

# Genome-wide association study of Atopic dermatitis in Hong Kong

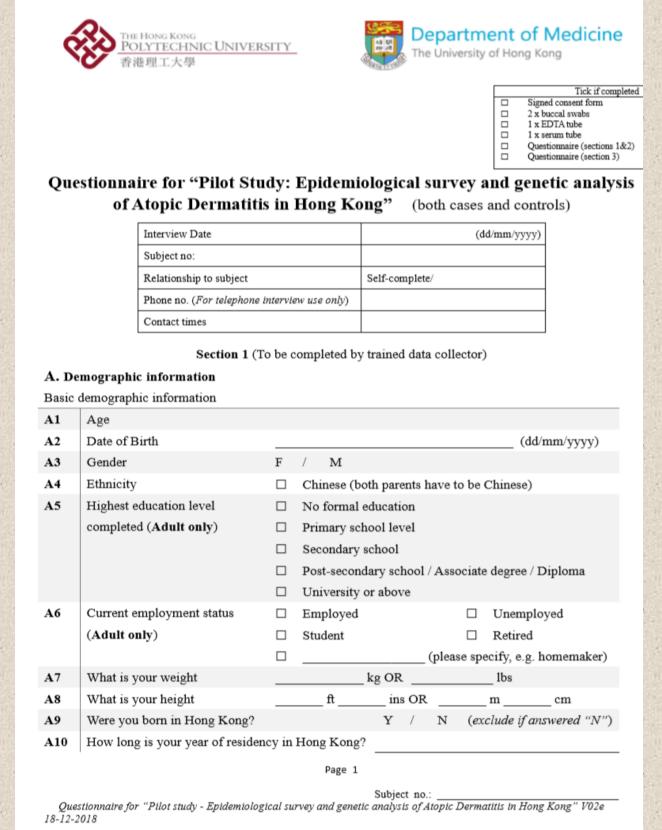
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#### Introduction

Atopic dermatitis (AD) is a common, incurable chronic inflammatory skin disorder that can significantly impact the quality of life of patients. There is an incomplete understanding of the underlying pathogenesis, with no genome-wide association study (GWAS) conducted in Hong Kong, and previous clinical studies were limited to childhood AD. The aim of this study was to carry out a pilot study to search for Hong Kong population specific genetic variants and environmental risk factors across childhood and adult AD.

We hypothesize that differences in subgroups of AD patients (eczema severity, age, IgE, eosinophil count, and age of onset), and different environmental risk factors maybe conferred by different genetic risk factors.

#### Method



43 child and 33 adult AD subjects recruited from Queen Mary 44 Hospital controls and were the Hong recruited from Kong University Polytechnic campus. Demographic and clinical details were collected via questionnaire interview Blood and buccal samples were taken from subjects to undergo total serum IgE, eosinophil count and GWAS for genetic variant profiling.

#### **Discussion**

- Novel child-onset factors:
  - 1. Caesarean section
  - 2. Non-smoking parent(s)
- 6 significant SNPs in association with the duration of AD:
  - 1. Astn2 gene: rs10982272, rs12380425, rs146392873, rs140693157, rs12378473
  - 2. Intron variant rs73123810
- 5 Significant SNPs in association with IgE:
  - 1. Intron variant rs10753681
  - 2. Unknown functions: rs12524729, rs2988685, rs4706269 and rs4707264

