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Date 2025/09/24

### Interests in Cheminformatics and Bioinformatics

#### Publications

#### Scientific Education

#### CV

## Interests in Cheminformatics and Bioinformatics

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- Software development of algorithmic solutions, scientific software packages and IT systems
- Molecular modeling and simulation, molecular recognition, QSAR/QSPR
- Model development, machine learning und computational intelligence
- Strategic developments

# Publications

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## Peer-Reviewed R&D Articles

- 2026 **Comparative Analysis of Chemical Structure String Representations for Neural Machine Translation**

K. Rajan, A. Zielesny, C. Steinbeck

In: Senn, W., et al. Artificial Neural Networks and Machine Learning. ICANN 2025 International Workshops and Special Sessions. ICANN 2025. Lecture Notes in Computer Science, vol 16072. pp. 8–16, 2026, Springer, Cham.

[\(online\)](#)

- 2025 **PySSA for Windows: End-User Protein Structure Prediction and Visual Analysis with ColabFold and PyMOL**

H. Kullik, M. Urban, J. Schaub, A. Loidl-Stahlhofen and A. Zielesny

Journal of Chemical Information and Modeling (2025), 65, 5839-5846

[\(online\)](#)

- 2024 **STOUT V2.0: SMILES to IUPAC name conversion using transformer models**

K. Rajan, A. Zielesny and C. Steinbeck

Journal of Cheminformatics (2024), 16:146

[\(online\)](#)

**An automated Calculation Pipeline for Differential Pair Interaction Energies with Molecular Force Fields using the Tinker Molecular Modeling Package**

F. Bänsch, M. Daniel, H. Lanig, C. Steinbeck, A. Zielesny

Journal of Cheminformatics (2024), 16:96

[\(online\)](#)

**Advancements in hand-drawn chemical structure recognition through an enhanced DECIMER architecture**

K. Rajan, H. O. Brinkhaus, A. Zielesny and C. Steinbeck

Journal of Cheminformatics (2024), 16:78

[\(online\)](#)

- 2023 **DECIMER.ai: an open platform for automated optical chemical structure identification, segmentation and recognition in scientific publications**

K. Rajan, H. O. Brinkhaus, M. I. Agea, A. Zielesny and C. Steinbeck

Nature Communications (2023), 14:5045

[\(online\)](#)

**Notes on molecular fragmentation and parameter settings for a dissipative particle dynamics study of a C<sub>10</sub>E<sub>4</sub>/water mixture with lamellar bilayer formation**

F. Bänsch, C. Steinbeck and A. Zielesny

Journal of Cheminformatics (2023), 15:23

[\(online\)](#)

- 2023 **Open data and algorithms for open science in AI-driven molecular informatics**  
H.-O. Brinkhaus, K. Rajan, J. Schaub, A. Zielesny, C. Steinbeck  
Current Opinion in Structural Biology (2023), 79:102542  
([online](#))
- MORTAR: a rich client application for in silico molecule fragmentation**  
F. Bänsch, J. Schaub, B. Sevindik, S. Behr, J. Zander, C. Steinbeck and A. Zielesny  
Journal of Cheminformatics (2023), 15:1  
([online](#))
- 2022 **Scaffold Generator: a Java library implementing molecular scaffold functionalities in the Chemistry Development Kit (CDK)**  
J. Schaub, J. Zander, A. Zielesny, C. Steinbeck  
Journal of Cheminformatics (2022), 14:79  
([online](#))
- Notes on the Treatment of Charged Particles for Studying Cyclotide/Membrane Interactions with Dissipative Particle Dynamics**  
F. Bänsch, C. Steinbeck and A. Zielesny  
Membranes (2022), 12(6):619  
([online](#))
- DECIMER - Hand-drawn molecule images dataset**  
H.-O. Brinkhaus, A. Zielesny, C. Steinbeck, K. Rajan  
Journal of Cheminformatics (2022), 14:36  
([online](#))  
Zenodo repository: [online](#)
- RanDepict - Random Chemical Structure Depiction Generator**  
H.-O. Brinkhaus, K. Rajan, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2022), 14:31  
([online](#))
- Performance of chemical structure string representations for chemical image recognition using transformers**  
K. Rajan, C. Steinbeck and A. Zielesny  
Digital Discovery (2022), 1:84-90  
([online](#))
- 
- 2021 **DECIMER 1.0: deep learning for chemical image recognition using transformers**  
K. Rajan, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2021), 13:61  
([online](#))
- Quantitative Estimation of Cyclotide-Induced Bilayer Membrane Disruption by Lipid Extraction with Mesoscopic Simulation**  
K. van den Broek, M. Epple, L. S. Kersten, H. Kuhn and A. Zielesny  
Journal of Chemical Information and Modeling (2021), 61, 3027-3040  
([online](#))

- 2021 **STOUT: SMILES to IUPAC names using neural machine translation**  
K. Rajan, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2021), 13:34  
([online](#))
- Description and Analysis of Glycosidic Residues in the Largest Open Natural Products Database**  
J. Schaub, A. Zielesny, C. Steinbeck and M. Sorokina  
Biomolecules (2021), 11, 486  
([online](#))
- DECIMER-Segmentation: Automated extraction of chemical structure depictions from scientific literature**  
K. Rajan, H. O. Brinkhaus, M. Sorokina, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2021), 13:20  
([online](#))
- Molecule Set Comparator (MSC) – A CDK-based open rich-client tool for molecule set similarity evaluations**  
K. Rajan, J.-M. Hein, C. Steinbeck and A. Zielesny  
Journal of Cheminformatics (2021), 13:5  
([online](#))
- 2020 **Too sweet: cheminformatics for deglycosylation in natural products**  
J. Schaub, A. Zielesny, C. Steinbeck and M. Sorokina  
Journal of Cheminformatics (2020), 12:67  
([online](#))
- DECIMER - Towards Deep Learning for Chemical Image Recognition**  
K. Rajan, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2020), 12:65  
([online](#))
- A review of optical chemical structure recognition tools**  
K. Rajan, H. O. Brinkhaus, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2020), 12:60  
([online](#))
- MFsim - an open Java all-in-one rich-client simulation environment for mesoscopic simulation**  
K. van den Broek, M. Daniel, M. Epple, J.-M. Hein, H. Kuhn, S. Neumann, A. Truszkowski and A. Zielesny  
Journal of Cheminformatics (2020), 12:29  
([online](#))
- 2019 **ErtlFunctionalGroupsFinder: automated rule-based functional group detection with the Chemistry Development Kit (CDK)**  
S. Fritsch, S. Neumann, J. Schaub, C. Steinbeck and A. Zielesny  
Journal of Cheminformatics (2019), 11:37  
([online](#))

