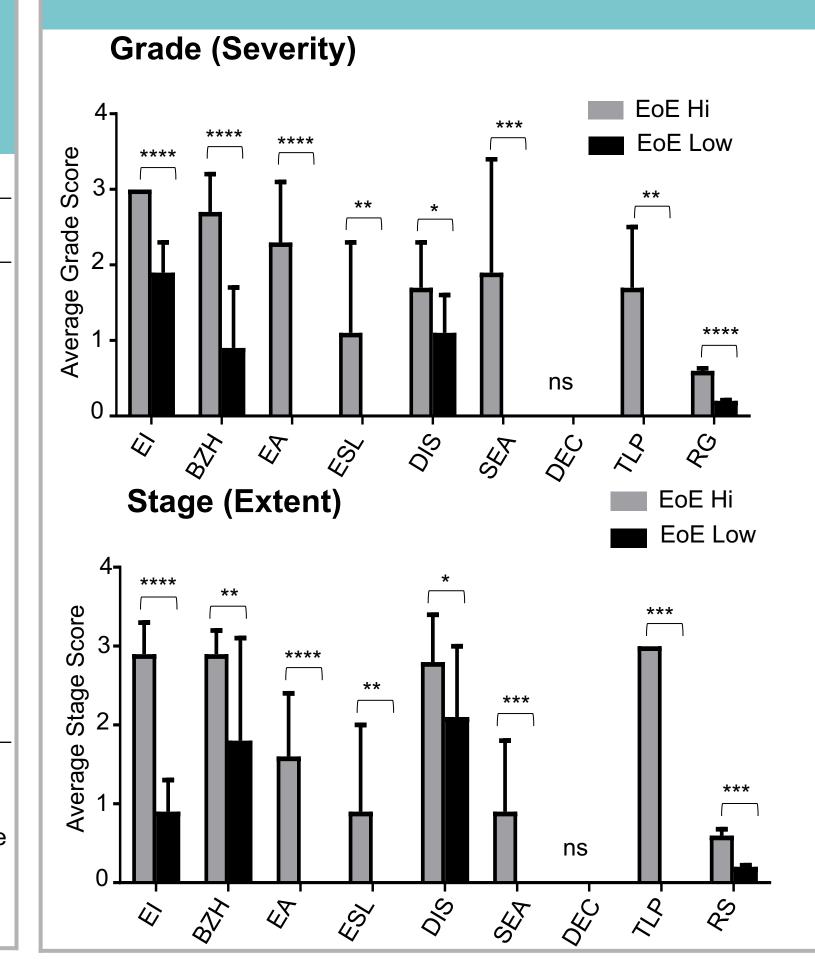
of biopsy. ⁺In addition to a diagnosis of EoE, 2 patients in the EoE-Hi group have diagnoses of eosinophilic colitis and 1 patient has a diagnosis of eosinophilic gastroenteritis.



Patients in the EoE-Hi Group have a **Greater Incidence of Furrows, Exudates** and Thickening on Endoscopy



