

CURRICULUM VITAE

1. MAILING ADDRESS

Dr. Lam-Son Phan Tran, Professor

Texas Tech University

Department of Plant and Soil Science

Institute for Genomics of Crop Abiotic Stress Tolerance (IGCAST)

Experimental Science Building II (Room 201A)

Texas Tech University, Lubbock, TX 79409, USA

Email: son.tran@ttu.edu

Web of Science ResearcherID (A-1346-2009):

<https://www.webofscience.com/wos/author/record/397751>

ORCID: <https://orcid.org/0000-0001-9883-9768>

Google Scholar ID: https://scholar.google.co.jp/citations?hl=en&user=PA-FRFsAAAAJ&view_op=list_works&sortby=pubdate

2. EDUCATION

- 1994-1997: Godollo University of Agricultural Sciences, Hungary (currently, Szent Istvan campus of Hungarian University of Agriculture and Life Sciences, Hungary)
- Qualification: **Doctor of Philosophy**
Ph.D. program: *Biological Bases of Agricultural Biotechnology*.
- 1988-1994: Godollo University of Agricultural Sciences, Hungary
Qualification: **Master of Agricultural Sciences** (specialized in *Biotechnology*)
- 1991-1992: Courses in International Trade organized in Godollo University of Agricultural Sciences, Hungary
Qualification: **Certificate in International Trade**
- 1987-1988: International Preparatory Institute, Hungary: 1987-1988
Qualification: Hungarian Language Training Course
- 1986-1987: Hanoi Languages University, Vietnam
Qualification: Hungarian Language Training Course

3. PROFESSIONAL CAREER

- 09/2020-current: Professor, Texas Tech University, Department of Plant and Soil Science, Institute for Genomics of Crop Abiotic Stress Tolerance (IGCAST), USA
- 04/2021-current: Senior Visiting Scientist, RIKEN, Japan
- 05/2019-05/2022: Visiting Scientist, Duy Tan University, Vietnam
- 10/2015-12/2020: Visiting Scientist, Ton Duc Thang University, Vietnam
- 03/2012-12/2020: Visiting Professor of Vietnam Academy of Agricultural Science
- 01/01/2009-31/08/2020: Unit Leader, Stress Adaptation Research Unit (former Signaling Pathway Research Unit), RIKEN Center for Sustainable Resource Science (former RIKEN Plant Science Center), Yokohama, Japan.
- 06/08/2007-31/12/2008: Senior Research Scientist, Soybean Genomics and Biotechnology Laboratory, Division of Plant Science, National Center for Soybean Biotechnology, University of Missouri-Columbia, MO, 65211, USA.
- 01/10/2003-31/07/2007: Special Researcher, Biological Resources Division, Japan International Research Center for Agricultural Sciences (JIRCAS), Japan.
- 10/10/2001-30/09/2003: JIRCAS fellow, Biological Resources Division, Japan International Research Center for Agricultural Sciences (JIRCAS), Japan.

- 01/03/2001-05/10/2001: JSPS postdoctoral fellow, Department of Microbial Cell Biology, Graduate School of Biological Science, Nara Institute of Science and Technology, Japan.
- 01/12/1998-30/11/2000: STA postdoctoral fellow, Applied Bacteriology Laboratory, Division of Microbiology, National Food Research Institute, Tsukuba, Japan.
- 01/07/1998-30/11/1998: Senior Researcher, Phylaxia company in conjunction with Department of Biotechnology and Molecular Genetics of Szent Istvan University, Hungary.
- 01/09/1997-30/06/1998: Researcher, Molecular Genetics Group, Hungarian Academy of Sciences, Department of Biotechnology and Molecular Genetics, Szent Istvan University, Hungary.
- 1994-1997: Ph.D. student, Teaching Assistant, Department of Biotechnology and Molecular Genetics, Szent Istvan University, Hungary.
- 1992-1994: Teaching Assistant and Demonstrator, Department of Biotechnology and Molecular Genetics, Szent Istvan University, Hungary.
- 1990-1992: part-time technician, Department of Biotechnology and Molecular Genetics, Szent Istvan University, Hungary.

4. TEACHING AND SUPERVISING EXPERIENCE

Teaching

- 09/2020-current: Professor, Texas Tech University, Department of Plant and Soil Science, Institute for Genomics of Crop Abiotic Stress Tolerance (IGCAST), USA
- 03/2012-12/2020: Visiting Professor of Vietnamese Academy of Agricultural Sciences
- 1992-1994: Teaching Assistant and Demonstrator in Department of Biotechnology and Molecular Genetics, Szent Istvan University, Hungary.

Supervising

- 01/2009-current: supervised/co-supervised 8 graduate students, supervised 7 postdoctoral researchers, and hosted more than 20 M.Sc and Ph.D students and visiting scholars from different countries for short-term visits.
- 2007-2008: participated in advising Ph.D students of Molecular Genetics and Soybean Genomics Laboratory, University of Missouri-Columbia, USA.
- 2006-2007: participated in advising M.Sc and Ph.D students of Biological Resources Division, JIRCAS-Laboratory of Plant Molecular Physiology, University of Tokyo, Japan.
- 1994-1997: participated in advising M.Sc and Ph.D students of Department of Biotechnology and Molecular Genetics, Szent Istvan University, Hungary.

5. CURRENT RESEARCH OF INTERESTS

- Functions of signaling molecules (e.g., acetate, ethanol, Ca, hormones, NO, H₂S, Ca, etc.), and their crosstalk in plant responses to abiotic stresses (e.g., drought, salinity heavy metal stresses, and pathogens).
- Signaling regulatory network controlling plant growth, productivity and responses to environmental stresses.
- Functional genomics of legume crops for improvement of productivity under adverse conditions.
- Discovery of novel genes for improvement of abiotic stress tolerance of crops (chickpea, common bean, cotton, maize, rice, soybean, and sorghum).
- Isolation and characterization of plant growth-promoting microorganisms that can confer abiotic tolerance and biotic resistance to various crops.
- Large-scale data mining of plant genomic sequences for functional and comparative genomics.

6. PROFESSIONAL DUTIES

Handling editor

- Editor of Plant Nano Biology (from 03/2022~)
- Associate Editor of AoB PLANTS (from 11/2020~)
- Associate Editor of Plant-Environment Interactions (from 7/2019~)
- Editor of Scientific Reports (from 10/2014~)
- Associate Editor of Plant Molecular Biology Reporter (from 04/2014~)
- Associate Editor of BMC Plant Biology (from 10/2013~)
- Associate Editor of BMC Genomics (from 09/2013~)
- Editor of International Journal of Molecular Sciences (from 05/2013~)
- Associate Editor of Frontiers in Plant Science (from 07/2016~)
- Academic Editor of BioMed Research International (former Journal of Biomedicine and Biotechnology) (from 10/2012~)
- Academic Editor of PloS One (from 02/2012~)
- Associate Editor of Critical Reviews in Biotechnology (04/2020-09/2021)

Board members

- Member of Editorial Board of Oil Crop Science (from 01/2023~)
- Member of Editorial Board of Plant Communications (from 01/2020~)
- Member of Editorial Board of Current Protein & Peptide Science (from 02/2019~)
- Member of the Bio2 (Functional Biology) Fellowship panel at the Research Foundation Flanders (01/2019-12/2019)
- Member of Editorial Board of Plant & Cell Physiology (from 01/2016~)
- Member of Editorial Board of Plant Science (04/2014-12/2016)
- Member of Editorial Board of Critical Reviews in Biotechnology (from 04/2012-03/2020)
- Member of Editorial Board of Biotechnology Advances (from 03/2012~)

Ad hoc reviewer

- **For journals:** Agricultural Water Management; Agronomy; Annals of Botany; Australian Journal of Botany; BMC Genomics; BMC Plant Biology; Bioinformatics; Critical Reviews in Biotechnology; DNA Research; Environmental and Experimental Botany; Environmental Pollution; Environmental Science: Nano; Environmental Science & Technology; FEBS Letters; Frontiers in Plant Science; Functional & Integrative Genomics; Genetics and Molecular Biology; GM Crops & Food: Biotechnology in Agriculture and the Food Chain; Genome Biology; Journal of Agronomy and Crop Science; Journal of Chromatography A; Journal of Experimental Botany; Journal of Hazardous Materials; Journal of Plant Growth Regulation; Journal of Plant Physiology; Journal of Plant Research; Land Degradation & Development; Molecular Biology Reports; Molecular Genetics and Genomics; Molecular Plants, Molecular Plant-Microbe Interactions; Mycorrhiza; Nature Communications; New Phytologist; Pedosphere; Photosynthetica; Physiological and Molecular Plant Pathology; Physiologia Plantarum; Phytochemistry; Planta; Plant Biotechnology Journal; Plant Cell; Plant, Cell & Environment; Plant & Cell Physiology; Plant Cell Reports; Plant Cell, Tissue and Organ Culture; Plant Communications; Plant Genome; Plant Journal; Plant Molecular Biology; Plant Molecular Biology Reporter; Plant Physiology; Plant Physiology and Biochemistry; Plant Science; Plant Signaling & Behavior; PloS Genetics; PloS One; Proceedings of the National Academy of Sciences; Renewable &

Sustainable Energy; Rice; Science of the Total Environment; Scientific Reports; Symbiosis; Theoretical and Applied Genetics; Tree Genetics & Genomes; Trends in Plant Science.

- **For funding agencies:** National Science Foundation-USA, United State-Israel Binational Science Foundation, the United States-Israel Binational Agricultural Research & Development Fund (BARD), French National Research Agency, ERA-CAPS-German Research Foundation, Academy of Sciences of the Czech Republic, National Science Centre of Poland, International Foundation for Science, Biotechnology Panel of Israeli Ministry of Agriculture and Rural Development, AgreeSkills programme (INRA-France), Research Foundation Flanders (Belgium), National Center of Science and Technology Evaluation (Kazakhstan), Millenium Science Initiative (Chile).

7. RESEARCH GRANTS & FUNDING

A total of ~\$US500,000 has been secured as PI or co-PI since joining Texas Tech University from 09/01/2020.

NAME (List/PD #1 first)	SUPPORTING AGENCY AND AGENCY ACTIVE AWARD/PENDING PROPOSAL NUMBER	TOTAL \$ AMOUNT	EFFECTIVE AND EXPIRATION DATES	% OF TIME COMMITTED	TITLE OF PROJECT
PD: Lam-Son Phan Tran Co-PDs: Luis Herrera-Estrella, Damar Lopez-Arredondo, Chien Ha	USDA-AFRI (Pending)	\$300,000	03/01/2024 to 02/28/2027	5%	Generating high carbon assimilation and drought-tolerant cotton by exploring transgenerational memory effect of acetic acid application
PD: Lam-Son Phan Tran Co-PDs: Madhusudana Janga, Mostofa Mohammed	USDA-AFRI (Pending)	\$650,000	03/01/2024 to 02/29/2028	8%	Silicon-mediated enhancement of heat tolerance in cotton
PD: Damar Lopez-Arredondo Co-PDs: Luis Herrera-Estrella, Lam-Son Phan Tran	USDA-AFRI (Pending)	\$650,000	01/01/2024 to 12/31/2026	3%	Harnessing microalgae diversity to discover novel bioherbicides and nematocides with new modes of action for sustainable agriculture
PD: Gunvant Patil co-PD: Lam-Son Phan Tran	USDA-AFRI (Pending)	\$492,218	03/01/2024 to 02/28/2028	3%	Frost-proof soybeans: Enhancing cold tolerance through optimized nutrient uptake and beneficial microorganisms at germination and post-germination stages
PD: Henry Nguyen, Co-PD: Lam-Son Phan Tran	USDA-AFRI (Pending)	\$500,000	04/01/2024 to 03/31/2027	3%	Understanding and improvement of soybean tolerance to individual and combined cold and waterlogging stresses at early growth stage
PD: Henry Nguyen, Co-PDs: Lam-Son Phan Tran , Yinping Jiao	USDA-AFRI (Pending)	\$850,000	03/01/2024 to 02/28/2027	3%	Decoding the potential of sorghum root systems and microbiome: Strategies for resilience and nitrogen-use-efficiency in hot and arid land
PD: Lam-Son Phan Tran	Multi-Regional Soybean Checkoff_Mid-South Regional project (Pending)	\$70,000	01/01/2024 to 09/30/2024	3%	Toward microbial syncom-dryland farming: Improving agricultural practices through microbiome-induced soybean resilience under hot and arid field conditions
PD: Henry Nguyen, Co-PDs: Lam-Son Phan Tran , Gunvant Patil, Vasit Sagan, C. Nathan Hancock, Kendall R. Kirk	Multi-Regional Soybean Checkoff-National Project (Pending)	\$450,000	01/01/2024 to 09/30/2024	3%	Genetic and physiological mechanisms underlying soybean resilience to drought and hot climate
PI: Lam-Son Phan Tran Co-PI: Chien Van Ha	The Texas State Support Committee (22-484TX) and Cotton Incorporated (22-618) (Current)	\$25,650	01/01/2024 – 12/31/2024	3%	Improvement of cotton growth and yield performance under drought stress with the employment of zinc oxide nanoparticles (Continuation for 3 rd year)