- 2018 **SPICES: a particle-based molecular structure line notation and support library for mesoscopic simulation**  
K. van den Broek, M. Daniel, M. Epple, H. Kuhn, J. Schaub and A. Zielesny  
Journal of Cheminformatics (2018), 10:35  
([online](#))
- Jdpd: an open Java simulation kernel for molecular fragment dissipative particle dynamics**  
K. van den Broek, H. Kuhn and A. Zielesny  
Journal of Cheminformatics (2018), 10:25  
([online](#))
- 2015 **Mesoscopic Simulation of Phospholipid Membranes, Peptides, and Proteins with Molecular Fragment Dynamics**  
A. Truszkowski, K. van den Broek, H. Kuhn, A. Zielesny and M. Epple  
Journal of Chemical Information and Modeling (2015), 55(5), 983-997  
([online](#))
- 2014 **A molecular fragment cheminformatics roadmap for mesoscopic simulation**  
A. Truszkowski, M. Daniel, H. Kuhn, S. Neumann, C. Steinbeck, A. Zielesny and M. Epple  
Journal of Cheminformatics (2014), 6:45  
([online](#))
- 2013 **Geometric, electronic and NMR properties of Hemicucurbit[n]urils and their anionic Complexes**  
H.-J. Buschmann and A. Zielesny  
Computational and Theoretical Chemistry (2013), 1022, 14-22  
([online](#))
- Molecular fragment dynamics study on the water-air interface behavior of non-ionic polyoxyethylene alkyl ether surfactants**  
A. Truszkowski, M. Epple, A. Fiethen, A. Zielesny and H. Kuhn  
Journal of Colloid and Interface Science (2013), 410, 140-145  
([online](#))
- 2011 **New developments on the cheminformatics open workflow environment CDK-Taverna**  
A. Truszkowski, K. V. Jayaseelan, S. Neumann, E. L. Willighagen, A. Zielesny and C. Steinbeck  
Journal of Cheminformatics (2011), 3, 54  
([online](#))
- 2010 **CDK-Taverna: an open workflow environment for cheminformatics**  
T. Kuhn, E. L. Willighagen, A. Zielesny and C. Steinbeck  
BMC Bioinformatics (2010), 11, 159  
([online](#))
- 2006 **Structure, Stability, Electronic Properties and NMR-Shielding of the Cucurbit[6]uril-Spermine-Complex**  
H.-J. Buschmann, A. Wego, A. Zielesny, E. Schollmeyer  
Journal of Inclusion Phenomena and Macrocyclic Chemistry (2006), 54, 241-246

- 2006 **Hemicucurbit[6]uril: A macrocyclic ligand with unusual complexing properties**  
 H.-J. Buschmann, A. Zielesny, E. Schollmeyer  
 Journal of Inclusion Phenomena and Macrocyclic Chemistry (2006), 54, 181-185
- Structure, Electronic Properties and NMR-Shielding of Cucurbit[n]urils**  
 H.-J. Buschmann, A. Wego, A. Zielesny, E. Schollmeyer  
 Journal of Inclusion Phenomena and Macrocyclic Chemistry (2006), 54, 85-88
- 2004 **Neuronal networks curtail adhesive development** (in German)  
 K. U. Koch, A. Zielesny  
 Adhaesion (2004), 48(1-2), 32-37
- 1995 **Ultrasonic absorption of nitrobenzene-isooctane mixtures of non-critical composition**  
 A. Zielesny, D. Woermann  
 J. Chem. Soc., Faraday Trans. (1995), 91 (21), 3889-3892
- Diffusivity of a non-ionic surfactant/water mixture of critical composition**  
 M. Lesemann, A. Zielesny, L. Belkoura, D. Woermann  
 J. Chem. Phys. (1995), 102 (1), 414-418
- 1994 **Crossover behaviour and critical amplitude of the viscosity of binary liquid mixtures of critical composition**  
 A. Zielesny, D. Woermann  
 J. Chem. Soc., Faraday Trans. (1994), 90 (15), 2215-2222
- Static light scattering experiments with non-ionic surfactant/water mixtures of critical composition**  
 A. Zielesny, L. Belkoura, D. Woermann  
 Ber. Bunsen-Ges. Phys. Chem. (1994), 98 (4), 579-584
- Viscosity of two non-ionic amphiphile-water mixtures of critical composition: Study of the systems triethyleneglycolmonohexyl ether (C<sub>6</sub>E<sub>3</sub>)-water and tetraethyleneglycolmono-octyl ether (C<sub>8</sub>E<sub>4</sub>)-water**  
 A. Zielesny, S. Limberg, D. Woermann  
 Ber. Bunsen-Ges. Phys. Chem. (1994), 98 (2), 195-201
- Viscosity and diffusivity of a binary liquid mixture of critical composition: study of the system 2-butoxyethanol/water**  
 A. Zielesny, J. Schmitz, S. Limberg, A. G. Aizpiri, S. Fusenig, D. Woermann  
 Int. J. Thermophys. (1994), 15 (1), 67-94
- 1990 **Temperature dependence of viscosity of polystyrene/cyclohexane mixtures of critical composition**  
 W. A. Goedel, A. Zielesny, L. Belkoura, T. Engels, D. Woermann  
 Ber. Bunsen-Ges. Phys. Chem. (1990), 94 (1), 17-24

## Patent

- 2005 **Method for the Provision of any Type of Storage Media containing pre-recorded structured Information**  
B. Achten, F. Augustin, O. Gebert, S. Kohl, C. Schaub, M. Schimeczek, A. Zielesny  
Bayer Business Service GmbH  
Patent WO2005086039A2 (2005-09-15), US2008021798A1, JP2007528543A  
Application EP2005001748W (2005-02-19)  
Priority DE102004010517A (2004-03-04), DE102004032231A (2004-07-02)

## Textbook

- 2016 **From Curve Fitting to Machine Learning: An illustrative Guide to scientific Data Analysis and Computational Intelligence**  
***Second revised and extended Edition***  
A. Zielesny  
[Springer: Intelligent Systems Reference Library, Volume 109, 2016](#)  
Supplementary information available on [GitHub](#).
- 2011 **From Curve Fitting to Machine Learning: An illustrative Guide to scientific Data Analysis and Computational Intelligence**  
A. Zielesny  
[Springer: Intelligent Systems Reference Library, Volume 18, 2011](#)  
Supplementary information available on [GitHub](#).

## Preprint R&D Articles (not peer-reviewed)

- 2025 **MARCUS: Molecular Annotation and Recognition for Curating Unravalled Structures**  
K. Rajan, V. Weißenborn, L. Lederer, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- 2024 **STOUT V2.0: SMILES to IUPAC name conversion using transformer models**  
K. Rajan, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- PySSA: end-user protein structure prediction and visual analysis with ColabFold and PyMOL**  
H. Kullik, M. Urban, J. Schaub, A. Loidl-Stahlhofen, A. Zielesny  
ChemRxiv. Preprint: [online](#)
- Advancements in Hand-Drawn Chemical Structure Recognition through an Enhanced DECIMER Architecture**  
K. Rajan, H. O. Brinkhaus, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- An automated Calculation Pipeline for Differential Pair Interaction Energies with Molecular Force Fields using the Tinker Molecular Modeling Package**  
F. Bänsch, M. Daniel, H. Lanig, C. Steinbeck, A. Zielesny  
ChemRxiv. Preprint: [online](#)
- 2023 **Complex formation of Cucurbit[6]uril with DABCO and DABCO derivatives: An experimental and DFT study**  
A. Wego, A. Zielesny  
ChemRxiv. Preprint: [online](#)
- DECIMER.ai - An open platform for automated optical chemical structure identification, segmentation and recognition in scientific publications**  
K. Rajan, H. O. Brinkhaus, M. I. Agea, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- 2022 **Open data and algorithms for open science in AI-driven molecular informatics**  
H. O. Brinkhaus, K. Rajan, J. Schaub, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- MORTAR - A Rich Client Application for in silico Molecule Fragmentation**  
F. Bänsch, J. Schaub, B. Sevindik, S. Behr, J. Zander, C. Steinbeck, A. Zielesny  
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- DECIMER - Hand-drawn molecule images dataset**  
H.-O. Brinkhaus, A. Zielesny, C. Steinbeck, K. Rajan  
ChemRxiv. Preprint: [online](#)  
Zenodo repository: [online](#)

