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As a result of increased healthcare requirements and the introduction of genetically modified foods, the problem of allergies is becoming a growing health problem [1]. The identification of potential allergenic proteins is usually done by scanning a database of allergenic proteins and locating known allergens with a high sequence similarity [2, 3]. However, this methodology will fail if a known allergen with similar sequence is not included in the database.

**Acknowledgements**: (Times New Roman, 10 pt, regular, single space)

We acknowledge the financial support …

**References**: (Times New Roman, 10 pt, regular, single space)

1. Sampson, H. J. Allergy Clin. Immunol., 115, 2005, 139-41.

2. Radauer, C. Int. Arch. Allergy Immunol. 173, 2017, 1-11.

3. Dimitrov, I., Naneva, L., Doytchinova, I., Bangov, I. Bioinformatics, 30, 2014, 846-851.

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