PI: Lam-Son Phan Tran Co-PI: Aarti Gupta	The Texas State Support Committee (24-921TX) (Current)	\$20,000	01/01/2024 – 12/31/2024	3%	Roles of plant growth-promoting bacteria in improving cotton growth and yield performance under high salinity (1st year)
PI: Chien Van Ha Co-PI: Lam-Son Phan Tran , Noureddine Abidi	The Texas State Support Committee (24-909TX) (Current)	\$25,000	01/01/2024 – 12/31/2024	3%	Enhancement of drought tolerance and reduction of fiber detachment force of cotton using co-application of acetic acid and ethanol (1st year)
PI: Damar Lopez-Arredondo Co-PIs: Luis Herrera-Estrella, Lam-Son Phan Tran	United Soybean Board (Project #2314-209-0501) (Current)	\$131,956	10/01/2023-09/30/2024	2% (20% allocation of credit)	Genetic improvement of soybean to boost weed control and phosphorus nutrition (Continuation for 2nd year)
PI: Mayank Gururani Co-PI: Lam-Son Phan Tran	United Arab Emirates University (UAEU) Program for Advanced Research, ID: 700027433) (Current)	\$95,000	04/01/2023-03/31/2025	5%	Study on the mechanism of brassinosteroids regulating symbiotic nodulation in soybean
PI: Luis Herrera-Estrella Co-PIs: Lam-Son Phan Tran , Damar Lopez-Arredondo, Gunvant Patil, Yinping Jiao	United Sorghum Checkoff Program (A23-0146) (Current)	\$1,000,000	01/01/2023-12/31/2027	3% (15% allocation of credit)	An integrated research program to accelerate sorghum breeding and management for improving weed control, abiotic stress tolerance and grain quality
PI: Lam-Son Phan Tran Co-PI: Mohamad Golam Mostofa	The Texas State Support Committee (22-484TX) and Cotton Incorporated (22-618) (Current)	\$25,650	01/01/2023 – 12/31/2023	3%	Improvement of cotton growth and yield performance under drought stress with the employment zinc oxide nanoparticles (Continuation for 2nd year)
PI: Damar Lopez-Arredondo Co-PIs: Luis Herrera-Estrella, Lam-Son Phan Tran	United Soybean Board (Project #2314-209-0501) (Finished)	\$108,979	10/01/2022-09/30/2023	2% (20% allocation of credit)	Genetic improvement of soybean to boost weed control and phosphorus nutrition
PI: Lam-Son Phan Tran Co-PI: Mohamad Golam Mostofa	United Soybean Board (Project #2313-209-0101) (Finished)	\$112,474	10/01/2022-09/30/2023	5%	Combining chemical and genetic strategies for improvement of growth and yield potential in soybean under drought stress (continuation for 2nd year , project #2220-172-0148)
PI: Lam-Son Phan Tran Co-PI: Mohamad Golam Mostofa	United Soybean Board (Project #2220-172-0148) (Finished)	\$99,656	10/2021 – 03/2023	5%	Combining chemical and genetic strategies for improvement of growth and yield potential in soybean under drought stress (1st year)
PI: Lam-Son Phan Tran Co-PIs: Mendu Venugopal, Dampanaboina Lavanya	BASF-TTU Project Revolution (Finished)	\$138,818	11/2021 – 03/2023	5%	Genome editing to improve photosynthetic efficiency and biomass yield in soybean
PI: Lam-Son Phan Tran Co-PI: Mohamad Golam Mostofa	The Texas State Support Committee (22-484TX) and Cotton Incorporated (22-618) (Finished)	\$23,650	1/2022 – 12/2022	3%	Improvement of cotton growth and yield performance under drought stress with the employment of zinc oxide nanoparticles (1st year)

8. AWARDS AND HONOURS

- Highly Cited Researcher 2023 (Thomson Reuters/Clarivate Analytics)
- The Texas Tech Parents Association's Barney E. Rushing, Jr., Faculty Distinguished Research Award – STEM, 2023
- Elected Fellow of *The National Academy of Sciences, India* (effective 01-January-2023)
- Highly Cited Researcher 2022 (Thomson Reuters/Clarivate Analytics)
- Letter of Appreciation from the President of RIKEN for receiving Highly Cited Researcher 2022
- Highly Cited Researcher 2021 (Thomson Reuters/Clarivate Analytics)

- Letter of Appreciation from the President of RIKEN for receiving Highly Cited Researcher 2021
- Highly Cited Researcher 2020 (Thomson Reuters/Clarivate Analytics)
- Letter of Appreciation from the President of RIKEN for receiving Highly Cited Researcher 2020
- Visiting Scientists Award from the Chinese Academy of Sciences for 2021
- Stanford University's lists of 2% scientists (both "career-long citation impact" and "citation impact during the single calendar year 2019") (PLoS Biology 18:e300098; <https://doi.org/10.1371/journal.pbio.3000918>)
- Elected Fellow of *The World Academy of Sciences (TWAS) for the advancement of science in developing countries* (effective 01-January-2020)
- Letter of Appreciation from the President of RIKEN for receiving Highly Cited Researcher 2019
- Highly Cited Researcher 2019 (Thomson Reuters/Clarivate Analytics)
- Letter of Appreciation from the President of RIKEN for receiving Highly Cited Researcher 2018
- Highly Cited Researcher 2018 (Thomson Reuters/Clarivate Analytics)
- Letter of Appreciation from the President of RIKEN for receiving Highly Cited Researcher 2016
- Highly Cited Researcher 2016 (Thomson Reuters/Clarivate Analytics)
- 10/2001-09/2003: JIRCAS Visiting Research Fellowship.
- 03/2001-09/2001: Postdoctoral fellowship from Japan Society for the Promotion of Science.
- 11/1998-11/2000: Postdoctoral fellowship from Science and Technology Agency, Japan.
- 09/1994-08/1997: Ph.D. scholarship from the Szent Istvan University and Ministry of Training and Education, Hungary.
- 04/1993: Honour Prize from the Hungarian Ministry of Environmental Protection on National Scientific Congress of University Students, Biotechnology Section, Szombathely, Hungary.
- 04/1993: Second Prize on National Scientific Congress of University Students, Biotechnology Section, Szombathely, Hungary.
- 12/1992: First Prize on Scientific Conference of Szent Istvan University's Students, Biotechnology Section, Godollo, Hungary.
- 08/1987-08/1994: Scholarship for Outstanding Achievement on University Admission Examination, awarded by the Vietnamese Government for B.Sc-M.Sc studies in Hungary.

PUBLICATIONS

I. Patents

Ha CV, **Tran LP**, Herrera-Estrella L, Christian M (2022). U.S. Provisional Pat. App. No. 63/388,878 (filed on 07/13/ 2022, 25%/each)

II. Research papers (*corresponding author)

2023

1. Rahman MM, Das AK, Sultana S, Ghosh PK, Islam MR, Keya SS, Ahmed M, Nihad SAI, Khan MAR, Lovell MC, Rahman MA, Ahsan SM, Anik TR, Fnu P, ***Tran LP**, Mostofa MG (2023). Biochar potentially enhances maize tolerance to arsenic toxicity by improving physiological and biochemical responses to excessive arsenate. **Biochar** 5:71.
2. Khan MAR, Mahmud A, Ghosh UK, Hossain MS, Siddiqui MN, Islam AKMA, Anik TR, Rahman MM, Sharma A, Abdelrahman M, Ha CV, Mostofa MG, ***Tran LP** (2023). Exploring the phenotypic and genetic variabilities in yield and yield-related traits of the diallel-crossed F₅ population of Aus rice. **Plants** 12:3601.
3. Tian Y, Liu Y, Uwaremwe C, Zhao X, Yue L, Zhou Q, Wang Y, **Tran LP**, Li W, Chen G, Sha Y, Wang R (2023). Characterization of three new plant growth-promoting microbes and effects of the interkingdom interactions on plant growth and disease prevention. **Plant Cell Rep** 42:1757-76.
4. Liu H, Liu S, Wang F, Liu Y, Liu Y, Sun J, McConkey KR, **Tran LP**, Dong Y, Yu L, Wang Q (2023). Identifying ecological compensation areas for ecosystem services degradation on the Qinghai-Tibet Plateau. **J Clean Prod** 423:138626.
5. Wang F, Liu S, Liu H, Zhao Y, Dong Y, **Tran LP**, Li W (2023). Resilience assessment of the nitrogen flow system in food production and consumption for sustainable development on the Qinghai-Tibet Plateau. **Sci Total Environ** 896:165223
6. Yin X, Xiang Y, Huang F-Q, Chen Y, Ding H, Du J, Chen X, Wang X, Wei X, Cai Y-Y, Guo D, Alolga RN, Kan X, Zhang B, Alejo-Jacuinde G, Li P, ***Tran LP**, Herrera-Estrella L, Lu X, Qi L-W (2020). Comparative genomics of the medicinal plants *Lonicera acranthoides* and *L. japonica* provides insight into genus genome evolution and hederagenin-based saponin biosynthesis. **Plant Biotech J** 21:2209-23.
7. Somaddar U, Mia S, Khalil MI, Sarker UK, Uddin MR, Kaysar MS, Chaki AK, Robin AHK, Hashem A, Abd_Allah EF, Ha CV, Gupta A, Park J-I, ***Tran LP**, Gopal Saha. Effect of reproductive stage-waterlogging on the growth and yield of upland cotton (*Gossypium hirsutum*). **Plants** 12:1548.
8. Anik TR, Mostofa MG, Rahman MM, Khan MAR, Ghosh PK, Sultana S, Das AK, Hossain MS, Keya SS, Rahman MA, Jahan N, ***Tran LP** (2023). Zn supplementation mitigates drought effects on cotton by improving photosynthetic performance and antioxidant defense mechanisms. **Antioxidants** 12:854.
9. Nguyen HM, Ha CV, Le VP, Bui HT, Wirschell M, Keya SS, Li W, Li M, Pham NT, Do AM, Le MQ, Anik TR, ***Tran LP** (2023). Improvement of photosynthetic performance by acetic acid to enhance drought tolerance in common bean (*Phaseolus vulgaris*). **J Plant Growth Regul** 42:7116-28.
10. Keya SS, Mostofa MG, Rahman MM, Das AK, Sultana S, Ghosh PK, Anik TR, Ahsan SM, Rahman MA, Jahan N, ***Tran LP** (2023). Salicylic acid application improves

- photosynthetic performance and biochemical responses to mitigate saline stress in cotton. **J Plant Growth Regul** 42:5881-94.
11. Zahedi SM, Hosseini MS, Karimi M, Gholami R, Amini M, Abdelrahman M, ***Tran LP** (2023). Chitosan-based Schiff base-metal (Fe, Cu, and Zn) complexes mitigate the negative consequences of drought stress on pomegranate fruits. **Plant Physiol Biochem** 196:952-64.
 12. Younes NA, Anik TR, Rahman MM, Wardany AA, Dawood MFA, ***Tran LP**, Abdel Latef A, Mostofa MG (2023). Effects of microbial biostimulants (*Trichoderma album* and *Bacillus megaterium*) on growth, quality attributes, and yield of onion under field conditions. **Heliyon** 9:e14203.
 13. Yi F, Song A, Cheng K, Liu J, Wang C, Shao L, Wu S, Wang P, Zhu J, Liang Z, Chang Y, Chu Z, Cai C, Zhang X, Wang P, Chen A, Xu J, Burritt DJ, Herrera-Estrella L, ***Tran LP**, Li W, Cai Y (2023). Strigolactones positively regulate Verticillium wilt resistance in cotton via crosstalk with other hormones. **Plant Physiol** 192:945-66.
 14. Li L., Li L, Cui S, Qian D, Lyu S, Liu W, Botella JR, Li H, Burritt DJ, **Tran LP**, Li W, Zhang Y (2023). *PDC1* is activated by ABF4 and inhibits seed germination by promoting ROS accumulation in *Arabidopsis*. **Environ Exp Bot** 266:105188.
 15. Wang F, Liu S, Liu H, Liu Y, Lu Y, Wang Q, Dong Y, Sun J, **Tran LP**, Li W (2023). Aggravation of nitrogen losses driven by agriculture and livestock husbandry development on the Qinghai-Tibet Plateau. **J Environ Manage** 326:116795.

2022

16. Van Ha C, Mostofa MG, Nguyen KH, Tran CD, Watanabe Y, Li W, Osakabe Y, Sato M, Toyooka K, Tanaka M, Seki M, Burritt DJ, Anderson C, Zhang R, Nguyen HM, Le VP, Bui HT, Mochida K, ***Tran LP** (2022). The histidine phosphotransfer AHP4 plays a negative role in *Arabidopsis* plant response to drought. **Plant J** 111:1732-52.
17. Mostofa MG, Abdelrahman M, Rahman MM, Tran CD, Nguyen KH, Watanabe Y, Itouga M, Li W, Wang Z, Mochida K, ***Tran LP** (2022). Karrikin receptor KAI2 coordinates salt tolerance mechanisms in *Arabidopsis thaliana*. **Plant Cell Physiol** 63:1927-42.
18. Abdelrahman M, Mostofa MG, Tran CD, El-Sayed M, Li W, Sulieman S, Tanaka M, Seki M, ***Tran LP** (2022). The karrikin receptor KARRIKIN INSENSITIVE2 positively regulates heat stress tolerance in *Arabidopsis thaliana*. **Plant Cell Physiol** 63:1914-26.
19. Nasr Esfahani M, Kusano M, Abdelrahman M, Nguyen KH, Watanabe Y, Mochida K, Burritt DJ, ***Tran LP** (2022). Differential metabolic rearrangements in the roots and leaves of *Cicer arietinum* caused by single or double nitrate and/or phosphate deficiencies. **Plant J** 111:1643-59.
20. Tian H, Watanabe Y, Nguyen KH, Tran CD, Abdelrahman M, Liang X, Kun X, Sepulveda C, Mostofa MG, Ha CV, Nelson DC, Mochida K, Tian C, Tanaka M, Seki M, Miao Y, ***Tran LP**, Li W (2022). KARRIKIN UPREGULATED F-BOX 1 negatively regulates drought tolerance in *Arabidopsis*. **Plant Physiol** 190:2671-87. (commented at <https://doi.org/10.1093/plphys/kiac413>).
21. Hossain MS, Abdelrahman M, Tran CD, Nguyen KH, Chu HD, Watanabe Y, Fujita M, ***Tran LP** (2022). Modulation of osmoprotection and antioxidant defense by exogenously applied acetate enhances cadmium stress tolerance in lentil seedlings. **Environ Pollut** 308:119687.
22. Wang F, Liu S, Liu H, Liu Y, Lu Y, Wang Q, Dong Y, **Tran LP**, Sun J, Zhao W (2022). Scenarios and sustainability of the economy–nitrogen–resource–environment system using a system dynamic model on the Qinghai-Tibet Plateau. **J Environ Manage** 318:115623.