- 2022 **Scaffold Generator - A Java library implementing molecular scaffold functionalities in the Chemistry Development Kit (CDK)**  
J. Schaub, J. Zander, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- RanDepict - Random Chemical Structure Depiction Generator**  
H.-O. Brinkhaus, K. Rajan, A. Zielesny, C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- 2021 **Performance of chemical structure string representations for chemical image recognition using transformers**  
K. Rajan, C. Steinbeck, A. Zielesny  
ChemRxiv. Preprint: [online](#)
- DECIMER 1.0: Deep Learning for Chemical Image Recognition using Transformers**  
K. Rajan, A. Zielesny and C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- Quantitative estimation of cyclotide-induced bilayer membrane disruption by lipid extraction with mesoscopic simulation**  
K. van den Broek, M. Epple, L. S. Kersten, H. Kuhn and A. Zielesny  
ChemRxiv. Preprint: [online](#)
- DECIMER Segmentation - Automated Extraction of Chemical Structure Depictions from Scientific Literature**  
K. Rajan, H. O. Brinkhaus, M. Sorokina, A. Zielesny and C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- 2020 **STOUT: SMILES to IUPAC names using Neural Machine Translation**  
K. Rajan, A. Zielesny and C. Steinbeck  
ChemRxiv. Preprint: [online](#)
- Too sweet: cheminformatics for deglycosylation in natural products**  
J. Schaub, A. Zielesny, C. Steinbeck and M. Sorokina  
Preprint: [online](#)
- DECIMER - Towards Deep Learning for Chemical Image Recognition**  
K. Rajan, A. Zielesny and C. Steinbeck  
ChemRxiv. Preprint: [online](#)

## Open-Source Software

- 2024 **MFsim (Versions 2.6.4.0, 2.6.3.0, 2.6.2.0)**  
*An open Java all-in-one rich-client simulation environment for mesoscopic simulation*  
Available on [GitHub](#).
- Jdpd (Versions 1.6.2.0)**  
*An open Java Simulation Kernel for Molecular Fragment Dissipative Particle Dynamics (DPD)*  
Available on [GitHub](#).
- 2023 **MFsim (Versions 2.6.1.0, 2.6.0.0)**  
*An open Java all-in-one rich-client simulation environment for mesoscopic simulation*  
Available on [GitHub](#).
- Jdpd (Versions 1.6.1.0)**  
*An open Java Simulation Kernel for Molecular Fragment Dissipative Particle Dynamics (DPD)*  
Available on [GitHub](#).
- 2022 **MFsim (Versions 2.5.0.0)**  
*An open Java all-in-one rich-client simulation environment for mesoscopic simulation*  
Available on [GitHub](#).
- Jdpd (Versions 1.6.0.0)**  
*An open Java Simulation Kernel for Molecular Fragment Dissipative Particle Dynamics (DPD)*  
Available on [GitHub](#).
- 2021 **MFsim (Versions 2.4.0.0, 2.3.0.0)**  
*An open Java all-in-one rich-client simulation environment for mesoscopic simulation*  
Available on [GitHub](#).
- Jdpd (Versions 1.5.0.0, 1.4.0.0)**  
*An open Java Simulation Kernel for Molecular Fragment Dissipative Particle Dynamics (DPD)*  
Available on [GitHub](#).
- 2020 **MSC (Version 1.0)**  
*Molecule Set Comparator*  
Available on [GitHub](#).

- 2020 **MFsim (Version 2.2.4.0)**  
*An open Java all-in-one rich-client simulation environment for mesoscopic simulation*  
Available on [GitHub](#).
- CIP - Computational Intelligence Packages (Version 3.1)**  
*Mathematica® 11 (or higher) open-source library for curve fitting, data smoothing, clustering and machine learning*  
Available on [GitHub](#).
- 2019 **ErtlFunctionalGroupsFinder (Versions 1.0.3.0, 1.0.2.0, 1.0.1.0, 1.0.0.0)**  
*Ertl algorithm for automated functional groups detection and extraction of organic molecules implemented on the basis of the Chemistry Development Kit (CDK)*  
Available on [GitHub](#).
- Jdpd (Versions 1.3.0.0, 1.2.0.0, 1.1.1.0)**  
*An open Java Simulation Kernel for Molecular Fragment Dissipative Particle Dynamics (DPD)*  
Available on [GitHub](#).
- 2018 **SPICES (Version 1.0.0.0)**  
*A particle-based Molecular Structure Line Notation and Support Library for Mesoscopic Simulation*  
Available on [GitHub](#).
- Jdpd (Versions 1.1.0.0, 1.0.2.0, 1.0.1.0 and 1.0.0.0)**  
*An open Java Simulation Kernel for Molecular Fragment Dissipative Particle Dynamics (DPD)*  
Available on [GitHub](#).
- CIP - Computational Intelligence Packages (Version 3.0)**  
*Mathematica® 11 (or higher) open-source library for curve fitting, data smoothing, clustering and machine learning*  
Available on [GitHub](#).
- MathCompiler (Version 1.0.0.0)**  
*A C# high-performance mathematical function compiler for efficient and fast calculation of function values of arbitrary complex single-line mathematical formulas at design and runtime.*  
Available on [GitHub](#).
- 2016 **CIP - Computational Intelligence Packages (Version 2.0)**  
*Mathematica® 10 (or higher) open-source library for curve fitting, data smoothing, clustering and machine learning*  
Available on [GitHub](#).
- 2012 **CIP - Computational Intelligence Packages (Version 1.1 and 1.2)**  
*Mathematica® 7 (or higher) open-source library for curve fitting, data smoothing, clustering and machine learning*  
Available on [GitHub](#).

- 2012 **CDK-Taverna (Version 2.0)**  
*Java open-source pipelining/workflow solution for cheminformatics through combination of different open-source projects (Taverna, CDK - Chemistry Development Kit, WEKA - Waikato Environment for Knowledge Analysis)*  
Websites: [Blog](#)
- 2011 **CIP - Computational Intelligence Packages (Version 1.0)**  
*Mathematica® 7 (or higher) open-source library for curve fitting, data smoothing, clustering and machine learning*  
Available on [GitHub](#).
- 2010 **MathComponent Freeware Library (Version 1.0.0.1)**  
*C# 2.0 free class collection for a fast, flexible and automated mathematical interplay of an application's classes and components at design and run time*  
Project support was stopped in 2018, see successor project "MathCompiler" above.