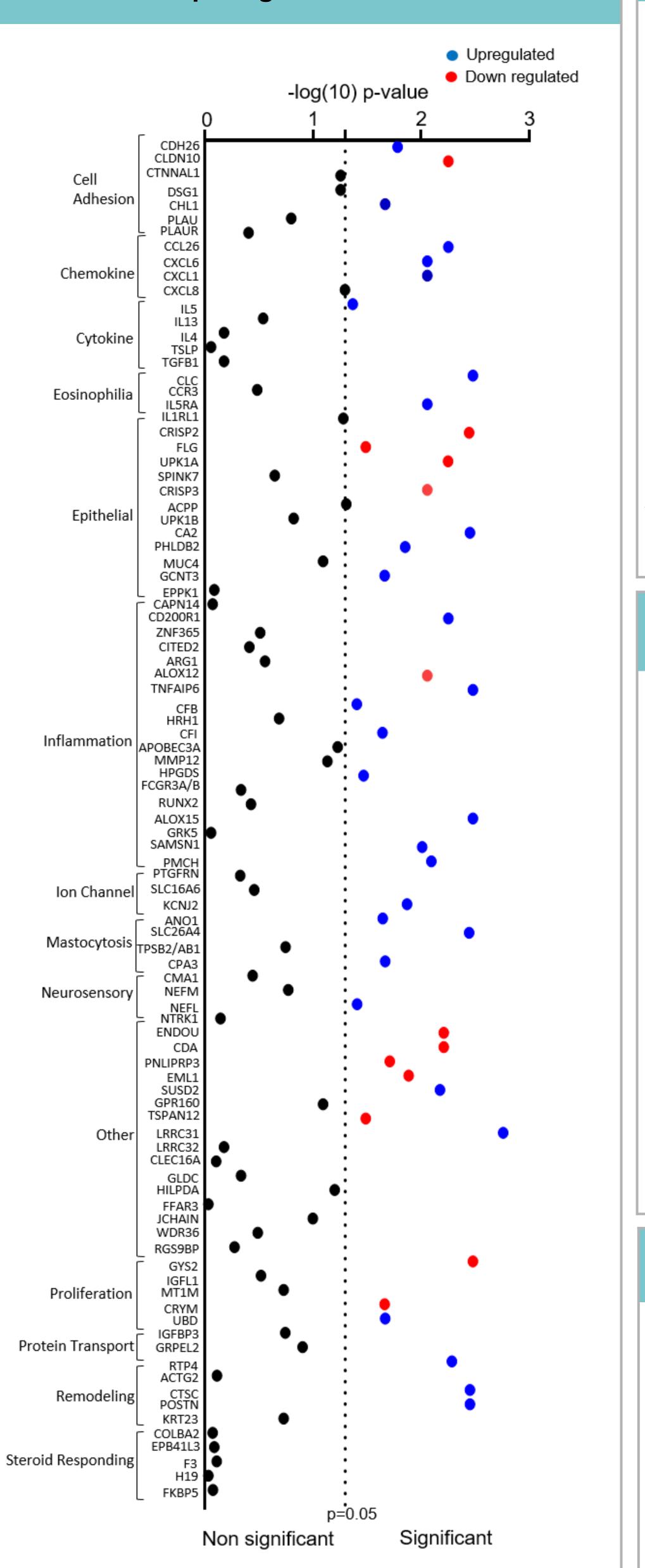
Total Score for Each of Individual Histologic Parameter was Higher (More Severe) in the EoE-Hi Group



EI: eosinophilic infiltration BZH: basal zone hyperplasia EA: eosinophil abscess ESL: eosinophil surface layering DIS: dilated intercellular spaces SEA: surface epithelial alteration DEC: dyskeratotic epithelial cells TLP: thickened lamina propria RG: relative grade RS: relative stage ****p ≤0.0001, ***p ≤0.001, **p ≤0.01, *p ≤0.05; ns, not significant.