23. Yu L, Liu S, Wang F, Liu Y, Liu H, Wang Q, **Tran LP**, Dong Y, Li W (2022). Strategies for agricultural production management based on land, water and carbon footprints on the Qinghai-Tibet Plateau. **J Clean Prod** 362:132563.
24. Feng Z, Liang X, Tian H, Jia J, Watanabe Y, Nguyen KH, Tran CD, Abdelrahman M, Xu, K, Mostofa MG, Ha CV, Mochida K, Tian C, Tanaka M, Seki M, Liang Z, Miao Y, ***Tran LP**, Li W (2022). SUPPRESSOR of MAX2 1 (SMAX1) and SMAX1-LIKE2 (SMXL2) negatively regulate drought resistance in *Arabidopsis thaliana*. **Plant Cell Physiol** 63:1900-13.
25. Yang Y, Fang X, Chen M, Wang L, Xia J, Wang Z, Fang J, **Tran LP**, Shangguan L (2022) Copper stress in grapevine: Consequences, responses, and a novel mitigation strategy using 5-aminolevulinic acid. **Environ Pollut** 307:119561.
26. Wang Q, Liu S, Wang F, Liu H, Liu Y, Lu Y, Sun J, **Tran LP**, Dong Y (2022). Quantifying carbon sequestration service flow associated with human activities based on network model on the Qinghai-Tibet Plateau. **Front Environ Sci** 10:900908 |
27. Keya SS, Mostofa MG, Rahman MM, Das AK, Rahman MA, Anik TR, Islam MR, Sultana S, Khan MAR, Islam MR, Watanabe Y, Mochida K, ***Tran LP** (2022). Effects of glutathione on waterlogging-induced damage in sesame crop. **Ind Crops Prod** 185:115092.
28. Li W, Sun Y, Li K, Tian H, Jia J, Zhang H, Wang Y, Wang H, Bi B, Guo J, **Tran LP**, Miao Y (2022). Sinapate esters mediate UV-B-induced stomatal closure by regulating nitric oxide, hydrogen peroxide, and malate accumulation in *Arabidopsis thaliana*. **Plant Cell Physiol** 63:1890-9.
29. Rahman MM, Mostofa MG, Das AK, Anik TR, Keya SS, Ahsan SM, Khan MAR, Ahmed M, Rahman MA, Hossain MM, ***Tran LP** (2022). Ethanol positively modulates photosynthetic traits, antioxidant defense and osmoprotectant levels to enhance drought acclimatization in soybean. **Antioxidants** 11:516.
30. La HV, Chu HD, Tran CD, Nguyen KH, Le QTN, Hoang CM, Cao BP, Phan ATC, Nguyen BD, Nguyen TQ, Nguyen LV, Ha CV, Le HT, Le HH, Le TD, ***Tran LP** (2022). Insights into the gene and protein structures of the *CaSWEET* family members in chickpea (*Cicer arietinum*), and their gene expression patterns in different organs under various stress and abscisic acid treatments. **Gene** 819:146210.
31. Das AK, Anik TR, Rahman MM, Keya SS, Islam MR, Rahman MA, Sultana S, Ghosh PK, Khan S, Ahamed T, Ghosh TK, **Tran LP**, Mostofa MG (2022). Ethanol treatment enhances physiological and biochemical responses to mitigate saline toxicity in soybean. **Plants** 11:272.
32. Yu L, Liu S, Wang F, Liu Y, Li M, Wang Q, Dong S, Zhao W, **Tran LP**, Sun Y, Li W, Dong Y, Beazley R, Qian H (2022). Effects of agricultural activities on energy-carbon-water nexus of the Qinghai-Tibet Plateau. **J Clean Prod** 331:129995.
33. Tian L, Chang J, Shi S, Ji L, Zhang J, Sun Y, Li X, Li X, Xie H, Cai Y, Chen D, van Veen JA, Kuramae EE, ***Tran LP**, Tian C (2022). Comparison of methane metabolism in the rhizomicrobiome of wild and related cultivated rice accessions reveals a strong impact of crop domestication. **Sci Total Environ** 803:150131.
34. Yin H, Zhou H, Wang W, ***Tran LP**, Zhang B (2022). Evidence for miRNAs involved in the high-altitude responses of sainfoin (*Onobrychis viciifolia*) grown in the Qinghai-Tibetan plateau. **J Plant Biochem Biotechnol** 31:533-44.

35. Luo Y, Teng S, Yin H, Zhang S, Tuo X, ***Tran LP** (2021). Transcriptome analysis reveals roles of anthocyanin- and jasmonic acid-biosynthetic pathways in rapeseed in response to high light stress. **Int J Mol Sci** **22**:13027.
36. Abdelrahman M, Nishiyama R, Tran CD, Kusano M, Nakabayashi R, Okazaki Y, Matsuda F, Chávez Montes RA, Mostofa MG, Li W, Watanabe Y, Fukushima A, Tanaka M, Seki M, Saito K, Herrera-Estrella L, ***Tran LP** (2021). Defective cytokinin signaling reprograms lipid and flavonoid gene-to-metabolite networks to mitigate high salinity in *Arabidopsis*. **Proc Natl Acad Sci U S A** **118**:e2105021118.
37. Mostofa MG, Ha CV, Rahman MM, Nguyen KH, Keya SS, Watanabe Y, Itouga M, Hashem A, Abd_Allah EF, Fujita M, ***Tran LP** (2021). Strigolactones modulate cellular antioxidant defense mechanisms to mitigate arsenate toxicity in rice shoots. **Antioxidants** **10**:1815.
38. Wan J, Wang R, Zhang P, Sun L, Ju Q, Huang H, Lü S, ***Tran LP**, Xu J (2021). MYB70 modulates seed germination and root system development in *Arabidopsis*. **iScience** **24**:103228.
39. Zahedi SM, Hosseini MS, Hoveizeh FN, Gholami R, Abdelrahman M, ***Tran LP** (2021). Exogenous melatonin mitigates salinity-induced damage in olive seedlings by modulating ion homeostasis, antioxidant defense and phytohormone balance. **Physiol Plant** **173**:1682-94.
40. Hoang XLT, Chuong NN, Hoa TTK, Doan H, Van PHP, Trang LDM, Huyen PNT, Le DT, ***Tran LP**, Thao NP (2021). The drought-mediated soybean GmNAC085 functions as a positive regulator of plant response to salinity. **Int J Mol Sci** **22**:8986.
41. Sun Y, Tian L, Chang J, Shi S, Zhang J, Xie H, Cai Y, Chen D, Kuramae EE, van Veen JA, Li W, ***Tran LP**, Tian C (2021). Rice domestication influences the composition and function of the rhizosphere bacterial chemotaxis systems. **Plant Soil** **466**:81-99.
42. Kumari P, Singh S, Yadav S, ***Tran LP** (2021). Influence of different types of explants in chickpea regeneration using thidiazuron seed-priming. **J Plant Res** **134**:1149-54.
43. Salahvarzi M, Nasr Esfahani M, Shirzadi N, Burritt DJ, ***Tran LP** (2021). Genotype- and tissue-specific physiological and biochemical changes of two chickpea (*Cicer arietinum*) varieties following a rapid dehydration. **Physiol Plant** **172**:1822-34.
44. Liu S, Sun Y, Wu X, Li W, Liu Y, ***Tran LP** (2021). Driving factor analysis of ecosystem services balance for watershed management in the Lancang River Valley, Southwest China. **Land** **10**:522.
45. Zhang S, Guo Y, Zhang Y, Guo J, Li K, Fu W, Jia Z, Li W, **Tran LP**, Jia KP, Miao Y (2021). Genome-wide identification, characterization and expression profiles of the *CCD* gene family in *Gossypium* species. **3 Biotech** **11**:249.
46. Younes NA, Rahman MM, Wardany AA, Dawood MFA, Mostofa MG, Keya SS, Abdel Latef A, ***Tran LP** (2021). Antioxidants and bioactive compounds in licorice root extract potentially contribute to improving growth, bulb quality and yield of onion (*Allium cepa*). **Molecules** **26**:2633
47. Tian L, Wang E, Lin X, Chang J, Chen H, Wang J, Chen D, ***Tran LP**, Tian C (2021). Wild rice harbors more root endophytic fungi than cultivated rice in the F1 offspring after crossbreeding. **BMC Genomics** **22**:278
48. Echeverria A, Larrainzar E, Li W, Watanabe Y, Sato M, Tran CD, Moler JA, Hirai MY, Sawada Y, ***Tran LP**, Gonzalez EM (2021). *Medicago sativa* and *Medicago truncatula* show contrasting root metabolic responses to drought. **Front Plant Sci** **12**:652143.
49. Mostofa MG, Rahman MM, Nguyen KH, Li W, Watanabe Y, Tran CD, Zhang M, Itouga M, Fujita M, ***Tran LP** (2021). Strigolactones regulate arsenate uptake, vacuolar-sequestration and antioxidant defense responses to resist arsenic toxicity in rice roots. **J Hazard Mater** **325**:125589.

50. Hosseini MS, Samsampour D, Zahedi SM, Zamanian K, Rahman M, Mostofa MG, ***Tran LP** (2021). Melatonin alleviates drought impact on growth and essential oil yield of lemon verbena by enhancing antioxidant responses, mineral balance, and abscisic acid content. **Physiol Plant** **172**:1363-75.
51. Siddiqui N, Mostofa MG, Rahman M, Tahjib-Ul-Arif, Das AK, Mohi-Ud-Din M, Rohman MM, Hafiz HR, Ansary MU, ***Tran LP** (2021). Glutathione improves rice tolerance to submergence: insights into its physiological and biochemical mechanisms. **J Biotechnol** **325**:109-18
52. Nasr Esfahani M, Inoue K, Nguyen KH, Chu HD, Watanabe Y, Kanatani A, Burritt DJ, Mochida K, ***Tran LP** (2021). Phosphate or nitrate imbalance induces stronger molecular responses than combined nutrient deprivation in roots and leaves of chickpea plants. **Plant Cell Environ** **44**:574-97.
53. Chen L, Yang H, Fang Y, Guo W, Chen H, Zhang X, Dai W, Chen S, Hao Q, Yuan S, Zhang C, Huang Y, Shan Z, Yang Z, Qiu D, Liu X, ***Tran LP**, Zhou X, Cao D (2021). Overexpression of *GmMYB14* improves high-density yield and drought tolerance of soybean through regulating plant architecture mediated by the brassinosteroid pathway. **Plant Biotech J** **19**:702-16.
54. Rahman M, Mostofa MG, Keya SS, Rahman A, Das AK, Islam R, Abdelrahman M, Bhuiyan SU, Naznin T, Ansary MU, ***Tran LP** (2021). Acetic acid improves drought acclimation in soybean: an integrative response of photosynthesis, osmoregulation, mineral uptake and antioxidant defense. **Physiol Plant** **172**:334-50.
55. Tahjib-Ul-Arif M, Sohag AAM, Mostofa MG, Polash MAS, Mahamud AGMSU, Afrin S, Hossain MA, Hossain MA, Murata Y, ***Tran LP** (2021). Comparative effects of ascobin and glutathione on copper homeostasis and oxidative stress metabolism in mitigation of copper toxicity in rice. **Plant Biol** **23**:162-9.

2020

56. Tian L, Shi S, Sun Y, ***Tran LP**, Tian C (2020). The compositions of rhizosphere microbiomes of wild and cultivated soybeans changed following the hybridization of their F1 and F2 generations. **Eur J Soil Biol** **101**:103249.
57. Niu L, Chu HD, Tran CD, Nguyen KH, Pham HX, Le DT, Li W, Wang W, Le TD, ***Tran LP** (2020). The *GATA* gene family in chickpea: structure analysis and transcriptional responses to abscisic acid and dehydration treatments revealed potential genes involved in drought adaptation. **J Plant Growth Regul** **39**: 1647-60.
58. Yin X, Fan H, Chen Y, Li LZ, Song W, Fan Y, Zhou W, Ma G, Alolga RN, Li W, Zhang B, Li P, ***Tran LP**, Lu X, Qi LW (2020). Integrative omic and transgenic analyses reveal the positive effect of ultraviolet-B irradiation on salvianolic acid biosynthesis through upregulation of *SmNAC1*. **Plant J** **104**:781-99.
59. Zahedi SM, Abdelrahman M, Hosseini MS, Yousefi R, ***Tran LP** (2020). Physical and biochemical properties of 10 wild almond (*Amygdalus scoparia*) accessions naturally grown in Iran. **Food Biosci** **37**:100721.
60. Yin H, Wang L, Shi Y, Qian C, Zhou H, Wang W, Ma X-F, ***Tran LP**, Zhang B (2020). The east asian winter monsoon acts as a major selective factor in the intraspecific differentiation of drought-tolerant *Nitraria tangutorum* in northwest China. **Plants** **9**:1100.
61. Li W, Gupta A, Tian H, Nguyen KH, Tran CD, Watanabe Y, Tian C, Li K, Yang Y, Guo J, Luo J, Miao Y, ***Tran LP** (2020). Different strategies of strigolactone and karrikin signals in regulating the resistance of *Arabidopsis thaliana* to water-deficit stress. **Plant Signal Behav** **15**:e1789321.