## Online Lectures

- 2011 **Chemical File Formats and Line Notations**  
A. Zielesny  
in C. Steinbeck (ed.), Introduction to Cheminformatics: The Biomedical & Life Sciences Collection, Henry Stewart Talks Ltd, London ([online](#)) , 2011 (last reviewed: 2022)
- Storing, searching and dissemination of chemical information**  
A. Zielesny  
in C. Steinbeck (ed.), Introduction to Cheminformatics: The Biomedical & Life Sciences Collection, Henry Stewart Talks Ltd, London ([online](#)) , 2011 (last reviewed: 2022)

## Other Articles

- 2025 **DECIMER.ai – Recognition of hand-drawn chemical molecules with Artificial Intelligence** (in German)  
K. Rajan, C. Steinbeck, A. Zielesny  
Research and Development at the Westphalian University  
Research Report 2024
- PySSA – Prediction and Analysis of Protein Structures with Artificial Intelligence** (in German)  
H. Kullik, M. Urban, J. Schaub, A. Loidl-Stahlhofen, A. Zielesny  
Research and Development at the Westphalian University  
Research Report 2024
- 2023 **Decomposition of molecules into biofunctional units** (in German)  
F. Bänsch, S. Behr, J. Schaub, B. Sevindik, C. Steinbeck, A. Zielesny  
Research and Development at the Westphalian University  
Research Report 2022
- Molecular interactions for biomolecular mesoscopic simulations** (in German)  
F. Bänsch, M. Daniel, C. Steinbeck, A. Zielesny  
Research and Development at the Westphalian University  
Research Report 2022
- 2022 **DECIMER – Artificial intelligence for optical chemical structure recognition**  
K. Rajan, C. Steinbeck, A. Zielesny  
Research and Development at the Westphalian University  
Research Report 2018 – 2021
- Mesoscopic simulation of large (bio)molecular systems**  
K. van den Broek, A. Zielesny  
Research and Development at the Westphalian University  
Research Report 2018 – 2021

- 2021 **DECIMER – Artificial intelligence for recognising the structures of chemical molecules** (in German)  
K. Rajan, C. Steinbeck, A. Zielesny  
Research and Development at the Westphalian University of Applied Sciences  
Research Report 2020
- 2019 **Mesoscopic Simulation of large (bio)molecular Systems** (in German)  
K. van den Broek, A. Zielesny  
Research and Development at the Westphalian University of Applied Sciences  
Research Report 2018
- 2015 **A virtual Microscope into the biomolecular Nanoworld** (in German)  
K. van den Broek, A. Truszkowski, A. Zielesny  
Research and Development at the Westphalian University of Applied Sciences  
Research Report 2014
- 2010 **Open-Source-Workflows** (in German)  
T. Kuhn, C. Steinbeck, A. Zielesny  
Nachrichten aus der Chemie (2010), 58 (1), 40-42
- 2006 **From Industrie to University** (in German)  
A. Zielesny  
Nachrichten aus der Chemie (2006), 54 (11), 1172-1173  
Gesellschaft Deutscher Chemiker, Berufsbilder in der Chemie, March 2008, page 44
- 2005 **Chemistry Software Package ChemOffice Ultra 2005**  
A. Zielesny  
J. Chem. Inf. Model. (2005), 45 (5), 1474-1477
- Platform for Scientific Information** (in German)  
C. Schaub, A. Zielesny  
Nachrichten aus der Chemie (2005), 53 (7/8), 786-788

## Conference Lectures and Poster Contributions

2025 **MARCUS - Molecular Annotation and Recognition for Curating Unravalled Structures**

K. Rajan, V. Weißenborn, L. Lederer, A. Zielesny and C. Steinbeck

Poster contribution to the „ 8th RSC-CICAG / RSC-BMCS Artificial Intelligence in Chemistry” from September 22 to 24, 2025, in Cambridge, United Kingdom

**The Molecule Fragmentation Framework (MORTAR): a rich client application for algorithmic substructure extraction**

F. Bänsch, C. Steinbeck, A. Zielesny, J. Schaub

Poster contribution to the „13th International Conference on Chemical Structures” from June 1 to 5, 2025, in Noordwijkerhout, The Netherlands

**PySSA: end-user protein structure prediction and visual analysis with ColabFold and PyMOL**

H. Kullik, M. Urban, J. Schaub, A. Loidl-Stahlhofen, A. Zielesny

Poster contribution to the „13th International Conference on Chemical Structures” from June 1 to 5, 2025, in Noordwijkerhout, The Netherlands

2024 **COCONUT 2.0 and automated literature mining using DECIMER.ai**

K. Rajan, V. Chandrasekhar, S. Kanakam, N. Sharma, J. Schaub, A. Zielesny, C. Steinbeck

Poster contribution to the „18th German Conference on Cheminformatics” from November 3 to 6, 2024, in Bad Soden am Taunus, Germany

**STOUT V2: Improved Translation Between SMILES/SELFIES and IUPAC Nomenclature Using Transformer Models**

K. Rajan, A. Zielesny, C. Steinbeck

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Poster contribution to the „18th German Conference on Cheminformatics” from November 3 to 6, 2024, in Bad Soden am Taunus, Germany

**MORTAR: a rich client application for in silico molecule fragmentation and fragment feature vector-based clustering**

J. Schaub, F. Bänsch, B. Sevindik, S. Behr, M. Rottmann, Z. Dagtekin, C. Steinbeck, A. Zielesny

Poster contribution to the „18th German Conference on Cheminformatics” from November 3 to 6, 2024, in Bad Soden am Taunus, Germany

**An automated Calculation Pipeline for Differential Pair Interaction Energies with Molecular Force Fields using the Tinker Molecular Modeling Package**

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2024 **PySSA: end-user protein structure prediction and visual analysis with ColabFold and PyMOL**

A. Zielesny, H. Kullik, M. Urban, J. Schaub, A. Loidl-Stahlhofen

Poster contribution to the „18th German Conference on Cheminformatics” from November 3 to 6, 2024, in Bad Soden am Taunus, Germany

**STOUT V2: Improved Translation Between SMILES/SELFIES and IUPAC Nomenclature Using Transformer Models**

K. Rajan, A. Zielesny, C. Steinbeck

Poster contribution to the „ACS Fall Meeting” from August 18 to 22, 2024, in Denver, USA

**Advancements in Hand-Drawn Chemical Structure Recognition through an Enhanced DECIMER Architecture**

K. Rajan, A. Zielesny, C. Steinbeck

Poster contribution to the „ACS Fall Meeting” from August 18 to 22, 2024, in Denver, USA

**The DECIMER.ai Project**

K. Rajan, A. Zielesny and C. Steinbeck

Poster contribution to the „Workshop Computer Vision for Science (CV4Science)” at the „IEEE / CVF Computer Vision and Pattern Recognition Conference (CVPR)”, June 17, 2024, Seattle, USA

**Also in Chemistry, Deep Learning Models Love Really Big Data**

K. Rajan, A. Zielesny, C. Steinbeck

Lecture at the „Beilstein Bozen Symposium 2024 - AI in Chemistry and Biology: Evolution or Revolution?” from June 4 to 6, 2024, in Rüdesheim, Germany

**An automated Calculation Pipeline for Differential Pair Interaction Energies with Molecular Force Fields using the Tinker Molecular Modeling Package**

