



## RESEARCH PROFILE

### Contact Information

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### Doctoral Degree

Plant Science-Genetics, Molecular Genetics  
North Dakota State University

### Research Interest Area(s)

Plant Genetics, Molecular Genetics, and Epigenomics

### Active Grants

- National Science Foundation – Delaware EPSCoR
- USDA – AFRI - NEWBio
- USDA – AFRI - NIFA - Capacity Building Grant
- NSF-Research Experiences for Undergraduates

### Professional Affiliations

*American Association for the Advancement  
of Science*  
*American Society for Plant Biology (ASPB)*  
*Crop Science Society of America*  
*International Bean Genomics Consortium*  
*Bean Improvement Cooperative*

### Honors and Awards

2014 Educator of the year (Higher Education), Delaware  
Biosciences Association

2014 Invited Speaker, International Plant Genomics  
Congress, London, UK

2013 Chair, W2150: Breeding Common Bean (*Phaseolus  
vulgaris* L.) for Resistance to Abiotic and Biotic Stresses,  
Sustainable Production, and Enhanced Nutritional Value

2012 Invited Speaker, Phaseomics: The Genome, Special  
Edition, an international conference on common bean  
genomics, Mexico

2012 Member, International Committee on Phaseomics-  
planning next steps of bean genomics

2011 Secretary, W2150: Breeding Common Bean (*Phaseo-  
lus vulgaris* L.) for Resistance to Abiotic and Biotic Stresses,  
Sustainable Production, and Enhanced Nutritional Value

2011 Faculty Excellence Award for Research/Creative  
Activities, Delaware State University



**VENU KALAVACHARLA**

Professor

Melmaiee, K., Todd, A., Lee, R., McClean, P., Schleuter, J., Jackson, S., and Kalavacharla, V. Identification of molecular markers associated with the deleted region in common bean Ur-3 mutants. *Australian Journal of Crop Science*, 7:354-360. 2013.

Liu, Z., \*Crampton, M., Todd, A and Kalavacharla, V. Identification of expressed resistance gene-like sequences by data mining in 454-derived transcriptomic sequences of common bean (*Phaseolus vulgaris* L.). *BMC Plant Biology*, 12:42. 2012.

Liu, L, Yu, L, Kalavacharla, V. and Liu, Z. A Bayesian model for gene family evolution. *BMC Bioinformatics*, 12:426. 2011.

Kunjeti, S., Evans., T, Marsh, A., Gregory, N., Kunjeti, S., Meyers, B., Kalavacharla, V. and Donofrio, N. RNA-seq reveals infection related global changes in *Phytophthora phaseoli*, the causal agent of lima bean downy mildew. *Molecular Plant Pathology*, 13(5): 454-466 Year?

Kalavacharla, V., Liu, Z., Meyers, B.C., Thimmapuram, J. and Melmaiee, K. Identification and analysis of common bean (*Phaseolus vulgaris* L.) transcriptomes by massively parallel pyrosequencing. *BMC Plant Biology*, 11:135. 2011.