62. Yin H, Zhou H, Wang W, ***Tran LP**, Zhang B (2020). Transcriptome analysis reveals potential roles of abscisic acid and polyphenols in adaptation of *Onobrychis viciifolia* to extreme environmental conditions in the Qinghai-Tibetan plateau. **Biomolecules** **10**:E967.
63. Zhang P, Wang R, Yang X, Ju Q, Li W, Lü S, ***Tran LP**, Xu J (2020). The R2R3-MYB transcription factor AtMYB49 modulates salt tolerance in *Arabidopsis* by modulating the cuticle formation and antioxidant defence. **Plant Cell Environ** **43**:1925-43.
64. Li P, Wen J, Chen P, Guo P, Ke Y, Wang M, Liu M, ***Tran LP**, Li J, Du H (2020). MYB superfamily in *Brassica napus*: evidence for hormone-mediated expression profiles, large expansion, and functions in root hair development. **Biomolecules** **10**:E875.
65. Mostofa MG, Rahman MM, Siddiqui MN, Fujita M, ***Tran LP** (2020). Salicylic acid antagonizes selenium phytotoxicity in rice: selenium homeostasis, oxidative stress metabolism and methylglyoxal detoxification. **J Hazard Mater** **394**:122572.
66. Li W, Nguyen KH, Tran CD, Watanabe Y, Tian C, Yin X, Li K, Yang Y, Guo J, Miao Y, Yamaguchi S, ***Tran LP** (2020). Negative roles of strigolactone-related SMXL6, 7 and 8 proteins in drought resistance in *Arabidopsis*. **Biomolecules** **10**:607.
67. Abdelrahman M, Nakabayashi R, Mori T, Ikeuchi T, Mori M, Murakami K, Ozaki Y, Matsumoto M, Uragami A, Tsujimoto H, ***Tran LP**, Kanno A (2020). Comparative metabolome and transcriptome analyses of susceptible *Asparagus officinalis* and resistant wild *A. kiusianus* revealed insights into stem blight disease resistance. **Plant Cell Physiol** **61**:1464-76.
68. Pour-Aboughadareh A, Etminan A, Abdelrahman M, Siddique KHM, ***Tran LS** (2020). Assessment of biochemical and physiological parameters of durum wheat genotypes at the seedling stage during polyethylene glycol-induced water stress. **Plant Growth Regul** **92**: 81-93.
69. Nghia DHT, Chuong NN, Hoang XLT, Nguyen NC, Tu NHC, Huy NVG, Ha BTT, Nam TNH, Thu NBA, **Tran LP**, Thao NP (2020). Heterologous expression of a soybean gene *RR34* conferred improved drought resistance of transgenic *Arabidopsis*. **Plants** **9**:494.
70. Li W, Nguyen KH, Chu HD, Watanabe W, Osakabe Y, Sato M, Toyooka K, Seo M, Tian L, Tian C, Yamaguchi S, Tanaka M, Seki M, ***Tran LP** (2020). Comparative functional analyses of DWARF14 and KARRIKIN INSENSITIVE2 in drought adaptation of *Arabidopsis thaliana*. **Plant J** **103**:111-27.
71. Guo W, Chen L, Chen H, Yang H, You Q, Bao A, Chen S, Hao Q, Huang Y, Qiu D, Shan Z, Yang Z, Yuan S, Zhang C, Zhang X, Jiao Y, ***Tran LP**, Zhou X, Cao D (2020). Overexpression of *GmWRI1b* in soybean stably improves plant architecture and associated yield parameters, and increases total seed oil production under field conditions. **Plant Biotech J** **18**:1639-41.
72. Tian L, Shi S, Ma L, ***Tran LS**, Tian C (2020). Community structures of the rhizomicrobiomes of cultivated and wild soybeans in their continuous cropping. **Microbiol Res** **232**:126390.
73. Hossain MS, Abdelrahman M, Tran CD, Nguyen KH, Chu HD, Watanabe Y, Hasanuzzaman M, Mohsin SM, Fujita M, ***Tran LP** (2020). Insights into acetate-mediated copper homeostasis and antioxidant defense in lentil under excessive copper stress. **Environ Pollut** **258**:113544.
74. Hoang XLT, Nguyen NC, Nguyen YNH, Watanabe Y, ***Tran LP**, Thao NP (2020). The soybean GmNAC019 transcription factor mediates drought tolerance in *Arabidopsis* in an abscisic acid-dependent manner. **Int J Mol Sci** **21**:286.
75. Sulieman S, Schulze J, Mühling KH, ***Tran LP** (2020). Is N-feedback involved in the regulation of nitrogenase activity in *Medicago truncatula*?. **J Plant Nutr Soil Sci** **183**:42-5.

76. Sulieman S, Kusano M, Ha CV, Watanabe Y, Abdalla MA, Abdelrahman M, Kobayashi M, Saito K, Mühling KH, ***Tran LS** (2019). Divergent metabolic adjustments in nodules are indispensable for efficient N₂ fixation of soybean under phosphate stress. **Plant Sci** **289**:110249.
77. Mostofa MG, Rahman MM, Ansary MMU, Fujita M, ***Tran LS** (2019). Interactive effects of salicylic acid and nitric oxide in enhancing rice tolerance to cadmium stress. **Int J Mol Sci** **20**:E5798.
78. Nguyen NC, Hoang XLT, Nguyen QT, Binh NX, Watanabe Y, Thao NP, ***Tran LS** (2019). Ectopic expression of *Glycine max GmNAC109* enhances drought tolerance and ABA sensitivity in *Arabidopsis*. **Biomolecules** **9**:E714.
79. Rahman MM, Mostofa MG, Rahman MA, Islam MR, Keya SS, Das AK, Miah MG, Kawser AQMR, Sowrav A, Hashem A, Tabassum B, Abd_Allah EF, ***Tran LS** (2019). Acetic acid: a cost-effective agent for mitigation of seawater-induced salt toxicity in mung bean. **Sci Rep** **9**:15186.
80. Shetty HS, Sharada MS, Jogaiah S, Rao A, Hansen M, Jørgensen HJL, ***Tran LS** (2019). Bioimaging structural signatures of the oomycete pathogen *Sclerospora graminicola* in pearl millet using different microscopic techniques. **Sci Rep** **9**:15175.
81. Salemi F, Nasr Esfahani M, ***Tran LS** (2019). Mechanistic insights into osmopriming-induced tolerance of early growth of alfalfa (*Medicago sativa* L.) under low water potential. **Ind Crops Prod** **19**:436-45.
82. Zahedi SM, Abdelrahman M, Hosseini MS, Hoveizeh NF, ***Tran LS** (2019). Alleviation of the effect of salinity on growth and yield of strawberry by foliar spray of selenium-nanoparticles. **Environ Pollut** **253**:246-58.
83. Tian L, Chang C, Ma L, Nasir F, Zhang J, Li W, ***Tran LS**, Tian C (2019). Comparative study of the mycorrhizal root transcriptomes of wild and cultivated rice in response to the pathogen *Magnaporthe oryzae*. **Rice** **12**:35.
84. Bao A, Chen H, Chen L, Chen S, Hao Q, Guo W, Qiu D, Shan Z, Yang Z, Yuan S, Zhang C, Zhang X, Li X, Liu B, Kong F, Zhou X, ***Tran LS**, Cao D (2019). CRISPR/Cas9-mediated targeted mutagenesis of *GmSPL9* genes alters plant architecture in soybean. **BMC Plant Biol** **19**:131.
85. Zhang P, Wang R, Ju Q, Li W, **Tran LS**, Xu J (2019). The R2R3-MYB transcription factor MYB49 regulates cadmium accumulation. **Plant Physiol** **180**:529-42.
86. Li W, Nguyen KH, Ha CV, Watanabe Y, ***Tran LS** (2019). Crosstalk between the cytokinin and MAX2 signaling pathways in growth and callus formation of *Arabidopsis thaliana*. **Biochem Biophys Res Commun** **511**:300-6.
87. Nguyen KH, Mostofa MG, Watanabe Y, Tran CD, Rahman MM, ***Tran LS** (2019). Overexpression of *GmNAC085* enhances drought tolerance in *Arabidopsis* by regulating glutathione biosynthesis, redox balance and glutathione-dependent detoxification of reactive oxygen species and methylglyoxal. **Environ Exp Bot** **161**:242-54.
88. Chang C, Tian L, Ma L, Weiqiang L, Nasir F, Li X, ***Tran LS**, Tian C (2019). Differential responses of molecular mechanisms and physiochemical characters in wild and cultivated soybeans against invasion by the pathogenic *Fusarium oxysporum* Schltdl. **Physiol Plant** **166**:1008-25.
89. Abdel Latef A, Mostofa MG, Rahman MM, Abdel-Farid IB, ***Tran LS** (2019). Extracts from yeast and carrot roots enhance maize performance under seawater-induced salt stress by altering physiol-biochemical characteristics of stressed plants. **J Plant Growth Regul** **38**:966-79.

90. Shabanian S, Nasr Esfahani M, Karamian R, ***Tran LS** (2019). Salicylic acid modulates cutting-induced physiological and biochemical responses to delay senescence in two gerbera cultivars. **Plant Growth Regul** 87:245-56.
91. Rahman MM, Mostofa MG, Rahman MA, Miah MG, Saha SR, Karim MA, Akter M, Keya SS, Islam M, ***Tran LS** (2019). Insight into salt tolerance mechanisms of the halophyte *Achras sapota*: an important fruit tree for agriculture in coastal areas. **Protoplasma** 256:181-91.
92. Jogaiah S, Kurjogi M, Abdelrahman M, Nagabhushana H, ***Tran LS** (2019). *Ganoderma applanatum*-mediated green synthesis of silver nanoparticles: structural characterization and *in vitro* and *in vivo* biomedical and agrochemical properties. **Arabian J Chemistry** 12:1108-20.

2018

93. Jemo M, Dhiba D, Hashem A, Abd_Allah EF, Alqarawi AA, ***Tran LS** (2018). Mycorrhizal fungal community structure in tropical humid soils under fallow and cropping conditions. **Sci Rep** 8:17061.
94. Tahjib-Ul-Arif M, Siddiqui MN, Sohag AAM, Sakil MA, Rahman MM, Polash MAS, Mostofa MG, ***Tran LS** (2018). Salicylic acid-mediated enhancement of photosynthesis attributes and antioxidant capacity contributes to yield improvement of maize plants under salt stress. **J Plant Growth Regul** 37:1318-30.
95. Chu HD, Nguyen KH, Watanabe Y, Le DT, Pham TLT, Mochida K, ***Tran LS** (2018). Identification, structural characterization and gene expression analysis of members of the nuclear factor-Y family in chickpea (*Cicer arietinum* L.) under dehydration and abscisic acid treatments. **Int J Mol Sci** 19:3290.
96. Tran CD, Chu HD, Nguyen KH, Watanabe Y, Tran KD, ***Tran LS** (2018). Genome-wide identification of the TCP transcription factor family in chickpea (*Cicer arietinum* L.) and their transcriptional responses to dehydration and exogenous abscisic acid treatments. **J Plant Growth Regul** 37:1286-99.
97. Shi S, Tian L, Nasir F, Li X, Li, W, ***Tran LS**, Tian C (2018). Impact of domestication on the evolution of rhizomicrobiome of rice in response to the presence of *Magnaporthe oryzae*. **Plant Physiol Biochem** 132:156-65.
98. Li W, Nishiyama R, Watanabe Y, Ha CV, Kojima M, An P, Tian C, Sakakibara H, ***Tran LS** (2018). Effects of overproduced ethylene on the contents of other phytohormones and expression of their key biosynthetic genes. **Plant Physiol Biochem** 128:170-7.
99. Le DT, Nguyen KL, Chu HD, Vu NT, Pham TTL, ***Tran LS** (2018). Functions of the evolutionarily conserved plant methionine-S-sulfoxide reductase without catalytic residue. **Protoplasma** 225:1741-50.
100. Tian L, Shi S, Ma L, Nasir F, Li X, ***Tran LS**, Tian C (2018). Co-evolutionary associations between root-associated microbiomes and root transcriptomes in wild and cultivated rice varieties. **Plant Physiol Biochem** 128:134-41.
101. Nguyen KH, Mostofa MG, Li W, Ha CV, Watanabe Y, Le DT, Thao NP, ***Tran LS** (2018). The soybean transcription factor GmNAC085 enhances drought tolerance in *Arabidopsis*. **Environ Exp Bot** 151:12-20.
102. Tian L, Shi S, Nasir F, Chang C, Li W, ***Tran LS**, Tian C (2018). Comparative analysis of the root transcriptomes of cultivated and wild rice varieties in response to *Magnaporthe grisea* infection revealed both common and species-specific pathogen responses. **Rice** 11:26.

103. Luo S, Tian L, Chang C, Wang S, Zhang J, Zhou X, ***Tran LS**, Tian C (2018). Grass and maize vegetation systems restore saline-sodic soils in the Songnen Plain of northeast China. **Land Degrad Develop** 29:1107-19.
104. Kurjogi M, Satapute P, Jogaiah S, Abdelrahman M, Daddam JR, Ramu V, ***Tran LS** (2018). Computational modeling of the staphylococcal enterotoxins and their interaction with natural antitoxin compounds. **Int J Mol Sci** 19:133.
105. Shabanian S, Nasr Esfahani M, Karamian R, ***Tran LS** (2018). Physiological and biochemical modifications by postharvest treatment with sodium nitroprusside extend vase life of cut flowers of two gerbera cultivars. **Postharvest Biol Technol** 137:1-8.
106. Kumari P, Singh S, Yadav S, ***Tran LS** (2018). Pretreatment of seeds with thidiazuron delimits its negative effects on explants and promotes regeneration in chickpea (*Cicer arietinum* L.). **Plant Cell Tissue Organ Cult** 133:103-14.
107. Jogaiah S, Abdelrahman M, **Tran LS**, Ito S (2018). Different mechanisms of *Trichoderma virens*-mediated resistance in tomato against *Fusarium* wilt involve the jasmonic and salicylic acid pathways. **Mol Plant Pathol** 19:870-82.
108. Abdel Latef A, Srivastava AK, El-sadek MSA, Kordrostami M, ***Tran LS** (2018). Titanium dioxide nanoparticles improve growth and enhance tolerance of broad bean plants under saline soil conditions. **Land Degrad Develop** 29:1065-73.