F. Bänsch, M. Daniel, H. Lanig, C. Steinbeck, A. Zielesny

Poster contribution to the „36th Molecular Modeling Workshop” from March 4 to 6, 2024, in Erlangen, Germany

2023 **DECIMER.ai developments and automated chemical literature mining for COCONUT 2.0**

K. Rajan, H.-O. Brinkhaus, M. I. Agea, V. C. Nainala, N. Sharma, A. Zielesny and C. Steinbeck

Lecture at the „ACS Fall Meeting” from August 13 to 17, 2023, in San Francisco, USA

**Training data diversification for data-driven OCSR applications with RanDepict**

H.-O. Brinkhaus, K. Rajan, M. I. Agea, A. Zielesny, C. Steinbeck

Poster contribution to the „ACS Fall Meeting” from August 13 to 17, 2023, in San Francisco, USA

**MORTAR – A Rich Client Application for in silico Molecule Fragmentation**

F. Bänsch, J. Schaub, B. Sevindik, S. Behr, J. Zander, C. Steinbeck, A. Zielesny

Poster contribution to the „35th Molecular Modeling Workshop” from March 13 to 15, 2023, in Erlangen, Germany

- 2023 **A Calculation Pipeline for Differential Molecule Pair Interaction Energies**  
F. Bänsch, M. Daniel, H. Lanig, C. Steinbeck, A. Zielesny  
Poster contribution to the „35th Molecular Modeling Workshop” from March 13 to 15, 2023, in Erlangen, Germany
- 2022 **The DECIMER (Deep IEarning for Chemical IMagE Recognition) project**  
K. Rajan, H.-O. Brinkhaus, A. Zielesny, C. Steinbeck  
Lecture at the „ ACS Fall Meeting” from August 21 to 25, 2022, in Chicago, USA
- The DECIMER (Deep IEarning for Chemical IMagE Recognition) project**  
K. Rajan, H.-O. Brinkhaus, A. Zielesny, C. Steinbeck  
Poster contribution to the „12th International Conference on Chemical Structures” from June 12 to 16, 2022, in Noordwijkerhout, The Netherlands
- The DECIMER (Deep IEarning for Chemical IMagE Recognition) project**  
K. Rajan, H.-O. Brinkhaus, A. Zielesny, C. Steinbeck  
Lecture at the „17th German Conference on Cheminformatics” from May 8 to 10, 2022, in Garmisch-Partenkirchen, Germany
- In silico deglycosylation with the Sugar Removal Utility (SRU)**  
J. Schaub, A. Zielesny, C. Steinbeck, M. Sorokina  
Poster contribution to the „17th German Conference on Cheminformatics” from May 8 to 10, 2022, in Garmisch-Partenkirchen, Germany
- MORTAR - An open rich-client framework for in silico molecule fragmentation**  
F. Bänsch, J. Schaub, C. Steinbeck, A. Zielesny  
Poster contribution to the „17th German Conference on Cheminformatics” from May 8 to 10, 2022, in Garmisch-Partenkirchen, Germany
- 2020 **A new Approach to DPD Repulsion Parameter Estimation**  
F. Bänsch, M. Daniel, H. Lanig, C. Steinbeck and A. Zielesny  
Poster contribution to the „34th Molecular Modeling Workshop” from February 17 to 19, 2020, in Erlangen, Germany
- Rule-based in-silico Fragmentation for the Analysis of Natural Product Chemical Space**  
J. Schaub, F. Bänsch, A. Zielesny and C. Steinbeck  
Poster contribution to the „34th Molecular Modeling Workshop” from February 17 to 19, 2020, in Erlangen, Germany
- 2019 **Towards a comprehensive open computational support cycle for Molecular Fragment Dissipative Particle Dynamics (DPD)**  
F. Bänsch, M. Daniel, J. Schaub, C. Steinbeck, A. Zielesny  
Poster contribution to the „15th German Conference on Cheminformatics” from November 3 to 5, 2019, in Mainz, Germany
- ErtlFunctionalGroupsFinder: automated rule-based functional group detection with the Chemistry Development Kit (CDK)**  
J. Schaub, S. Fritsch, S. Neumann, C. Steinbeck, A. Zielesny  
Poster contribution to the „15th German Conference on Cheminformatics” from November 3 to 5, 2019, in Mainz, Germany

- 2018 **Steps towards an open all-in-one rich-client Environment for particle-based Mesoscopic Simulation**  
 K. van den Broek, H. Kuhn, A. Zielesny and M. Epple  
 Poster contribution to the „14th German Conference on Cheminformatics” from November 11 to 13, 2018, in Mainz, Germany  
**PSMILES – A particle-based Molecular Structure Representation for Mesoscopic Simulation**  
 K. van den Broek, M. Daniel, M. Epple, J. Schaub, H. Kuhn and A. Zielesny  
 Poster contribution to the „11th International Conference on Chemical Structures” from May 27 to 31, 2018, in Noordwijkerhout, The Netherlands
- 2017 **Mesoscopic Simulations: New Membrane Models & Studying the Mechanism of the Cyclotide Kalata B1 and it’s Mutants**  
 K. van den Broek, H. Kuhn, A. Zielesny and M. Epple  
 Poster contribution to the „13th German Conference on Cheminformatics” from November 5 to 7, 2017, in Mainz, Germany
- Improved Plasma Membrane Models as Test Systems for the Membrane Disrupting Activity of Kalata B1**  
 K. van den Broek, S. Albrecht, H. Kuhn, A. Zielesny and M. Epple  
 Poster contribution to the „31th Molecular Modeling Workshop” from March 27 to 29, 2017, in Erlangen, Germany
- 2016 **Mesoscopic simulation of the membrane disrupting activity of the cyclotide Kalata B1**  
 K. van den Broek, H. Kuhn, A. Zielesny and M. Epple  
 Poster contribution to the „12th German Conference on Cheminformatics” from November 6 to 8, 2016, in Fulda, Germany
- Mesoscopic Simulation of the Membrane Disrupting Activity of the Cyclotide Kalata B1**  
 K. van den Broek, H. Kuhn, A. Zielesny and M. Epple  
 Poster contribution to the „30th Molecular Modeling Workshop” from April 4 to 6, 2016, in Erlangen, Germany
- 2015 **Mesoscopic Simulation of Biomolecular Systems**  
 A. Zielesny, K. van den Broek, M. Epple, H. Kuhn, A. Truszkowski  
 Lecture at the „11th German Conference on Cheminformatics” from November 8 to 10, 2015, in Fulda, Germany  
 Journal of Cheminformatics 2016, 8(Suppl 1):O8
- 2014 **Molecular simulations of peptides and proteins with Molecular Fragment Dynamics**  
 A. Truszkowski, A. Fiethen, H. Kuhn, A. Zielesny and M. Epple  
 Poster contribution to the „10th International Conference on Chemical Structures” and „10th German Conference on Cheminformatics” from June 1 to 5, 2014, in Noordwijkerhout, The Netherlands

2014 **Molecular Fragment Dynamics Study of the Interaction between Zinc Ricinoleate and the Complexing Agent Methylglycinediacetic Acid as a new System for Enzyme Purification**

K. van den Broek, A. Fiethen, A. Truszkowski, A. Zielesny and H. Kuhn

Poster contribution to the „10th International Conference on Chemical Structures” and „10th German Conference on Chemoinformatics” from June 1 to 5, 2014, in Noordwijkerhout, The Netherlands

