

# Kirill Schenstniy, Ph.D.

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## **Education**

- 2017 – 2021 **Doctor of Natural Sciences (*Dr. rer. nat.*) in Biology**  
The Center for Plant Molecular Biology (ZMBP), Eberhard Karls University of Tuebingen, Germany  
Dissertation: Analyses of TALE-induced resistance and putative susceptibility genes in tomato  
Principal Investigator: Prof. Dr. Thomas Lahaye ([link](#))
- 2014 – 2016 **Master of Science in Plant Breeding & Genetic Resources**  
Wageningen University & Research, the Netherlands
- 2010 – 2012 **Bachelor of Science in Agronomy**  
Dnipro State Agrarian and Economic University, Ukraine

## **Research Experience**

- 2025 – Present **Department of Chemical and Biomolecular Engineering, University of California, Berkeley**  
*Postdoctoral Researcher* – lab of Prof. Dr. Markita Landry ([link](#))
  - Development and optimization of gene-editing tools for transient plant transformation.
- 2024 – 2025 **Plant Development and Physiology Lab, Tokyo Metropolitan University, Japan**  
*Visiting Postdoctoral Researcher / Lecturer* – lab of Prof. Dr. Takashi Okamoto ([link](#))
  - Studying cell death execution during early zygote & fertilization-independent egg cell development;
  - DNA-free RNP-mediated gene editing in rice zygotes and non-fertilized egg cells;
  - *in vitro* fusion of germ cells from different plant species: rice × wild rice and rice × wheat.
- 2021 – 2024 **Institute for Molecular Physiology, Henrich Heine University of Duesseldorf, Germany**  
*Postdoctoral Researcher* – lab of Prof. Dr. Wolf Frommer ([link](#))
  - DNA-free RNP-mediated gene editing in rice zygotes and non-fertilized egg cells;
  - Nanoparticle-mediated delivery of gene-editing tools into plant cells;
  - Development of selectable markers for plant transformation.
- 2021 – 2021 **Department of General Genetics, ZMBP, Eberhard Karls University of Tuebingen, Germany**  
*Research Assistant* – lab of Prof. Dr. Suayib Uestuen ([link](#))
  - CRISPR/Cas9-mediated editing of autophagy-related genes in tomato.
- 2017 – 2020 **Department of General Genetics, ZMBP, Eberhard Karls University of Tuebingen, Germany**  
*Research Assistant* – lab of Prof. Dr. Thomas Lahaye ([link](#))
  - Dissecting genetic pathways mediated by NLR and executor (E) proteins in tomato;
  - Identification of putative susceptibility genes (TALE targets) in tomato;
  - CRISPR/Cas9-mediated editing of immunity-related genes in tomato.
- 2016 – 2016 **Plant Breeding, Wageningen University & Research, the Netherlands**  
*Research Fellow* – lab of Prof. Dr. Luisa Trindade ([link](#))
  - Genetic analysis and QTL mapping of cell wall components and digestibility in maize.
- 2015 – 2016 **Plant Breeding, Wageningen University & Research, the Netherlands**  
*Research Fellow* – lab of Prof. Dr. Yuling Bai ([link](#))
  - Identifying the role of potato "helper" NLR proteins in mediation of resistance to *P. infestans*.

## **Research Contributions**

1. Kumaran, G., Pathak, P.K., Quandoh, E., Mursalimov, S., Devi, J., Alkalai-Tuvia, S., Leong, J.X., **Schenstniy, K.**, Levin, E., Üstün, S., Michaeli, S. (2023). Autophagy restricts tomato fruit ripening via a general role in ethylene repression. *bioRxiv* [in revision]. <https://doi.org/10.1101/2023.12.20.572633>
2. **Schenstniy, K.**, Zhang, Z., Liu, B., Nakamura, M., Schepler-Luu, V., Loo, E.P.I., Yang, B., Frommer, W.B. (2023). Loss-of-function mutation in the polyamine transporter gene *OsLAT5* as a selectable marker for genome editing. *bioRxiv* [in revision]. <https://doi.org/10.1101/2023.12.12.571390>
3. Schepler-Luu, V., Sciallano, C., Stiebner, M., Ji, C., Boulard, G., Diallo, A., Auguy, F., Char, S. N., Arra, Y., **Schenstniy, K.**, Buchholzer, M., Loo, E.P.I., Bilaro, A.L., Lihepanyama, D., Mkuya, M., Murori, R., Oliva, R., Cunnac, S., Yang, B., Szurek, B., Frommer, W.B. (2023). Genome editing of an African elite rice variety confers resistance against endemic and emerging *Xanthomonas oryzae* pv. *oryzae* strains. *eLife*, 12, e84864. <https://doi.org/10.7554/eLife.84864>

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4. **Schenstniy, K.**, Strauß, A., Dressel, A., Morbitzer, R., Wunderlich, M., Andrade, A.G., Phan, T.-T.-T., Aguilera, P.d.l.A., Brancato, C., Berendzen, K. W., Lahaye, T. (2022). The tomato resistance gene *Bs4* suppresses leaf watersoaking phenotypes induced by AvrHah1, a transcription activator-like effector from tomato-pathogenic xanthomonads. *New Phytologist*, 236(5), 1856-1870. <https://doi.org/https://doi.org/10.1111/nph.18456>
5. Monino-Lopez, D., Nijenhuis, M., Kodde, L., Kamoun, S., Salehian, H., **Schenstniy, K.**, Stam, R., Lokossou, A., Abd-El-Haliem, A., Visser, R.G.F., Vossen, J. H. (2021). Allelic variants of the NLR protein Rpi-chc1 differentially recognize members of the *Phytophthora infestans* PexRD12/31 effector superfamily through the leucine-rich repeat domain. *The Plant Journal*, 107(1), 182-197. <https://doi.org/https://doi.org/10.1111/tpj.15284>

## **Honours, Awards, and Scholarships**

- 2023 [CEPLAS Seed Fund](#) – Funding for the development of independent and innovative projects (10.000 EUR / ≈11.000 USD).
- 2014 [Worldwide Studies Program](#) – Scholarship covering tuition fees at Wageningen University & Research (33.000 EUR / ≈36.000 USD)