2017

109. Li W, Nguyen KH, Chu HD, Ha CV, Watanabe Y, Osakabe Y, Leyva-González MA, Sato M, Toyooka K, Voges L, Tanaka M, Mostofa MG, Seki M, Seo M, Yamaguchi S, Nelson DC, Herrera-Estrella L, ***Tran LS** (2017). The karrikin receptor KAI2 promotes drought resistance in *Arabidopsis thaliana*. **PLoS Genet** 13:e1007076 [highlighted in “Multifactorial response to drought”, **Science**, 2017, 358:1267 by Pamela J. Hines].
110. Jemo M, Sulieman S, Bekkaoui F, Olomide OAK, Hashem A, Abd_Allah EF, Alqarawi AA, ***Tran LS** (2017). Comparative analysis of the combined effects of different water and phosphate levels on growth and biological nitrogen fixation of nine cowpea varieties. **Front Plant Sci** 8:2111.
111. Siddiqui MN, Mostofa MG, Abu Sayed Md, Srivastava AS, Hasan MS, ***Tran LS** (2017). Impact of salt-induced toxicity on growth and yield-potential of local wheat cultivars: oxidative stress and ion toxicity are among the major determinants of salt-tolerant capacity. **Chemosphere** 187:385-94.
112. Abdel Latef A, Srivastava AK, Saber H, Alwaleed EA, ***Tran LS** (2017). *Sargassum muticum* and *Jania rubens* seaweeds regulate amino acid metabolism to improve growth and alleviate salinity in chickpea. **Sci Rep** 7:10537.
113. Nasr Esfahani M, Inoue K, Chu HD, Nguyen KH, Ha CV, Watanabe Y, Burritt DJ, Herrera-Estrella L, Mochida K, ***Tran LS** (2017). Comparative transcriptome analysis of nodules of two *Mesorhizobium*-chickpea associations with differential symbiotic efficiency under phosphate deficiency. **Plant J** 91:911-926.
114. Nguyen HM, Sako K, Matsui A, Suzuki Y, Mostofa MG, Ha CV, Tanaka M, **Tran LS**, Habu Y and Seki M (2017). Ethanol enhances high-salinity stress tolerance by detoxifying reactive oxygen species in *Arabidopsis thaliana* and rice. **Front Plant Sci** 8:1001.
115. Abdelrahman M, Ali H, El-Sayed M, Tanaka, ***Tran LS** (2017). Isolation and characterization of Cepa2, a natural alliospiroside A, from shallot (*Allium cepa* L. Aggregatum group) with anticancer activity. **Plant Physiol Biochem** 116:167-73.

116. Akram S, Siddiqui MN, Hussain BMN, Bari MAA, Mostofa MG, Hossain MA, ***Tran LS** (2017). Exogenous glutathione modulates salinity tolerance of soybean [*Glycine max* (L.) Merrill] at reproductive stage. **J Plant Growth Regul** 36:877-888.
117. Mostofa MG, Hossain MA, Siddiqui MN, Fujita M, ***Tran LS** (2017). Phenotypical, physiological and biochemical analyses provide insight into selenium-induced phytotoxicity in rice plants. **Chemosphere** 178:212-23.

2016

118. Govind SR, Jogaiah S, Abdelrahman M, Shetty HS, ***Tran LS** (2016). Exogenous trehalose treatment enhances the activities of defense-related enzymes and triggers resistance against downy mildew disease of pearl millet. **Front Plant Sci** 7:1593.
119. Jogaiah S, Kurjogi M, Govind SR, Hunthriker SS, Basappa VA, ***Tran LS** (2016). Isolation and evaluation of proteolytic actinomycete isolates as novel inducers of pearl millet downy mildew disease protection. **Sci Rep** 6:30789.
120. Nasr Esfahani M, Kusano M, Nguyen KH, Watanabe Y, Ha CV, Saito K, Sulieman S, Herrera-Estrella L, ***Tran LS** (2016). Adaptation of the symbiotic *Mesorhizobium*–chickpea relationship to phosphate deficiency relies on reprogramming of whole plant metabolism. **Proc Natl Acad Sci USA** 113:E4610-9.
121. Li W, Nguyen KH, Watanabe Y, Yamaguchi S, ***Tran LS** (2016). *OaMAX2* of *Orobanchae aegyptiaca* and *Arabidopsis AtMAX2* share conserved functions in both development and drought responses. **Biochem Biophys Res Commun** 478:521-6.
122. Chu HD, Nguyen KL, Watanabe Y, Le DT, ***Tran LS** (2016). Expression analyses of soybean genes encoding methionine-*R*-sulfoxide reductase under various conditions suggest a possible role in the adaptation to stress. **Appl Biol Chem** 59:681-7.
123. Jogaiah S, Shetty HS, Ito S, ***Tran LS** (2016). Enhancement of downy mildew disease resistance in pearl millet by the G_app7 bioactive compound produced by *Ganoderma applanatum*. **Plant Physiol Biochem** 105: 109-17.
124. Ahmad P, Abdel Latef A, Hashem A, Abd_Allah E, Gucel S, ***Tran LS** (2016). Nitric oxide mitigates salt stress by regulating levels of osmolytes and antioxidant enzymes in chickpea. **Front Plant Sci** 7:347.
125. Abdelrahman M, Abdel-Motaal F, El-Sayed M, Jogaiah S, Shigyo M, Ito S, ***Tran LS** (2016). Dissection of *Trichoderma longibrachiatum*-induced defense in onion (*Allium cepa* L.) against *Fusarium oxysporum* f. sp. *cepa* by target metabolite profiling. **Plant Sci** 246:128-38.
126. Abdel Latef A, ***Tran LS** (2016). Impacts of priming with silicon on the growth and tolerance of maize plants to alkaline stress. **Front Plant Sci** 7:243.
127. Nguyen KH, Ha CV, Nishiyama R, Watanabe Y, Leyva-González MA, Fujita Y, Tran UT, Tanaka M, Li W, Seki M, Schaller GE, Herrera-Estrella L, ***Tran LS** (2016). *Arabidopsis* type B cytokinin response regulators ARR1, ARR10, and ARR12 negatively regulate plant responses to drought. **Proc Natl Acad Sci USA** 113:3090-5.
128. Li W, Yamaguchi S, Khan AM, An P, Liu X, ***Tran LS** (2016). Roles of gibberellins and abscisic acid in regulating germination of *Suaeda salsa* dimorphic seeds under salt stress. **Front Plant Sci** 6:1235.

2015

129. Mostofa MG, Saegusa D, Fujita M, ***Tran LS** (2015). Hydrogen sulfide regulates salt tolerance in rice by maintaining Na⁺/K⁺ balance, mineral homeostasis and oxidative metabolism under excessive salt stress. **Front Plant Sci** 6:1055.

130. Mostofa MG, Ansary MU, Watanabe A, Fujita M, ***Tran LS** (2015). Hydrogen sulfide modulates cadmium-induced physiological and biochemical responses to alleviate cadmium toxicity in rice. **Sci Rep** 5:14078.
131. Mao H, Wang H, Liu S, Li Z, Yang X, Yan J, Li J, **Tran LS**, Qin F (2015). A miniature inverted-repeat transposable element in a *NAC* gene is associated with drought tolerance in maize. **Nat Commun** 6:8326.
132. Ha CV, Watanabe Y, Tran UT, Le DT, Tanaka M, Nguyen KH, Seki M, Nguyen DV, ***Tran LS** (2015). Comparative analysis of root transcriptomes from two contrasting drought-responsive Williams 82 and DT2008 soybean cultivars revealed differentially expressed genes under dehydration stress. **Front Plant Sci** 6:551.
133. Nguyen KH, Ha CV, Watanabe Y, Tran UT, Esfahani MN, Nguyen DV, ***Tran LS** (2015). Correlation between differential drought tolerability of two contrasting drought-responsive chickpea cultivars and differential expression of a subset of *CaNAC* genes under normal and dehydration conditions. **Front Plant Sci** 6:449.
134. Mostofa MG, Hossain MA, Fujita M, ***Tran LS** (2015). Physiological and biochemical mechanisms associated with trehalose-induced copper stress tolerance in rice. **Sci Rep** 5:11433.
135. Du H, Liang Z, Zhao S, Nan M-G, **Tran LS**, Lu K, Huang Y-B, Li J-N (2015). The evolutionary history of R2R3-MYB proteins across 50 eukaryotes: new insights into subfamily classification and expansion. **Sci Rep** 5:11037.
136. Mostofa MG, Fujita M, ***Tran LS** (2015). Nitric oxide mediates hydrogen peroxide- and salicylic acid-induced salt tolerance in rice (*Oryza sativa* L.) seedlings. **Plant Growth Regul** 77:265-277.
137. Ahmad P, Sarwat M, Bhat NA, Wani MR, Kazi AG, ***Tran LS** (2015). Alleviation of cadmium toxicity in *Brassica juncea* L. (Czern. & Coss.) by calcium application involves various physiological and biochemical strategies. **PLoS One** 10:e0114571.
138. Narendra Babu A, Jogaiah S, Ito S, Nagaraj AK, ***Tran LS** (2015). Improvement of growth, yield and early blight disease resistance of tomato plants by rhizosphere bacteria is correlated with their beneficial traits and induced biosynthesis of antioxidant peroxidase and polyphenol oxidase. **Plant Sci** 231:62-73.
139. Sulieman S, Ha CV, Esfahani MN, Watanabe Y, Nishiyama R, Pham CTB, Nguyen DV, ***Tran LS** (2015). DT2008: A promising new genetic resource for improved drought tolerance in soybean when solely dependent on symbiotic N₂ fixation. **Biomed Res Int** 2015:687213.
140. Thu NBA, Hoang TLX, Dung NHT, Thao NP, ***Tran LS** (2015). Differential expression of two-component system-related drought-responsive genes in two contrasting drought-tolerant soybean cultivars DT51 and MTD720 under well-watered and drought conditions. **Plant Mol Biol Report** 33:1569-610.

2014

141. Ha CV, Esfahani MN, Watanabe Y, Sulieman S, Mochida K, Nguyen DV, ***Tran LS** (2014). Genome-wide identification and expression analysis of the *CaNAC* family members in chickpea during development, dehydration and ABA treatments. **PLoS One** 9:e114107
142. Guttikonda SK, Valliyodan B, Neelakandan AK, **Tran LS**, Kumar R, Quach TN, Voothuluru P, Guttierrez-Gonzalez JJ, Aldrich DL, Pallardy SG, Sharp RE, Ho THD, Nguyen HT (2014). Overexpression of *AtDREB1D* transcription factor improves drought tolerance in soybean. **Mol Biol Rep** 41:7995-8008.

143. Nasr Esfahani M, Sulieman S, Schulze J, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2014). Mechanisms of physiological adjustment of N₂ fixation in chickpea (*Cicer arietinum* L.) during early stages of water deficit: single or multi-factor controls. **Plant J** 79:964-80.
144. Thu NBA, Hoang TLX, Doan H, Nguyen TH, Bui D, Thao NP, ***Tran LS** (2014). Differential expression analysis of a subset of *GmNAC* genes in shoots of two contrasting drought-responsive soybean cultivars DT51 and MTD720 under normal and drought conditions. **Mol Biol Rep** 41:5563-9.
145. Thu NBA, Nguyen QT, Hoang TLX, Thao NP, ***Tran LS** (2014). Evaluation of drought tolerance of the Vietnamese soybean cultivars provides potential genetic resources for genetic engineering. **Biomed Res Int** 2014:809736.
146. Sulieman S, Schulze J, ***Tran LS** (2014). N-feedback regulation is synchronized with nodule carbon alteration in *Medicago truncatula* under excessive nitrate or low phosphorus conditions. **J Plant Physiol** 171:407-10.
147. Quach TN, **Tran LS**, Valliyodan B, Nguyen HT, Kumar R, Neelakandan AK, Guttikonda SK, Sharp RE, Nguyen HT (2014). Functional analysis of water stress-responsive soybean *GmNAC003* and *GmNAC004* transcription factors in lateral root development in *Arabidopsis*. **PLoS One** 9:e84886.
148. Ha CV, Leyva-Gonzalez MA, Osakabe Y, Tran TU, Nishiyama R, Watanabe Y, Tanaka M, Seki M, Yamaguchi S, Dong NV, Yamaguchi-Shinozaki K, Shinozaki K, Herrera-Esterella L, ***Tran LS** (2014). Positive regulatory role of strigolactone in plant responses to drought and salt stress. **Proc Natl Acad Sci USA** 111:851-6.
149. Esfahani MN, Sulieman S, Schulze J, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2014). Approaches for enhancement of N₂ fixation efficiency of chickpea (*Cicer arietinum* L.) under limiting nitrogen conditions. **Plant Biotech J** 12:387-97.

