Construction of a High-density Genetic Linkage Map and QTL Analysis of Resistance to Cucumber Mosaic Virus in *Luffa cylindrica* (L.) Roem. Based on Specific Length Amplified Fragment Sequencing

Lina Lou (✉ linabeibei@163.com)

Institute of Vegetable Crops, Jiangsu Academy of Agricultural Sciences

Zhe Liu

Institute of vegetable crops, Jiangsu Academy of Agricultural Sciences

Xiaohong Liu

Institute of Vegetable Crops, Jiangsu Academy of Agricultural Sciences

Xiaojun Su

Institute of Vegetable Crops, Jiangsu Academy of Agricultural Sciences

Research article

Keywords: *Luffa cylindrica* L., Cucumber mosaic virus (CMV), high-density genetic map, SNPs, QTL analysis

Posted Date: November 2nd, 2019

DOI: <https://doi.org/10.21203/rs.2.15852/v3>

License: © ⓘ This work is licensed under a Creative Commons Attribution 4.0 International License.

[Read Full License](#)

The authors have withdrawn this preprint from Research Square