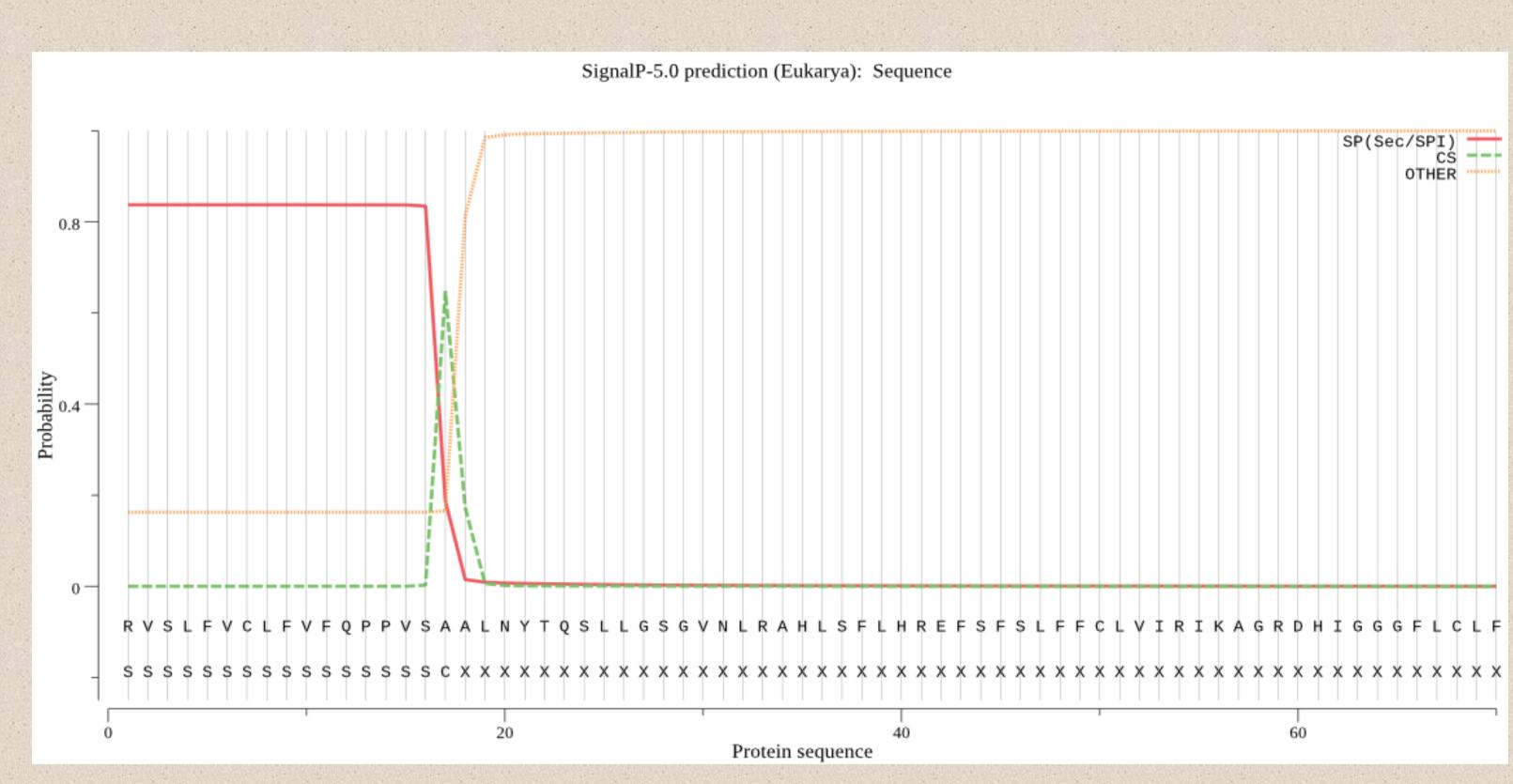


Figure 1. Likelihood of rs12378473 being the signal peptide transported and cleaved by Sec/SPI at each amino acid position.

## Results

Through subgrouping, SCORAD correlated AD more severe or SCORAD predisposition: immunological high subgroup (p=0.039), high eosinophil subgroup (p=0.006) subgroup (p=0.008). This study has identified novel child-onset factors such as caesarean section and non-smoking parent(s). GWAS has identified 6 candidate single-nucleotide polymorphism (SNPs) in association with the duration of AD and 5 potential SNPs in association with IgE.

## Conclusion

This pilot study supported our hypothesis that different AD subgroups based on AD severity, age of onset, IgE levels, and eosinophil count, may be associated with different environmental risk factors. Further fine mapping and functional studies shall be conducted to elucidate the exact roles of these candidate genes and environmental risk factors. Larger sample sizes are needed to validate the conclusion.

# Reference

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- 2. Hon, K.L., A.K. Leung, and B. Barankin, Barrier repair therapy in atopic dermatitis: an overview. Am J Clin Dermatol, 2013. 14(5): p. 389-99.