2013

150. Thao NP, Thu NBA, Hoang TLX, ***Tran LS** (2013). Differential expression analysis of a subset of drought-responsive *GmNAC* genes in two soybean cultivars differing in drought tolerance. **Int J Mol Sci** 14:23828-41.
151. Liu S, Wang X, Wang H, Xin H, Yang X, Yan J, Li J, **Tran LS**, Shinozaki K, Yamaguchi-Shinozaki K, Qin F (2013). Genome-wide analysis of *ZmDREB* genes and their association with natural variation in drought tolerance at seedling stage of *Zea mays* L. **PLoS Genet** 9:e1003790.
152. Ha CV, Le DT, Nishiyama R, Watanabe Y, Sulieman S, Tran UT, Mochida K, Dong NV, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2013). The auxin response factor transcription factor family in soybean: genome-wide identification and expression analyses during development and water stress. **DNA Res** 20:511-24.
153. Jogaiah S, Mostafa A, **Tran LS**, Ito S (2013). Characterization of rhizosphere fungi that mediate resistance in tomato against bacterial wilt disease. **J Exp Bot** 64:3829-42.
154. Le DT, Tarrago L, Watanabe Y, Kaya A, Lee BC, Tran UT, Nishiyama R, Fomenko DE, Gladyshev VN, ***Tran LS** (2013). Diversity of plant methionine sulfoxide reductases b and evolution of a form specific for free methionine sulfoxide. **PloS One** 8:e65637
155. Sulieman S, Ha CV, Schulze J, ***Tran LS** (2013). Growth and nodulation of symbiotic *Medicago truncatula* at different levels of phosphorus availability. **J Exp Bot** 64:2701-12.
156. Nishiyama R, Watanabe Y, Leyva-Gonzalez MA, Ha CV, Fujita Y, Tanaka M, Seki M, Yamaguchi-Shinozaki K, Shinozaki K, Herrera-Estrella L, ***Tran LS** (2013). *Arabidopsis*

- AHP2, AHP3 and AHP5 histidine phosphotransfer proteins function as redundant negative regulators of drought stress response. **Proc Natl Acad Sci USA** **110**:4840-5.
157. Sulieman S, Schulze J, ***Tran LS** (2013). Comparative analysis of the symbiotic efficiency of *Medicago truncatula* and *Medicago sativa* under phosphorus deficiency. **Int J Mol Sci** **14**:5198-213.
158. Mochida K, Yoshida T, Sakurai T, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2013). TreeTFDB: an integrative database of the transcription factors from six economically important tree crops for functional predictions and comparative and functional genomics. **DNA Res** **20**:151-62.
159. Ha CV, Le DT, Nishiyama R, Watanabe Y, Dong NV, Tran UT, ***Tran LS** (2013). Characterization of the newly developed soybean cultivar DT2008 in relation to the model variety W82 reveals a new genetic resource for comparative and functional genomics for improved drought tolerance. **Biomed Res Int** **2013**:759657

2012

160. Le DT, Nishiyama R, Watanabe Y, Tanaka M, Seki K, Ham HL, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2012). Differential gene expression in soybean leaf tissues at late developmental stages under drought stress revealed by genome-wide transcriptome analysis. **PLoS One** **7**:e49522.
161. Nam YJ, **Tran LS**, Kojima M, Sakakibara H, Nishiyama R, Shin R (2012). Regulatory roles of cytokinins and cytokinin signaling in response to potassium deficiency in *Arabidopsis*. **PLoS One** **7**:e47797.
162. Le DT, Aldrich DL, Valliyodan B, Watanabe Y, Ha CV, Nishiyama R, Guttikonda SK, Quach TN, Gutierrez-Gonzalez JJ, ***Tran LS**, Nguyen HT (2012). Evaluation of candidate reference genes for normalization of quantitative RT-PCR in soybean tissues under various abiotic stress conditions. **PLoS One** **7**:e46487.
163. Choudhary PS, Oral HV, Bhardwaj R, Yu J-Q, ***Tran LS** (2012). Interaction of brassinosteroids and polyamines enhances copper stress tolerance in *Raphanus sativus*. **J Exp Bot** **63**:5659-75.
164. Le DT, Nishiyama R, Watanabe Y, Vankova R, Tanaka M, Seki K, Ham HL, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2012). Identification and expression analysis of cytokinin metabolic genes in soybean under normal and drought conditions in relation to cytokinin levels. **PLoS One** **7**:e42411.
165. Choudhary PS, Kanwar M, Bhardwaj R, Yu J-Q, ***Tran LS** (2012). Chromium stress mitigation by polyamine-brassinosteroid application involves phytohormonal and physiological strategies in *Raphanus sativus* L. **PLoS One** **7**:e33210.
166. Nishiyama R, Le DT, Watanabe Y, Matsui A, Tanaka M, Seki M, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2012). Transcriptome analyses of a salt-tolerant cytokinin-deficient mutant reveal differential regulation of salt stress response by cytokinin deficiency. **PLoS One** **7**:e32124.
167. Kumar R, **Tran LS**, Neelakandan AK, Nguyen HT (2012). Higher plant cytochrome *b5* polypeptides modulate fatty acid desaturation. **PLoS One** **7**:e31370.

2011

168. Qin F, Kodaira KS, Maruyama K, Mizoi J, **Tran LS**, Fujita Y, Morimoto K, Shinozaki K, Yamaguchi-Shinozaki K (2011). *SPINDLY*, a negative regulator of gibberellic acid signaling, is involved in the plant abiotic stress response. **Plant Physiol** **157**:1900-13.

169. Kodaira KS, Qin F, **Tran LS**, Maruyama K, Fujita Y, Kidokoro S, Shinozaki K, Yamaguchi-Shinozaki K (2011). The *Arabidopsis* Cys2/His2-type zinc-finger proteins AZF1 and AZF2 negatively regulate ABA-repressive and auxin-inducible genes as transcriptional repressors under abiotic stress conditions. **Plant Physiol** **157**:742-56.
170. Nishiyama R, Watanabe Y, Fujita Y, Le DT, Kojima M, Werner T, Vankova R, Yamaguchi-Shinozaki K, Shinozaki K, Kakimoto T, Sakakibara H, Schmölling T, ***Tran LS** (2011). Analysis of cytokinin mutants and regulation of cytokinin metabolic genes reveals important regulatory roles of cytokinins in drought, salt and ABA responses, and ABA biosynthesis. **Plant Cell** **23**:2169-83.
171. Mochida K, Yoshida T, Sakurai T, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2011). *In silico* analysis of transcription factor repertoires and prediction of stress-responsive transcription factors from six major Gramineae plants. **DNA Res** **18**:321-32. (Editor's choice in the issue)
172. Le DT, Nishiyama R, Watanabe Y, Mochida K, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2011). Genome-wide survey and expression analysis of the plant-specific NAC transcription factor family in soybean during development and dehydration stress. **DNA Res** **18**: 263-76.
173. Le DT, Nishiyama R, Watanabe Y, Mochida K, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2011). Genome-wide expression profiling of soybean two-component system genes in soybean root and shoot tissues under dehydration stress. **DNA Res** **18**:17-29.
174. Kimura K, **Tran LS**, Funane K (2011). Loss of poly- γ -glutamic acid synthesis of *Bacillus subtilis* (natto) due to IS4*BsuI* translocation to *swrA* gene. **Food Sci Technol Res** **17**:447-51.

2010

175. Neelakandan AK, Nguyen HT, Kumar R, **Tran LS**, Guttikonda SK, Quach TN, Aldrich DL, Nes WD, Nguyen HT (2010). Molecular characterization and functional analysis of *Glycine max* sterol methyl transferase 2 genes involved in plant membrane sterol biosynthesis. **Plant Mol Biol** **74**:503-18.
176. Mochida K, Yoshida T, Sakurai T, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2010). Genome-wide analysis of two-component system and prediction of stress-responsive TCS members in soybean. **DNA Res** **17**:303-24.
177. Gutierrez-Gonzalez JJ, Guttikonda SK, **Tran LS**, Aldrich DL, Zhong R, Yu O, Nguyen HT, Sleper AD (2010). Differential expression of isoflavone biosynthetic genes in soybean during water deficits. **Plant Cell Physiol** **51**:936-48.
178. Mochida K, Yoshida T, Sakurai T, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2010). LegumeTFDB: An integrative database of *Glycine max*, *Lotus japonicus* and *Medicago truncatula* transcription factors. **Bioinformatics** **26**:290-91.

2009 and before

179. Mochida K, Yoshida T, Sakurai T, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2009). *In silico* analysis of transcription factor repertoire and prediction of stress responsive transcription factors in soybean. **DNA Res** **16**:353-69.
180. ***Tran LS**, Quach T, Guttikonda SK, Aldrich DL, Rajesh K, Neelakandan A, Valliyodan B, Nguyen HT (2009). Molecular characterization of stress-inducible *GmNAC* genes in soybean. **Mol Gen Genomics** **281**:647-64.

181. Kimura K, **Tran LS**, Do TH, Itoh Y (2009). Expression of the *pgsB* encoding the poly- γ -DL-glutamate synthetase of *Bacillus subtilis* (natto). **Biosci Biotechnol Biochem** **73**:1149-55.
182. Qin F, Sakuma Y, **Tran LS**, Maruyama K, Kikodoro S, Fujita Y, Fujita M, Sawano Y, Miyazono K, Tanokura M, Shinozaki K, Yamaguchi-Shinozaki K (2008). *Arabidopsis* DREB2A interacting proteins function as RING E3 ligases and negatively regulate plant drought stress responsive gene expression. **Plant Cell** **20**:1693-07.
183. **Tran LS**, Urao T, Qin F, Maruyama K, Kakimoto T, Shinozaki K, Yamaguchi-Shinozaki K (2007). Functional analysis of AHK1/ATHK1 and cytokinin receptor histidine kinases in response to ABA, drought and high salinity stresses in *Arabidopsis*. **Proc Natl Acad Sci USA** **104**:20623-8.
184. Nakashima K, **Tran LS**, Nguyen VD, Fujita M, Maruyama K, Todaka D, Ito Y, Hayashi N, Shinozaki K, Yamaguchi-Shinozaki K (2007). Functional analysis of NAC-type transcription factor OsNAC6 involved in abiotic and biotic stress-responsive gene expression in rice. **Plant J** **51**:617-30.
185. Qin F, Kakimoto M, Sakuma Y, Osakabe Y, Maruyama K, **Tran LS**, Shinozaki K, Yamaguchi-Shinozaki K (2007). Regulation and functional analysis of *ZmDREB2A* in response to drought and heat stresses in *Zea mays* L. **Plant J** **50**:54-69.
186. **Tran LS**, Nakashima K, Sakuma Y, Osakabe Y, Qin F, Simpson SD, Maruyama K, Fujita Y, Shinozaki K, Yamaguchi-Shinozaki K (2006). Co-expression of the stress-inducible zinc finger homeodomain ZFHD1 and NAC transcription factors enhances expression of the *ERD1* gene in *Arabidopsis*. **Plant J** **49**:46-63.
187. **Tran LS**, Nakashima K, Sakuma Y, Simpson SD, Fujita Y, Maruyama K, Fujita M, Seki M, Shinozaki K, Yamaguchi-Shinozaki K (2004). Isolation and functional analysis of *Arabidopsis* stress-inducible NAC transcription factors that bind to a drought-responsive *cis*-element in the *early responsive to dehydration stress 1* promoter. **Plant Cell** **16**:2481-98.
188. Fujita M, Fujita F, Maruyama K, Seki M, Hiratsu K, Ohme-Takagi M, **Tran LS**, Yamaguchi-Shinozaki K, Shinozaki K (2004). A dehydration-induced NAC protein, RD26, is involved in a novel ABA-dependent stress-signaling pathway. **Plant J** **39**:863-76.
189. Kimura K, **Tran LS**, Uchida I, Itoh Y (2004). Characterization of *Bacillus subtilis* γ -glutamyltransferase and its involvement in the degradation of capsule poly- γ -glutamate. **Microbiology** **150**: 4115-23.
190. Kimura K, **Tran LS**, Itoh Y (2004). Roles and regulation of the glutamate racemase isogenes, *racE* and *yprC*, in *Bacillus subtilis*. **Microbiology** **150**:2911-20.
191. **Tran LP**, Nagai T, Itoh Y (2000). Divergent structure of the *comQXPA* quorum-sensing components: molecular basis of strain-specific communication mechanism in *Bacillus subtilis*. **Mol Microbiol** **37**:1159-71.
192. Nagai T, **Tran LS**, Inatsu Y, Itoh Y (2000). A new IS4 family insertion sequence, *IS4Bsu1*, responsible for genetic instability of poly- γ -glutamic acid production in *Bacillus subtilis*. **J Bacteriol** **182**:2387-92.
193. **Tran LS**, Szabo L, Ponyi T, Orosz L, Sik T, Holczinger A (1999). Phage abortive infection of *Bacillus licheniformis* ATCC 9800; Identification of the *abiBL11* gene, localisation and sequencing of its promoter region. **Appl Microbiol Biotechnol** **52**:845-52.
194. ***Tran LS**, Szabo L, Orosz L, Sik T, Holczinger A (1998). Construction of a single-copy integration vector and its use to study gene expression in *Bacillus licheniformis*. **Microbiology** **144**:2573-8.

195. ***Tran LS**, Szabo L, Fulop L, Orosz L, Sik T, Holczinger A (1998). Isolation of a β -galactosidase-encoding gene from *Bacillus licheniformis*. Purification and characterization of the recombinant enzyme expressed in *Escherichia coli*. **Curr Microbiol** 37:39-43.
196. Fulop L, **Tran LS**, Pragai Z, Felfoldi F, Ponyi T (1996). Cloning and expression of a β -1,4-endoglucanase gene from *Cellulomonas* sp. CelB7 in *Escherichia coli*; purification and characterisation of the recombinant enzyme. **FEMS Microbiol Letters** 145: 355-60.
197. Pragai Z, **Tran SL**, Nagy T, Fulop L, Holczinger A, Sik T (1994). Transposon Tn917PFI mutagenesis in *Bacillus licheniformis*. **Microbiology** 140:3091-7.

III. Reviews, Addenda, Book Chapters, Book & Special Issue Editions

Reviews

2023

198. Khan TA, Kappachery S, Karumannil S, AlHosani M, Almansoori N, Almansoori H, Yusuf M, ***Tran LP**, Gururani MA (2023). Brassinosteroid signaling pathways: Insights into plant responses under abiotic stress. **Int J Mol Sci** 24: 17246.
199. Sharma A, Gupta A, Ramakrishnan M, Ha CV, Zheng B, Bhardwaj M, ***Tran LP** (2023). Roles of abscisic acid and auxin in plants during drought: A molecular point of view. **Plant Physiol Biochem** 204:108129.
200. Liu S, Dong Y, McConkey KR, **Tran LP**, Wang F, Liu H, Wu G (2023). Scientific concept and practices of life community of mountains, rivers, forests, farmlands, lakes, grasslands, and deserts in China. **Ambio** 52:1939-51.
201. Kandhol N, Pandey S, Singh VP, Herrera-Estrella L, Bucop JL, ***Tran LP**, Tripathi DK (2023). Bacterial community and root endodermis: a complementary relationship. **Trends Plant Sci** 28:749-51.
202. Hossain MM, Sultana F, Li W, ***Tran LP**, Mostofa MG (2023) *Sclerotinia sclerotiorum* (Lib.) de Bary: Insights into the pathogenomic features of a global pathogen. **Cells** 12:1063.
203. Kandhol N, Singh VP, Herrera-Estrella L, ***Tran LP**, Tripathi DK (2023). Nanocarrier spray: A non-transgenic approach for crop engineering. **Trends Plant Sci** 28:259-61.

