

**SUSAN R. STRICKLER**



Comparative and phylogenetic studies of wild and cultivated tomato species, including diversity studies to select accessions for pathogen response screening  
Identification of candidate genes involved in coffee humidity stress response

**Career Fair at Girl's Day Out "Jobs in Plant Biology"** Spring 2016 - 2020, 2022  
Cortland YWCA

**BTI Bioinformatics Course Instructor** Spring, 2012 - present  
Boyce Thompson Institute for Plant Research

**Summer Intern Bioinformatics Course Instructor** Summer, 2010 - present  
Boyce Thompson Institute for Plant Research

**Tech Savvy "Is Cauliflower the New Kale? What Bioinformatics Tells Us."** Mar 2017  
Cortland YWCA

**UPLB Bioinformatics Workshop Instructor** Oct 2014  
University of Philippines Los Banos

**AGA2013: Speciation Continuum and Non-model Population Genomics Workshop  
Teaching Assistant**

M. Scanlon (PI), M. Wu, A. Stroock (co-PI), H. Weatherspoon (co-PI), Y. Sun (co-PI), S. Strickler (program coordinator). Award number 1922551

**TRIAD Foundation** Jan 2021

**U.S. Patent No. 10,557,145: “FLGII-28 sensitivity 3 (FLS3) protein and methods of use”.**  
Inventors: G. B. Martin, S. Hind, S. Strickler. Filed 2014 by the Boyce Thompson Institute, Ithaca, NY. Converted to PCT Application No.PCT/US15/39520, filed July 8, 2015.

**U.S. Provisional Application (62/213,409): “Isolated nucleic acids and quantitative trait loci (QTL) from and methods of use thereof for increasing resistance to bacterial speck disease in tomato and other plants”.** Inventors: G. B. Martin, S. Hind, S. Strickler, B. Zhilong. Filed 2015 by the Boyce Thompson Institute, Ithaca, NY.

## CONFERENCES

### **Botany July 2022**

Workshop: *De novo* genome assembly and annotation with an emphasis on phylogenetic and population genetic studies

### **Botany July 2021**

Workshop: *De novo* genome assembly and annotation with an emphasis on phylogenetic and population genetic studies

Oral presentation: Improving Coffea genome assemblies with long-read data.

Oral presentation: Sol Genomics Network Workshop

**Plant and Animal Genome Conference** Jan 2015

Oral presentation: Genome assembly strategies of *Coffea arabica*.

*Lycopersicoides* reference genome facilitates insights into tomato specialized metabolism and immunity. *Plant J.* (2022), doi:10.1111/tpj.15770.

4. G. Moghe, **S. Strickler**, metaPathwayMap: A tool to predict metabolic pathway neighborhoods from structural classes of untargeted metabolomics peaks. *bioRxiv* (2022), p. 2022.03.15.484337.
5. J. H. Boyle, **S. Strickler**, A. Twyford, A. Ricono, **A. Powell**, **J. Zhang**, H. Xu, H. J. Dagleish, G. Jander, A. A. Agrawal, J. R. Puzey, Temporal matches and mismatches between monarch butterfly and milkweed population changes over the past 12,000 years. *bioRxiv* (2022), p. 2022.02.25.481796.
6. **G. Torres-Silva**, L. N. F. Correia, D. S. Batista, A. D. Koehler, S. V. Resende, E. Romanel, D. Cassol, A. M. R. Almeida, **S. R. Strickler**, C. D. Specht, W. C. Otoni, Transcriptome Analysis of *Melocactus glaucescens* (Cactaceae) Reveals Metabolic Changes During in vitro Shoot Organogenesis Induction. *Front. Plant Sci.* **12**, 697556 (2021).
7. **G. Torres-Silva**, L. N. F. Correia, A. D. Koehler, D. S. Batista, D. V. Faria, S. V. Resende, **S. R. Strickler**, J. Fouracre, E. Romanel, C. D. Specht, W. C. Otoni, Expression of *Melocactus glaucescens* SERK1 sheds new light on the mechanism of areolar activation in cacti. *Plant Cell Tissue Organ Cult.* (2021), doi:10.1007/s11240-021-02137-9.
8. L. Feiz, Y. Asakura, L. Mao, **S. R. Strickler**, Z. Fei, M. Rojas, A. Barkan, D. B. Stern, CFM1, a member of the CRM-domain protein family, functions in chloroplast group II intron splicing in *Setaria viridis*. *Plant J.*

15. **S. S.-E.-A. Zaidi, R. Z. Naqvi, M. Asif, S. Strickler, S. Shakir, M. Shafiq, A. M. Khan, I. Amin, B. Mishra, M. S. Mukhtar, B. E. Scheffler, J. A. Scheffler, L. A. Mueller, S. Mansoor,** Molecular insight into cotton leaf curl geminivirus disease resistance in cultivated cotton (*Gossypium hirsutum*). *Plant Biotechnol. J.* **18**, 691–706 (2020).
16. C. Mazo-Molina, S. Mainiero, B. J. Haefner, R. Bednarek, J. Zhang, A. Feder, K. Shi, **S. R. Strickler**, G. B. Martin, Ptr1 evolved convergently with RPS2 and Mr5 to mediate recognition of AvrRpt2 in diverse solanaceous species. *Plant J.* **103**, 1433–1445 (2020).
17. H. W. Choi, L. Wang, **A. F. Powell, S. R. Strickler**, D. Wang, D. A. Dempsey, F. C.

26. E. J. S. Michel, A. M. Hotto, **S. R. Strickler**, D. B. Stern, B. Castandet, A Guide to the Chloroplast Transcriptome Analysis Using RNA-Seq. *Methods Mol. Biol.* **1829**, 295–313 (2018).
27. **R. Z. Naqvi, S. S.-E.-A. Zaidi, K. P. Akhtar, S. Strickler**



**BTI News 2/4/2016**