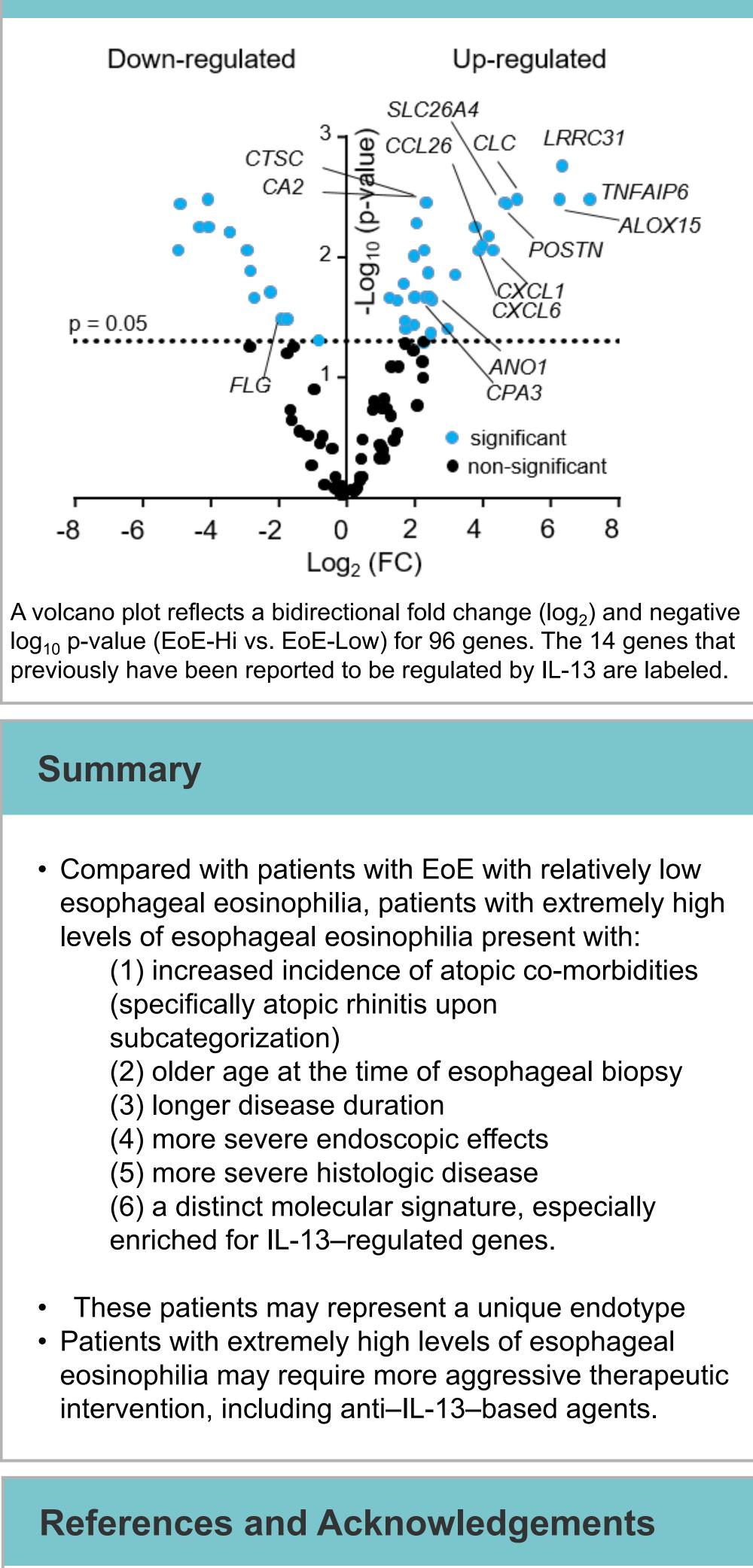
value

Transcriptomic Analysis Revealed 30 Upregulated and 13 Downregulated Genes When Comparing EoE-Hi vs EoE-Low



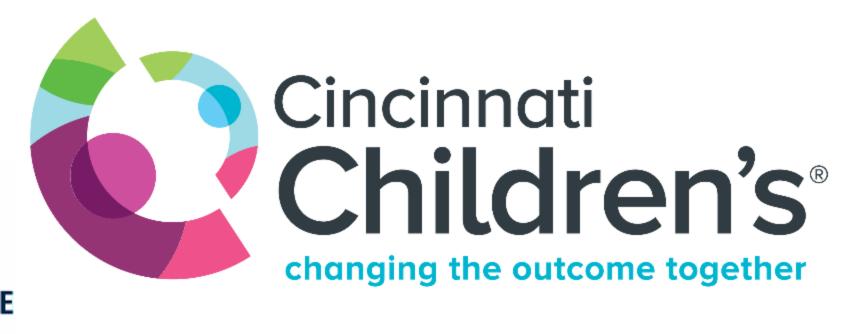
When individual expression of the EoE Diagnostic Panel genes was compared between the 2 groups, there were 43 genes with significantly different expression (p < 0.05, fold change >2.0). This graph displays gene expression by category and negative $\log_{10} p$ -

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IL-13 Related Pathways Associate with **Esophageal Eosinophil Levels**

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