2022

204. Gupta A, Bhardwaj M, ***Tran LP** (2022). Integration of auxin, brassinosteroid and cytokinin in the regulation of rice yield. **Plant Cell Physiol** 63:1848-56.
205. Kandhol N, Singh VP, White JC, ***Tran LP**, Tripathi DK (2022). Plant growth hormones and nanomaterial interface: Linking development to defence. **Plant Cell Physiol** 63:1840-47.
206. Tripathi DK, Yadav SR, Mochida K, ***Tran LP** (2022). Plant growth regulators: true managers of plant life. **Plant Cell Physiol** 63:1757-60.
207. Abdelrahman M, Tran L-SP, Shigyo M (2022) Editorial: Physiological and molecular perspectives of stress tolerance in vegetables. **Front Plant Sci** 13:1004093.
208. Sun H, Li W, Burritt JD, Tian H, Zhang H, Liang X, Miao Y, Mostofa MG, ***Tran LP** (2022). Strigolactones interact with other phytohormones to modulate plant root growth and development. **The Crop Journal** 10:1517-27.

209. Mostofa MG, Rahman MM, Ghosh TK, Kabir AH, Abdelrahman M, Khan AMR, Mochida K, ***Tran LP** (2022). Potassium in plant physiological adaptation to abiotic stresses. **Plant Physiol Biochem** **186**:279-89.
210. Alolga RN, Wang F, Zhang X, Li J, ***Tran L-SP**, Yin X (2022). Bioactive compounds from the Zingiberaceae family with known antioxidant activities for possible therapeutic uses. **Antioxidants** **11**:1281.
211. Kandhol N, Singh VP, Herrera-Estrella L, ***Tran LP**, Tripathi DK (2022). Arsenite: the umpire of arsenate perception and responses in plants. **Trends Plant Sci** **27**:420-2.
212. Sulieman S, Abdelrahman M, ***Tran LP** (2022). Carbon metabolic adjustment in soybean nodules in response to phosphate limitation: A metabolite perspective. **Environ Exp Bot** **196**:104810.

2021

213. Gupta A, Bhardwaj M, ***Tran LP** (2021). The JASMONATE ZIM-DOMAIN family proteins are the important node in jasmonic acid-abscisic acid crosstalk in regulating plant response to drought. **Curr Protein Pept Sci** **22**:759-66.
214. Rahman M, Mostofa MG, Keya SS, Siddiqui MN, Ansary MMU, Das AK, Rahman MA, ***Tran LP** (2021). Adaptive mechanisms of halophytes and their potential in improving salinity tolerance in plants. **Int J Mol Sci** **22**:10733.
215. Saha G, Mostofa MG, Rahman MM, ***Tran LP** (2021). Silicon-mediated heat tolerance in higher plants: A mechanistic outlook. **Plant Physiol Biochem** **166**:341-7.
216. Hoang XLT, Prerostova S, Thu NBA, Thao NP, Vankova R, ***Tran LP** (2021). Histidine kinases: diverse functions in plant development and responses to environmental conditions. **Annu Rev Plant Biol** **72**:297-323.
217. Mostofa MG, Rahman MM, Ansary MMU, Keya SS, Abdelrahman M, Miah MG, ***Tran LP** (2021). Silicon in mitigation of abiotic stress-induced oxidative damage in plants. **Crit Rev Biotechnol** **41**:918-34.

2020

218. Patel MK, Kumar M, Li W, Luo Y, Burritt DJ, Alkan N, ***Tran LP** (2020). Enhancing salt tolerance of plants: from metabolic reprogramming to exogenous chemical treatments and molecular approaches. **Cells** **9**:E2492.
219. Nasir F, Li W, ***Tran LP**, Tian J (2020). Does karrikin signaling shape the rhizomicrobiome via the strigolactone biosynthetic pathway? **Trends Plant Sci** **25**:1184-87.
220. Guo W, Chen L, Herrera-Estrella L, Cao D, ***Tran LP** (2020). Altering plant architecture to improve performance and resistance. **Trends Plant Sci** **25**:1154-70.
221. Gupta A, Bhardwaj M, ***Tran LP** (2020). Jasmonic acid at the crossroads of plant immunity and *Pseudomonas syringae* virulence. **Int J Mol Sci** **21**:7482.
222. Abdelrahman M, Ishii T, El-Sayed M, ***Tran LP** (2020). Heat sensing and lipid reprogramming as a signaling switch for heat stress responses in wheat. **Plant Cell Physiol** **61**:1399-1407.
223. Zhang B, Jin X, Yin H, Zhang D, Zhou H, Zhang X, ***Tran LP** (2020). Natural products, traditional uses and pharmacological activities of the genus *Biebersteinia* (Biebersteiniaceae) **Plants** **9**:595.
224. Tian L, Lin X, Tian J, Ji L, Chen Y, ***Tran LP**, Tian C (2020). Research advances of beneficial microbiota associated with crop plants. **Int J Mol Sci** **21**:E1792.

225. Gupta A, Sinha R, Fernandes JL, Abdelrahman M, Burritt DJ, ***Tran LP** (2020). Phytohormones regulate convergent and divergent responses between individual and combined drought and pathogen infection. **Crit Rev Biotechnol** 40:320-40.
226. Abdelrahman M, Burritt DJ, Gupta A, ***Tran LP** (2020). Heat stress effects on source-sink relationships and metabolome dynamics in wheat. **J Ext Bot** 71:543-54.

2019

227. Li W, Herrera-Estrella L, ***Tran LS** (2019). Do cytokinins and strigolactones crosstalk during drought adaptation?. **Trends Plant Sci** 24:669-72.
228. Patel MK, Pandey S, Burritt DJ, ***Tran LS** (2019). Plant responses to low-oxygen stress: interplay between ROS and NO signaling pathways. **Environ Exp Bot** 161:134-42.
229. Bao A, Chen H, Zhou X, Cao D, ***Tran LS** (2019). CRISPR/Cas9 system and its applications in crop genome editing. **Crit Rev Biotechnol** 39:321-36.

2018

230. Abiala MA, Abdelrahman M, Burritt DJ, ***Tran LS** (2018). Salt stress tolerance mechanisms and potential applications of legumes for sustainable reclamation of salt-degraded soils. **Land Degrad Develop** 29:3812-22.
231. Mostofa MG, Weiqiang L, Nguyen KH, Fujita M, ***Tran LS** (2018). Strigolactones in plant adaptation to abiotic stresses: an emerging avenue of plant research. **Plant Cell Environ** 41:2227-43.
232. Abdelrahman M, Al-Sadi AM, Pour-Aboughadareh A, Burritt DJ, ***Tran LS** (2018). Genome editing using CRISPR/Cas9-targeted mutagenesis: an opportunity for yield improvements of crop plants grown under environmental stresses. **Plant Physiol Biochem** 131:31-6.
233. Mostofa MG, Ghosh A, Fujita M, ***Tran LS** (2018). Methylglyoxal – a signaling molecule in plant abiotic stress responses. **Free Radic Biol Med** 122:96-109.
234. Abdelrahman M, Jogaiah S, Burritt DJ, ***Tran LS** (2018). Legume genetic resources and transcriptome dynamics under abiotic stress conditions. **Plant Cell Environ** 41:1972-83.
235. Srivastava S, Upadhyay MK, Srivastava AK, Abdelrahman M, Suprasanna P, ***Tran LS** (2018). Cellular and subcellular phosphate transport machinery in plants. **Int J Mol Sci** 19:1914.
236. Abdelrahman M, El-Sayed MA, Hashem A, Abd_Allah EF, Alqarawi AA, Burritt DJ, ***Tran LS** (2018). Metabolomics and transcriptomics in legumes under phosphate deficiency in relation to nitrogen fixation by root nodules. **Front Plant Sci** 9:922.
237. Cao D, Li Y, Liu B, Kong F, ***Tran LS** (2018). Adaptive mechanisms of soybean grown on salt-affected soils. **Land Degrad Develop** 29:1054-64.
238. Nasir F, Tian L, Chang C, Gao Y, ***Tran LS**, Tian C (2018). Toward understanding mechanistic defense of rice (*Oryza sativa*) in response to *Magnaporthe oryzae*'s infection. **Semin Cell Dev Biol** 83:95-105.
239. Abdelrahman M, Burritt DJ, ***Tran LS** (2018). The use of metabolomic quantitative trait locus mapping and osmotic adjustment traits for the improvement of crop yields under environmental stresses. **Semin Cell Dev Biol** 83:86-94.

2017

240. Abdelrahman M, El-Sayed M, Jogaiah S, Burritt DJ, ***Tran LS** (2017). "STAY-GREEN" trait and phytohormone signaling network under heat stress. **Plant Cell Rep** 36:1009-25.

241. Hoang XLT, Nhi DNH, Thu NBA, Thao NP, ***Tran LS** (2017). Transcription factors and their roles in signal transduction in plants under abiotic stresses. **Curr Genomics** **18**:483-497.
242. AbuQamar S, Moustafa K, ***Tran LS** (2017). Mechanisms and strategies of plant defense against *Botrytis cinerea*. **Crit Rev Biotechnol** **37**:262-74.

2016

243. AbuQamar S, Moustafa K, ***Tran LS** (2016). ‘Omics’ and plant responses to *Botrytis cinerea*. **Front Plant Sci** **7**:1658.
244. Hoque TS, Hossain MA, Mostofa MG, Burritt DJ, Fujita M, ***Tran LS** (2016). Methylglyoxal: a potent signaling molecule in plant abiotic stress responses and tolerance. **Front Plant Sci** **7**:1341.
245. Li W, Herrera-Estrella L, ***Tran LS** (2016). The yin-and-yang of cytokinin homeostasis and drought acclimation/adaptation. **Trends Plant Sci** **21**:548-50.
246. Guihéneuf F, Khan A, ***Tran LS** (2016). Genetic engineering: a promising tool to engender physiological, biochemical, and molecular stress resilience in green microalgae. **Front Plant Sci** **7**:420.
247. Singh A, Pandey A, Srivastava AK, ***Tran LS**, Girdhar PK (2016). Plant protein phosphatase 2C: genomic diversity to functional multiplicity and importance in stress management. **Crit Rev Biotechnol** **36**:1023-35.
248. Sytar O, Brestic M, Zivcak M, **Tran LS** (2016). The contribution of buckwheat genetic resources to health and dietary diversity. **Curr Genomics** **17**:193-206.
249. Srivastava AK, Suprasanna P, ***Tran LS** (2016). Multifaceted role of aquaporins as molecular conduits in plant responses to abiotic stresses. **Crit Rev Biotechnol** **36**:389-98.

2015

250. Li W, ***Tran LS** (2015). Are karrikins involved in plant abiotic stress responses?. **Trends Plant Sci** **20**:535-8.
251. Thao NP, Khan MIR, Thu NBA, Hoang XLT, Asgher M, Khan NA, ***Tran LS** (2015). Role of ethylene and its crosstalk with other signaling molecules in plant responses to heavy metal stress. **Plant Physiol** **169**:73-84.
252. Sulieman S, ***Tran LS** (2015). Phosphorus homeostasis in legume nodules as an adaptive strategy to phosphorus deficiency. **Plant Sci** **239**:36-43.
253. Hossain MA, Bhattachajee S, Armin SM, Burritt DJ, Fujita M, ***Tran LS** (2015). Hydrogen peroxide-priming modulates abiotic oxidative stress tolerance: insights from ROS detoxification and scavenging. **Front Plant Sci** **6**:420.
254. Gururani MA, Venkatesh J, ***Tran LS** (2015). Regulation of photosynthesis during abiotic stress-induced photoinhibition. **Mol Plant** **8**:1304-20.

2014

255. Osakabe Y, Osakabe K, Shinozaki K, ***Tran LS** (2014). Response of plant to water stress. **Front Plant Sci** **5**:86.
256. Osakabe Y, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2014). ABA control of plant macroelement membrane transport systems in response to water deficit and high salinity. **New Phytol** **202**:35-49 (Tansley review).

257. Ha S, ***Tran LS** (2014). Understanding plant responses to phosphorus starvation for improvement of plant tolerance to phosphorus deficiency by biotechnological approaches. **Crit Rev Biotechnol** 34:16-30.

2013

258. Osakabe Y, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2013). Sensing the environment: key roles of membrane-localized kinases in plant perception and response to abiotic stress. **J Exp Bot** 64:445-58.
259. Sulieman S, ***Tran LS** (2013). Asparagine: an amide of particular distinction in the regulation of symbiotic nitrogen fixation of legumes. **Crit Rev Biotechnol** 33:309-27.
260. Jogaiah S, Govind SR, ***Tran LS** (2013). Systems biology-based approaches towards understanding drought tolerance in food crops. **Crit Rev Biotechnol** 33:23-39.

2012

261. Ma Y, Qin F, ***Tran LS** (2012). Contribution of genomics to gene discovery in plant abiotic stress responses. **Mol Plant** 5:1176-8.
262. Choudhary PS, Yu J-Q, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2012). Benefits of brassinosteroid crosstalk. **Trends Plant Sci** 17:594-605.
263. Ha S, Vankova R, Yamaguchi-Shinozaki K, Shinozaki K, ***Tran LS** (2012). Cytokinins: metabolism and function in plant adaptation to environmental stresses. **Trends Plant Sci** 17:172-9.
264. Thao NP, ***Tran LS** (2012). Potentials toward genetic engineering of drought-tolerant soybean. **Crit Rev Biotechnol** 32:349-62.

2011

265. Choudhary PS, ***Tran LS** (2011). Phytosterols: Perspectives in human nutrition and clinical therapy. **Curr Med Chem** 18:4513-37.
266. Hadiarto T, ***Tran LS** (2011). Progress studies of drought-responsive genes in rice. **Plant Cell Rep** 30:297-310.

2010 and before

267. ***Tran LS**, Mochida K (2010). Functional genomics of soybean for improvement of productivity in adverse conditions. **Funct Integr Genomics** 10:447-62.
268. Le DT, Choi J-D, ***Tran LS** (2010). Amino acids conferring herbicide resistance in tobacco acetohydroxyacid synthase. **GM Crops** 1:62-7.
269. ***Tran LS**, Nishiyama R, Yamaguchi-Shinozaki K, Shinozaki K (2010). Potential utilization of NAC transcription factors to enhance abiotic stress tolerance in plants by biotechnological approach. **GM Crops** 1:34-41.
270. Manavalan LP, Guttikonda SK, ***Tran LS**, Nguyen HT (2009). Physiological and molecular approaches to improve drought resistance in soybean. **Plant Cell Physiol** 50:1260-76.