**Extension of molecular fragment based mesoscopic simulation to the biopolymer realm**

A. Truszkowski, A. Fiethen, H. Kuhn, A. Zielesny and M. Epple

Poster contribution to the „28th Molecular Modeling Workshop” from March 17 to 19, 2014, in Erlangen, Germany

**Molecular fragment dynamics study on the water-air interface behavior of non-ionic polyoxyethylene alkyl ether surfactants**

A. Truszkowski, A. Fiethen, H. Kuhn, T. Wiebringhaus, A. Zielesny and M. Epple

Poster contribution to the „9th German Conference on Chemoinformatics” from November 10 to 12, 2013, in Fulda, Germany

Journal of Cheminformatics 2014, 6(Suppl 1):P9

[\(online\)](#)

2013 **A molecular fragment dynamics (MFD) study on the phase behavior of non-ionic surfactants**

A. Truszkowski, A. Fiethen, H. Kuhn, A. Zielesny and M. Epple

Poster contribution to the „9th Zsigmondy Colloquium” from March 6 to 8, 2013, in Essen, Germany

2012 **Molecular simulations of peptides and proteins with Molecular Fragment Dynamics (MFD)**

A. Truszkowski, A. Fiethen, H. Kuhn, A. Zielesny and M. Epple

Poster contribution to the „8th German Conference on Chemoinformatics” from November 11 to 13, 2012, in Goslar, Germany

Journal of Cheminformatics (2013), 5(Suppl 1): P4

[\(online\)](#)

**Adsorption of amino acids in MFI-type zeolite: a computational and experimental study**

K. Stueckenschneider, J. Merz, A. Zielesny, G. Schembecker

Poster contribution to the „15. European Congress on Biotechnology” from September 23 to 26, 2012, in Istanbul, Turkey

**Adsorption of Alanine and Phenylalanine on MFI-type Zeolite: DFT Calculations and Experimental Results**

K. Stueckenschneider, A. Zielesny, G. Schembecker

Poster contribution to the „26th Molecular Modeling Workshop” from March 12 to 14, 2012, in Erlangen, Germany

**Adsorption of amino acids on MFI-type zeolite: Computational and experimental results (in German)**

K. Stueckenschneider, J. Merz, A. Zielesny and G. Schembecker

Chemie Ingenieur Technik (2012), 84, No. 8, 1413

[\(online\)](#)

- 2011 **Adsorption of amino acids on MFI-type zeolite: DFT calculations and experimental results**  
K. Stückenschneider, A. Zielesny and G. Schembecker  
Poster contribution to the „7th German Conference on Chemoinformatics” from November 6 to 8, 2011, in Goslar, Germany  
Journal of Cheminformatics (2012), 4(Suppl 1): P38  
([online](#))
- Reaction enumeration and machine learning enhancements for the open-source pipelining solution CDK-Taverna 2.0**  
A. Truszkowski, S. Neumann, A. Zielesny, E. L. Willighagen and C. Steinbeck  
Poster contribution to the „9th International Conference on Chemical Structures (ICCS)” from June 5 to 9, 2011, in Noordwijkerhout, The Netherlands
- 2010 **CDK-Taverna 2.0: migration and enhancements of an open-source pipelining solution**  
A. Truszkowski, S. Neumann, A. Zielesny, E. Willighagen and C. Steinbeck  
Poster contribution to the „6th German Conference on Chemoinformatics” from November 7 to 9, 2010, in Goslar, Germany  
Journal of Cheminformatics (2011), 3 (Suppl 1): P5  
([online](#))
- 2009 **Molecular fragments chemoinformatics**  
H. Kuhn, S. Neumann, C. Steinbeck, C. Wittekindt and A. Zielesny  
Poster contribution to the „5th German Conference on Chemoinformatics” from November 8 to 10, 2009, in Goslar, Germany  
Journal of Cheminformatics (2010), 2 (Suppl 1): P14  
([online](#))
- 2008 **Creating Chemo- & Bioinformatics Workflows: Further Developments within the CDK-Taverna Project**  
T. Kuhn, A. Zielesny and C. Steinbeck  
Poster contribution to the „4th German Conference on Chemoinformatics” from November 9 to 11, 2008, in Goslar, Germany  
Chemistry Central Journal (2009), 3 (Suppl 1): P42  
([online](#))
- Creating Chemo- & Bioinformatics Workflows: Further Developments within the CDK-Taverna Project**  
T. Kuhn, A. Zielesny and C. Steinbeck  
Poster contribution to the „8th International Conference on Chemical Structures (ICCS)” from June 1 to 5, 2008, in Noordwijkerhout, The Netherlands
- 2007 **Creating Chemo- and Bioinformatics Workflows – Building a Soft Computing Framework for the CDK-Taverna Project**  
T. Kuhn, A. Zielesny and C. Steinbeck  
Poster contribution to the „3. German Conference on Chemoinformatics“ from November 11 to 13, 2007, in Goslar, Germany,  
Chemistry Central Journal (2008), 2 (Suppl 1): P27  
([online](#))

- 2007 **Intelligent IT-Systems? Challenges, Fakes and Hard Science**  
S. Neumann, A. Zielesny  
Proceedings of the „2007 International Chemical Information Conference & Exhibition (ICIC)”, October 21 to 24, 2007, Sitges/Barcelona, Spain
- Understanding Cucurbit[6]uril-Spermine**  
H.-J. Buschmann, A. Wego, A. Zielesny, E. Schollmeyer  
Poster contribution to the 393. WE-Heraeus-Seminar „Trends in Molecular Biophysical Spectroscopy – Electronic Structure, Function, and Dynamics of Biomolecules“ from April 26 to 28, 2007, in Bad Honnef, Germany
- 2006 **Creating Chemo- and Bioinformatics Workflows – The CDK-Taverna Project**  
T. Kuhn, E. Willighagen, A. Zielesny and C. Steinbeck  
Poster contribution to the „2nd German Conference on Chemoinformatics“ from November 12 to 14, 2006, in Goslar, Germany
- 2005 **Understanding the Cucurbit[6]uril-Spermine-Complex: Geometry and Electronic Interactions**  
H.-J. Buschmann, A. Wego, A. Zielesny, E. Schollmeyer  
Poster contribution to the „1st German Conference on Chemoinformatics“ from November 13 to 15, 2005, in Goslar, Germany
- Structure, Electronic Properties and NMR-Shielding of Cucurbit[n]urils**  
H.-J. Buschmann, A. Wego, A. Zielesny, E. Schollmeyer  
Poster contribution to the „1st German Conference on Chemoinformatics“ from November 13 to 15, 2005, in Goslar, Germany
- A Case Study for Information Integration and Analysis with Wistract® - A new Platform for Scientific Information**  
C. Schaub, A. Zielesny  
Proceedings of the „2005 International Chemical Information Conference & Exhibition (ICIC)”, October 16 to 19, 2005, Nîmes, France
- 2004 **Combinatorial Formulation and Testing of Adhesives**  
K. U. Koch, H. Frenz, A. Zielesny  
Poster contribution to the „International Conference on High Throughput Formulation Technologies“ of the DECHEMA from December 7 to 8, 2004, in Frankfurt am Main, Germany
- Wistract® – Platform for Scientific Information**  
O. Gebert, C. Schaub, A. Zielesny  
Poster contribution to the „18th CIC-Workshop: New Developments in Chemoinformatics“ of the section Chemistry-Information-Computer (CIC) of the German Chemical Society from November 14 to 16, 2004, at Boppard am Rhein, Germany
- 2003 **Prediction of Curing Times of Adhesives with Neural Networks (in German)**  
K. U. Koch, A. Zielesny  
Poster contribution to the „Annual Conference 2003“ of the German Chemical Society from October 6 to 11, 2003, in Munich

