

**International Journal of Current  
Microbiology and Applied Sciences  
(IJCMAS) NAAS RATING-5.38, ICV-95.39**  
**ISSN: 2319-7692 (Print) ISSN 2319-7706 (Online)**

**An International, Monthly, Online, Free Access, Peer Reviewed,  
Indexed, fast track Scientific Research Journal**

**Certificate of Publication**

This is to certify that the following article reviewed by editorial board and published in International Journal of Current Microbiology and Applied Sciences (IJCMAS) ISSN: 2319-7692 (Print) ISSN 2319-7706 (Online).

Int.J.Curr.Microbiol.App.Sci.2019.8.(6):158-173

<https://doi.org/10.20546/ijcmas.2019.806.020>

Utilization and Characterization of Genome-wide SNP Markers for Assessment of Ecotypic Differentiation in *Arabidopsis thaliana*

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*International Journal of Current Microbiology and Applied Sciences*

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