Addenda

271. ***Tran LS**, Mochida K (2010). A platform for functional prediction and comparative analyses of transcription factors of legumes and beyond. **Plant Signal Behav** 5:550-2. (Addendum to **Mochida K et al. Bioinformatics** 26: 90-1).
272. ***Tran LS**, Mochida K (2010). Identification and prediction of abiotic stress responsive transcription factors involved in abiotic stress signaling in soybean. **Plant Signal Behav** 5: 255-7. (Addendum to **Mochida K et al. DNA Res** 16:353-69).
273. ***Tran LS**, Shinozaki K, Yamaguchi-Shinozaki K (2010). Role of cytokinin responsive two-component system in ABA and osmotic stress signalings. **Plant Signal Behav** 5: 148-50. (Addendum to **Tran LS et al. Proc Natl Acad Sci USA** 104:20623-8).

Book chapters

274. Hoang XLT, Nguyen YNH, Thao NP, Le VA, ***Tran LS** (2020). NAC transcription factors in drought and salinity tolerance, in “Salt and Drought Stress Tolerance in Plants: Signaling Networks and Adaptive Mechanisms”, Eds. Hasanuzzaman M and Tanveer M, Springer International Publishing AG, Switzerland, pp. 351-66.
275. Chuong NN, Nghia DHT, Hoang XLT, Thi VAL, **Tran LS**, Thao NP (2020). Type 2C protein phosphatases in plant signaling pathways under abiotic stress, in “Protein Phosphatases and Stress Management in Plants - Functional Genomic Perspective”. Ed. Ed. Pandey GK, Springer International Publishing AG, Switzerland, pp.67-82.
276. Mochida K, **Tran LS** (2020). Genome information resources to improve plant biomass productivity, in “Climate Change, Photosynthesis and Advanced Biofuels”, Ed. Kumar A, Yau Y-Y, Ogita S and Scheibe R, Springer International Publishing AG, Switzerland, pp.323-38.
277. Bao A, ***Tran LS**, Cao D (2020). CRISPR/Cas9-based gene editing in soybean, in “Legume Genomics: Methods and Protocols”, **Methods in Molecular Biology** series, Eds. Garg R, Jain M. Springer Science+Business Media, Humana Press, New York, 2107:349-64.
278. Ong Q, Le VA, Thao NP, ***Tran LS** (2019). Transcriptional factor databases for legume plants, in “The Model Legume *Medicago truncatula*”, Ed. de Bruijn F.J., Wiley-Blackwell, John Wiley & Sons, Hoboken, NJ 07030-5774, USA, pp.1131-6.
279. Mochida K, ***Tran LS** (2017). Transcription factors in *Jatropha*, in “The Jatropha Genome (Compendium of Plant Genomes)”, Ed. Tsuchimoto S, Springer International Publishing AG, Switzerland, pp.47-60.
280. Li W, ***Tran LS** (2017). Effects of ethylene on seed germination of halophyte plants under salt stress, in “Ethylene Signaling”, **Methods in Molecular Biology** series, Eds. Binder B, Schaller GE. Springer Science+Business Media, Humana Press, New York, 1573: 253-9.
281. Thu NBA, Mai TT, Hoang XLT, Thao NP, Sulieman S, ***Tran LS** (2017). Cytokinin signaling in plant responses to abiotic stresses, in “Mechanism of Plant Hormone Signaling under Stress”, Ed. Pandey GK, Wiley-Blackwell, John Wiley & Sons, Hoboken, NJ 07030-5774, USA, pp.71-100.
282. Sulieman S, Thao NP, ***Tran LS** (2015). Does elevated CO₂ provide real benefits for N₂-fixing leguminous symbioses?, in “Legume Nitrogen Fixation in a Changing Environment – Achievements and Challenges”, Eds. Sulieman S and **Tran LS**. Springer International Publishing AG, Switzerland, pp. 89-112.
283. Mochida K, Ha CV, Sulieman S, Dong NV, ***Tran LS** (2015). Databases of transcription factors in legumes, in “Biological Nitrogen Fixation”, Ed. de Bruijn FJ. Wiley-Blackwell, John Wiley & Sons, Hoboken, NJ 07030-5774, USA, pp. 817-21.
284. Hoang XLT, Thu NBA, Thao NP, ***Tran LS** (2014). Transcription factors in abiotic stress responses -their potentials in crop improvement, in “Improvement of Crops in the Era of

- Climatic Change”, Vol. II, Eds. Ahmad P, Wani MR, Azooz MM and **Tran LS**. Springer-Verlag, New York, NY 10013, USA, pp. 337-66.
285. Xin H, Qin F, ***Tran LS** (2012). Transcription factors involved in environmental stress response in plants, in “Environmental Adaptations and Stress Tolerance of Plants in the Era of Climatic Change”, Eds. Ahmad P and Prasad MNV. Springer-Verlag, New York, NY 10013, USA, pp. 279-95.
 286. **Tran LS**, Nguyen HT (2009). Future biotechnology of legumes, in "Nitrogen Fixation in Crop Production", Eds. Emerich WD and Krishnan H. The American Society of Agronomy, Crop Science Society of America, Soil Science Society of America, Madison, WI, USA, pp. 265-308.
 287. **Tran LS**, Nakashima K, Shinozaki K, Yamaguchi-Shinozaki K (2007). Plant gene networks in osmotic stress response: from genes to regulatory networks, in “Osmosensing and Osmosignaling” **Methods in Enzymology** series, Eds. Häussinger D, Sies H. Elsevier, **428**:109-28.

Book editions

288. “Exogenous Priming and Engineering of Plant Metabolic and Regulatory Genes: Stress Mitigation Strategies in Plants”, 2024, Eds. Patel MK, **Tran LS**, Pandey S, Mishra A, Elsevier, Academic Press, Cambridge, MA, USA (under preparation).
289. “Signaling Molecules in Plant Development and Abiotic Stress Management”, 2023, Eds. Tripathi DK, Chauhan DK, **Tran LS**, Elsevier, Academic Press, Cambridge, MA, USA (under preparation).
290. “Salinity Responses and Tolerance in Plants, Volume 2 - Exploring RNAi, Genome Editing and Systems Biology”, 2018, Eds. Kumar V, Wani SH, Suprasanna P, **Tran LS**, Springer International Publishing AG, Switzerland.
291. “Salinity Responses and Tolerance in Plants, Volume 1 - Targeting Sensory, Transport and Signaling Mechanisms”, 2018, Eds. Kumar V, Wani SH, Suprasanna P, **Tran LS**, Springer International Publishing AG, Switzerland.
292. “Plant Micronutrient Use Efficiency - Molecular and Genomic Perspectives in Crop Plants”, 2018, Eds. Hossain MA, Kamiya T, Burritt DJ, **Tran LS**, Fujiwara T, Elsevier, Academic Press, Cambridge, MA, USA.
293. “Glutathione in plant growth, development and stress tolerance”, 2017, Eds. Hossain MA, Mostofa MG, Vivancos D, Fujita M, Burritt DJ, **Tran LS**, Springer International Publishing AG, Switzerland.
294. “Plant Macronutrient Use Efficiency - Molecular and Genomic Perspectives in Crop Plants”, 2017, Eds. Hossain MA, Kamiya T, Burritt DJ, **Tran LS**, Fujiwara T, Elsevier, Academic Press, Cambridge, MA, USA.
295. “Legume N₂ Fixation in Soils with Low Phosphorus Availability - Adaptation and Regulatory Implication”, 2017, Eds. Sulieman S, **Tran LS**, Springer International Publishing AG, Switzerland.
296. “Drought Stress Tolerance in Plants, Vol 2: Molecular and Genetic Perspectives”, 2016, Eds. Hossain MA, Wani SH, Bhattachajee S, Burritt DJ, **Tran LS**, Springer International Publishing AG, Switzerland.
297. “Drought Stress Tolerance in Plants, Vol 1: Physiology and Biochemistry”, 2016, Eds. Hossain MA, Wani SH, Bhattachajee S, Burritt DJ, **Tran LS**, Springer International Publishing AG, Switzerland.
298. “Legume Nitrogen Fixation in a Changing Environment”, 2015, Eds. Sulieman S, **Tran LS**, Springer International Publishing AG, Switzerland.

299. “Improvement of Crops in the Era of Climatic Change” (Vol. II), 2014, Eds. Ahmad P, Wani MR, Azooz MM, **Tran LS**, Springer-Verlag, New York, NY 10013, USA.
300. “Phytohormones: A Window to Metabolism, Signaling and Biotechnological Applications”, 2014, Eds. **Tran LS**, Choudhary PS, Springer-Verlag, New York, NY 10013, USA.
301. “Improvement of Crops in the Era of Climatic Change” (Vol. I), 2013, Eds. Ahmad P, Wani MR, Azooz MM, **Tran LS**, Springer-Verlag, New York, NY 10013, USA.

Special Issue editions for journals

302. “Alteration of Growth, Physiological, Biochemical and Molecular Responses of Phototropic Organisms to Environmental Stress”, 2022-2023, **Cells**, Eds. Ha CV, **Tran LS**, (https://www.mdpi.com/journal/cells/special_issues/12IGPSQ785)
303. “Redox Biology and Crop Health”, 2022, **Front Plant Sci**, Eds. Bhattacharjee S, **Tran LS**, (https://www.frontiersin.org/research-topics/40250/redox-biology-and-crop-health?utm_source=F-RTM&utm_medium=TED1&utm_campaign=PRD_TED1_T1_RT-TITLE)
304. “Plant Metabolites and Their Reprogramming for Plant Tolerance under Environmental Stress 2.0”, 2021, **Int J Mol Sci**, Eds. **Tran LS**, Patel MK, (https://www.mdpi.com/journal/ijms/special_issues/Metabolites_Reprogramming)
305. “Functions of Transcription Factors in Plant Growth, Performance and Responses to Environmental Stresses 2.0”, 2022, **Int J Mol Sci**, Eds. Thao NP, **Tran LS**, (https://www.mdpi.com/journal/ijms/special_issues/Transcription_Factors2)
306. “Plant growth regulators- signaling and crosstalk: revisiting their roles in plants under changing environment” (2022), **Plant Cell Physiol**, Eds. Yadav SR, Tripathi DK, Mochida K, **Tran LS**.
307. Topical Collection “Phytohormone Signaling and Crosstalk in Regulation of Plant Growth and Environmental Stress Responses” (ongoing), **Int J Mol Sci**, Eds. **Tran LS**, Mostofa MG (https://www.mdpi.com/journal/ijms/special_issues/Plant_Phytohormone)
308. “Functions of Transcription Factors in Plant Growth, Performance and Responses to Environmental Stresses 2.0”, 2022, **Int J Mol Sci**, Eds. Thao NP, **Tran LS** (https://www.mdpi.com/journal/ijms/special_issues/Transcription_Factors_Environmental)
309. “Physiological and Molecular Perspectives of Stress Tolerance in Vegetables”, 2021, **Front Plant Sci**, Eds. Shigyo M, Abdelrahman M, **Tran LS**, (<https://www.frontiersin.org/research-topics/18970/physiological-and-molecular-perspectives-of-stress-tolerance-in-vegetables>)
310. “Plant Metabolites and Their Reprogramming for Plant Tolerance under Environmental Stress”, 2021, **Int J Mol Sci**, Eds. **Tran LS**, Patel MK, (https://www.mdpi.com/journal/ijms/special_issues/Plant_Reprogramming)
311. “Functions of Transcription Factors in Plant Growth, Performance and Responses to Environmental Stresses”, 2021, **Int J Mol Sci**, Eds. **Tran LS**, Thao NP, (https://www.mdpi.com/journal/ijms/special_issues/Transcription_Factors_Environmental)
312. “Phytohormones and Their Crosstalk during Plant Growth, Development and Environmental Stress Adaptation: New Prospects and Challenges to Cope with Global Climate Change”, 2019, **Int J Mol Sci**, Eds. **Tran LS**, Mostofa MG (https://www.mdpi.com/journal/ijms/special_issues/phytohormones)
313. “Phytohormones and Their Crosstalk during Plant Growth, Development and Environmental Stress Adaptation”, 2018, **Int J Mol Sci**, Eds. **Tran LS**, Mostofa MG,

- Pollman S
(http://www.mdpi.com/journal/ijms/special_issues/phytohormones_plant_growth)
314. “Phytohormones and Their Crosstalk during Plant Growth, Development and Environmental Stress Adaptation”, 2018, **Int J Mol Sci**, Eds. **Tran LS**, Mostofa MG, Pollman S
(http://www.mdpi.com/journal/ijms/special_issues/phytohormones_plant_growth)
 315. “Enhancement of Plant Productivity in the Post-genomics Era”, 2016, **Curr Genomics**, Eds. **Tran LS**, Thao NP (<http://benthamscience.com/journals/current-genomics/volume/17/issue/4/>).
 316. “Plant Quality Improvement and Nutrigenomics”, 2016, **Curr Genomics**, Eds. **Tran LS**, Kumar R (<http://benthamscience.com/journals/current-genomics/volume/17/issue/3/>).
 317. “Regulation of Plant Mineral Nutrition: Transport, Sensing and Signalling”, 2015, **Int J Mol Sci**, Eds. Rouached H, **Tran LS**.
 318. “Plants Coping Abiotic and Biotic Stresses: A Tale of Diligent Management”, 2014, **Biomed Res Int**, Lead Guest Editor **Tran LS** and Guest Editors Pal S, Rachmilvetch S, Rouached H, Libault M.
 319. “Symbiotic Nitrogen Fixation in Legume Nodules: Metabolism and Regulatory Mechanism”, 2014, **Int J Mol Sci**, Eds. **Tran LS**, Sulieman S.