- 2003 **Iterated Rank based Methods for Clustering**  
S.W. Perrey, H. Brinck, A. Zielesny  
Proceedings of the CSB 2003: IEEE Computer Society Bioinformatics Conference (2003, Stanford), 478-479
- 2002 **Neural-Network based IT Methods for Job-Man Matching – the personalized Internet Labour Portal** (in German)  
H. M. Stindt, A. Zielesny  
Workshop „Modern Services for the Labour Market“ of the „Hartz-Committee“ of the German Government, May 17, 2002, in Berlin
- 2000 **The ChemInformatics Integration Challenge**  
M. Diedrich, W. Jilge, M. Neumann, H. Rakel, A. Zielesny  
Proceedings of the „Drug Discovery Technology (DDT) 2000“, Boston, USA
- 1999 **Concept and Realization of Bayer's Integrated Chemistry Information System on the Corporate Intranet**  
A. Zielesny  
Proceedings of the „Chemistry and the Internet (ChemInt) 1999“, Washington DC, USA
- 1998 **Development of a Web-based chemical information workspace at Bayer: Review and perspectives for R&D**  
A. Zielesny, W. Jilge  
Proc. Int. Chem. Inf. Conf. (1998), 112-119 (Proceedings of the „1998 International Chemical Information Conference & Exhibition (ICIC)“, Nîmes, France)
- 1997 **Strategies and paradigms of scientific end-user information at Bayer**  
A. Zielesny, W. T. Donner, W. Jilge  
Proceedings of the idt 97, Paris, France
- 1996 **Scientific PC-based end-user information workspace: New developments at Bayer**  
A. Zielesny, M. Becker, A. Köhler, J. Sendelbach, C. Zirz, W. T. Donner  
Proc. Int. Chem. Inf. Conf. (1996), 125-129 (Proceedings of the „1996 International Chemical Information Conference & Exhibition (ICIC)“, Nîmes, France)
- 1995 **Use of the Beilstein Information at Bayer** (in German)  
C. Zirz, J. Sendelbach, A. Zielesny  
Proceedings of the „17. Online-Tagung der DGD“, Frankfurt am Main, May 16 to 18, 1995

# Scientific Education

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## *Cooperative Dissertation*

- **Development of deep learning applications for the automated extraction of chemical information from scientific literature**

Student: Henning Otto Brinkhaus  
Year: 2023

- **Development and implementation of *in silico* molecule fragmentation algorithms for the cheminformatics analysis of natural product spaces**

Student: Jonas Schaub  
Year: 2023

- **DARLING: Deep leARning for chemicalL Information processing**

Student: Kohulan Rajan  
Year: 2021

- **Development of an Open Biomolecular Mesoscopic Simulation Environment for the Study of Cyclotide/Membrane Interactions**

Student: Karina van den Broek  
Year: 2021

- **Simulation of Peptides, Proteins and Biomembranes with Molecular Fragment Dynamics (MFD)**

Student: Andreas Truszkowski  
Year: 2015

- **Open Source Workflow Engine for Cheminformatics: From Data Curation to Data Analysis**

Student: Thomas Kuhn  
Year: 2009

## **Master Thesis**

- **Standardizing the Natural Product Chemical Space**

Student: Samuel Behr  
Year: 2023

- **Exploring the natural product space with fragment fingerprinting and clustering**

Student: Betül Sevindik  
Year: 2023

- **Implementation of Molecular Scaffold Tree Functionality in the Chemistry Development Kit (CDK)**

Student: Julian Zander  
Year: 2022

- **Machine learning models for biomolecular interaction parameters**

Student: Lisa Sophie Kersten  
Year: 2021

- **Evaluation of machine learning approaches to support biomolecular simulation**

Student: Ann-Kathrin Brüggemann  
Year: 2021

- **Molecule Set Comparator – A Cheminformatics Tool for the Evaluation of Machine Learning based Biomolecule Recognition**

Student: Jan-Mathis Hein  
Year: 2020

- **Design of a computational Information and Analysis Platform for Lead Discovery**

Student: Jonas Schaub  
Year: 2019

- **Decision Support for Acute Lymphoblastic Leukemia (ALL) Diagnostics**

Student: Felix Bänsch  
Year: 2018

- **Implementation of an algorithm for the identification of functional groups in organic molecules for the Chemistry development Kit (CDK)**

Student: Sebastian Fritsch  
Year: 2018

- **Investigation of the Interaction between the Cyclotide Cycloviolacin O2 and Phospholipid Bilayers on the basis of Mutation Studies with Molecular Fragment Dynamics (MFD)**

Student: Sarah Albert  
Year: 2017

- **Molecular Fragment Dynamics Study of Biological Membranes and Zinc Ricinoleate Layers for Enzyme Purification**

Student: Karina van den Broek  
Year: 2014

- **Enabling molecular Biosciences: Re-engineering, reaction enumeration and machine learning enhancements for the open workflow platform CDK-Taverna**

Student: Andreas Truszkowski  
Year: 2011

## **Bachelor Thesis**

- **Protein structure prediction evaluation using PySSA**

Student: Martin Urban  
Year: 2025

- **AI-assisted Development of a Descriptor Enrichment Module for Natural Product Analysis with the Molecular Fragmentation System MORTAR**

Student: Manuel Schauer  
Year: 2025

- **Development of the end-user protein structure prediction system PySSA**

Student: Hannah Kullik  
Year: 2025

- **Exhaustive fragmentation of Natural Products using the MORTAR-System**

Student: Tom Weiß  
Year: 2025

- **Implementation of a modified ART 2-A Network for Clustering of Natural Product Spaces**

Student: Zeynep Dagtekin  
Year: 2024

- **Successive algorithmic fragmentation of alkyl and cycloalkyl groups in natural products**

Student: Maximilian Rottmann  
Year: 2023

- **Development of a Rich-Client Graphical User Interface (GUI) for Force Field based Biomolecular Mesoscopic Parameter Estimation**

Student: Veit Hucke  
Year: 2021

- **Extension of the Molecular Fragmentation Framework MORTAR**

Student: Samuel Behr  
Year: 2021

- **Development of a Rich-Client Graphical User Interface (GUI) for Machine Learning based Biomolecular Mesoscopic Parameter Estimation**

Student: Jana Wilms  
Year: 2021

- **Molecular Fragmentation Heuristics for Natural Products**

Student: Betül Sevindik  
Year: 2021

- **Automated molecular fragmentation for mesoscopic simulation based on functional groups**

Student: Julian Zander  
Year: 2019

- **Design and Generation of Animated Property Diagrams for Biomolecular Simulation**

Student: Jan-Mathis Hein  
Year: 2018

- **Simulation of realistic biomolecular Membrane Models with Dissipative Particle Dynamics**

Student: Lisa Sophie Kersten  
Year: 2018

- **Flexible Design of Protein Fragmentation Patterns for Biomolecular Simulation**

Student: Katrin Tietz  
Year: 2018

- **Development of an Open Software Component for Validation and Visualization of Fragment-Based Topologic Representation of complex Biomolecular Structures (Particle SMILES)**

Student: Jonas Schaub  
Year: 2017

- **Descriptor calculation for structural Bioinformatics based on the Chemistry Development Kit (CDK)**

Student: Felix Jonathan Bänsch  
Year: 2016

- **Investigation of the Interaction between Peptides/Proteins and Phospholipid Bilayers with Molecular Fragment Dynamics (MFD)**

Student: Sarah Albert  
Year: 2015

- **Development of parallelized machine learning algorithms for biological data analysis with Mathematica**

Student: Kolja Berger  
Year: 2014

- **Design and development of a rich client application for the calculation of chemical descriptors based on the Chemistry Development Kit (CDK)**

Student: Sebastian Fritsch  
Year: 2013

- **Design and development of a rich client application for compound selection based on the Chemistry Development Kit (CDK)**

Student: Hendrik Götdeke  
Year: 2013

- **Programming a neural Hopfield-Network for pattern recognition**

Student: Carina Goretzky  
Year: 2012

- **Dependency structures in protein expression data of invasive breast cancer**

Student: Florian Boecker  
Year: 2012

- **Extension of the Chemistry Development Kit (CDK) for biomedical Modelling of Quantitative Structure-Property Relationships**

Student: Julian Jorzik  
Year: 2011

- **Development and industrial validation of a combinatorial chemistry related enumeration software library for synthesis of potential pharmaceutical drugs on the basis of the Chemistry Development Kit (CDK)**

Student: Andreas Truszkowski  
Year: 2009

## **Diploma Thesis**

- **Synthesis of organozinc complexes for reaction with pollutants in gas phase: Molecular Modelling study of the reaction mechanism**

Student: Thomas Veidt  
Year: 2010

- **Alignment of molecular structures represented by Gaussian functions**

Student: Christian Geiger  
Year: 2008

- **Molecular Fragment Dynamics Simulations for the Aggregation of Phospholipids at solid Surfaces**

Student: Christoph Engels  
Year: 2007

- **Quantum-chemical Investigations on Cucurbit[6]uril and  $\alpha$ -Cyclodextrin Complexes**

Student: Jan-Niklas Schäfer  
Year: 2006

- **Software Design Aspects of an electronic Notebook for Materials Sciences**

Student: Holger Schulte  
Year: 2006

- **Development and implementation of algorithms and software components for the comparison and representation of biological sequences**

Student: Thomas Kuhn  
Year: 2005

- **Development, implementation and validation of an ART-based Neural Network software component for the solution of open-categorical classification problems in high dimensional scientific information spaces**

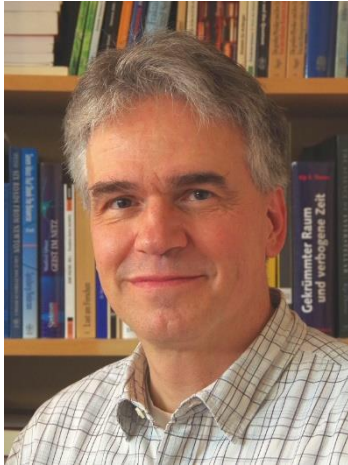
Student: Stefan Neuman  
Year: 2004

- **Integration of complex reaction mechanisms to simulation models and development of a model for hydrogenation in Trickle-Bed reactors**

Student: Emrah Ergörül  
Year: 2004

- **Partial purification, characterization and genotype determination of the acetylcholine esterase of different types of phorodon humuli**

Student: Sandra Bittner  
Year: 2003



## Prof. Dr. Achim Zielesny

Westphalian University of Applied Sciences  
Institute for Bioinformatics and Chemoinformatics (IBCI)

August-Schmidt-Ring 10  
D-45665 Recklinghausen, Germany

Phone +49-2361-915530  
E-Mail achim.zielesny@w-hs.de

Day and place of Birth | 1964/08/26 in Düsseldorf, Germany

## School, Job Training and Military Service

1984	German Abitur and degree as a biological technical assistant
1984/1985	Laboratory assistant in chemical analytics at the Medical Services Research Institute of the German Federal Armed Forces

## Academic Education

1986 – 1993	Study of chemistry at the University of Cologne, diploma and doctoral thesis at the Institute of Physical Chemistry (Prof. Dr. D. Woermann) of the University of Cologne  Scholarship holder of the German National Academic Foundation (1989 – 1991), Final-Degree Award of the German Chemical Industry Association (1993)
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## Job

1991 – 1994	Scientific assistant at the University of Cologne
1994 – 2001	Employee of the Bayer AG, Leverkusen (project and group manager in the central research, the computer science and the managerial human resources division)
Since 2001	Professor of Chemistry, Chemoinformatics and Bioinformatics at the Westphalian University of Applied Sciences

## Miscellaneous

2004	Founder of the <a href="#">GNWI</a> , a company for scientific computing solutions, Dortmund, Germany
Since 2005	Founder and chairman of the <a href="#">Institute for Bio- and Chemoinformatics</a> at the Westphalian University of Applied Sciences
2005 – 2012	Board member of the <a href="#">Computers in Chemistry (CIC)</a> section of the German Chemical Society (GDCh)
2005 – 2012	Member of the Scientific Advisory Board of the German Conference on Chemoinformatics (GCC)
2007 – 2020	Personal tutor of the <a href="#">German National Academic Foundation</a> (Studienstiftung des deutschen Volkes)
Since 2012	Board member of the <a href="#">Westphalian Institute of Health (WIGE)</a> and person in charge of the research focus Biomedical Modelling and Simulation
2016 – 2021	Founding member of the Life Sciences section of the Graduate Institute for Applied Research at Universities of Applied Sciences in North Rhine-Westphalia (Graduierteninstitut NRW)
Since 2017	Member of the selection committee for the Gmelin-Beilstein-Denkmuende award of the German Chemical Society (GDCh)
2018	Scientific coordinator of the 32nd Molecular Modelling Workshop (MMWS) 2018 in Erlangen, Germany
2019	Research stay at the Friedrich Schiller University Jena
Since 2021	Associated member of section Life Sciences and Health Technologies of the <a href="#">Doctoral Studies Program in North Rhine-Westphalia (Promotionskolleg NRW)</a>
Since 2021	Member of the <a href="#">Advisory Board Industry</a> of the <a href="#">Chemistry Consortium (NFDI4Chem)</a> in the <a href="#">NFDI</a> (Nationale Forschungsdateninfrastruktur)
Since 2021	Member of the organizational committee of the <a href="#">International Workshop on Open Molecular Informatics (IWOMI)</a>
2023	Research stay at the Friedrich Schiller University Jena
2024 – 2025	Member of the founding committee of the <i>Institute for Applied Artificial Intelligence</i> of the Westphalian University of Applied Sciences