

## Web-Based Simulation Tools in Chemistry

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### 1. Introduction

Online simulation platforms are web-based applications that enable users to engage with virtual models copying the real-world systems or scenarios directly through a web browser. These platforms facilitate interactive learning and training experiences without the need for specialized software installations. By providing a safe environment for experimentation, it allows users to manipulate variables, observe outcomes, and gain insights into complex processes.<sup>[1]</sup> Simulation is both an independent discipline and a versatile tool applied across various fields. Similar to mathematics and logic, it has its own theories and methods that researchers refine to enhance accuracy and effectiveness. While continuous advancements improve simulation models, its greatest value lies in practical applications. It is widely utilized in disciplines like chemistry, physics, engineering, and economics to test hypotheses, optimize processes, and predict.<sup>[2]</sup>

The integration of artificial intelligence (AI) and machine learning (ML) into online simulation platforms has further expanded their capabilities, enabling predictive modeling and automated analysis of complex chemical systems. For instance, AI-driven tools can accelerate drug discovery by screening millions of compounds virtually, while ML algorithms refine spectroscopic data interpretation.<sup>[3,4]</sup> These advancements democratize access to high-end computational resources, bridging gaps between theoretical research and practical applications. As cloud-based platforms evolve, they promise to transform chemical education and industrial R&D by offering scalable, collaborative, and cost-effective solutions.<sup>[2]</sup>

### 2. Discussion

#### 2.1 Online Simulation Platforms in the field of Chemistry

With the continuous advancements in the internet, web technologies, and computational tools, numerous online platforms have been developed to enhance the efficiency and convenience of studying complex molecular interactions. These platforms provide students and researchers in the field of chemistry with innovative tools to evaluate, analyze, and simulate intricate chemical processes with greater accuracy and speed. Among the most impactful online simulations are molecular docking, virtual labs, spectroscopy and reaction simulations, and cloud-based computational chemistry tools. By integrating artificial intelligence, machine learning, and cloud computing, these platforms revolutionize chemical research and education, enabling breakthroughs in drug discovery, material science, and industrial chemistry.

##### 2.1.1 Virtual Laboratory

A virtual lab is a computer-based simulation environment that replicates real-world laboratory experiments and procedures, allowing users to conduct scientific investigations, analyze data, and test hypotheses without physical laboratory equipment. These platforms are particularly useful in fields like chemistry, physics, biology, engineering, and medicine. For example, PhET Interactive Simulations offers a range of interactive science and math simulations, allowing users to engage in experiments that enhance their understanding of complex concepts. Similarly, Labster provides virtual lab experiences in biology and other sciences, allowing students to safely conduct experiments without the risks and limitations of a real-world lab. Overall, virtual labs make experiments more accessible, offering a valuable resource for both learners and researchers.<sup>[5,6,7,8]</sup>

### 2.1.2 Molecular Docking

Molecular docking is a computational technique used to predict how ligands, such as small molecules or proteins, interact with receptors like proteins or nucleic acids. It estimates the ligand's position within a binding site and is essential in drug discovery and molecular research. A key application of molecular docking is virtual screening, where compounds are tested against a target protein to identify potential drug candidates. The process involves predicting ligand orientation and stability using a scoring function, with the goal of accurately replicating experimental binding modes and ranking favorable poses. Despite challenges in achieving high accuracy due to external factors influencing binding interactions, molecular docking remains a vital tool in drug discovery and biomolecular studies.<sup>[4][9]</sup>

Several molecular docking software tools are integral to the drug discovery process. Glide (Schrödinger) is recognized for its precision and efficiency in predicting ligand-protein interactions, offering two distinct precision modes, Standard Precision (SP) and Extra Precision (XP), to tailor studies. AutoDock Vina (Scripps Research Institute) utilizes genetic algorithms and empirical scoring functions to predict ligand-receptor interactions, with recent advancements in its versions, including QuickVina2 and Smina, which enhance its search and scoring capabilities. GOLD (Cambridge Crystallographic Data Centre) employs a genetic algorithm to optimize ligand binding poses, accommodating flexible ligands and protein structures to improve docking accuracy. LeDock (University of Paris Diderot and CNRS) integrates geometric matching and optimization algorithms to match ligands to protein binding sites, with the added advantage of being an open-source tool that allows for customization and integration into larger computational workflows. These tools are essential in virtual screening, lead optimization, and predicting ligand binding interactions, providing researchers with reliable simulations to support drug discovery and development. Avogadro is an advanced open-source molecular editor and visualizer used in computational chemistry, molecular modeling, bioinformatics, and materials science. It features high-quality rendering, a powerful plugin architecture, real-time bond and atom manipulation, and an intuitive 3D molecular modeling interface.<sup>[10,11,12,13] [3]</sup>

### 2.1.3 Spectroscopy and Reaction Simulation

Computational chemistry employs spectroscopy and reaction simulation to understand molecular behavior and reaction mechanisms scientifically. Computational spectroscopy simulates electromagnetic radiation's interaction with matter, predicting and interpreting spectral data. This allows researchers to analyze vibrational, rotational, and electronic transitions within molecules, aiding in compound identification and structural elucidation. Quantum mechanical calculations enable accurate spectral modeling, enhancing the understanding of experimental observations and facilitating the design of new materials and drugs.<sup>[14]</sup>

Reaction simulation uses computational methods to model chemical reactions at the molecular level. This involves calculating potential energy surfaces and reaction pathways to predict reaction rates, intermediates, and products. Simulations provide insights into reaction mechanisms, allowing for optimization of reaction conditions and catalyst design. This approach is particularly useful for complex systems where experimental investigation is difficult.<sup>[15]</sup> The accuracy of both techniques relies heavily on the chosen level of theory and basis set used in the quantum mechanical calculations. Different levels of theory offer varying degrees of accuracy and computational cost, requiring careful consideration based on the specific system and research question.<sup>[16]</sup> Furthermore, the interpretation of results requires a strong understanding of both computational methods and experimental techniques to ensure accurate and meaningful conclusions.

### 2.1.4 Cloud-based computational chemistry tools

Cloud-based computational chemistry tools have significantly enhanced the accessibility and scalability of chemical simulations. These platforms leverage cloud computing resources, providing high-performance computing capabilities without the need for extensive local infrastructure.<sup>[17]</sup> This

allows researchers to perform complex calculations, such as electronic structure computations, molecular dynamics simulations, and data analysis, via web interfaces. This accessibility fosters collaboration and resource sharing among researchers across the globe.

Examples of cloud-based computational chemistry tools include Gaussian OnDemand and Schrödinger LiveDesign. Gaussian OnDemand provides access to the Gaussian software package for electronic structure modeling, enabling researchers to study the properties and behaviors of molecular systems.<sup>[19]</sup> Schrödinger LiveDesign offers a collaborative environment for designing and analyzing chemical compounds, integrating various computational methods to support drug discovery and materials science.<sup>[18]</sup> The use of cloud-based platforms has democratized to high-performance computing, enabling researchers with limited resources to perform computationally intensive simulations. This has led to significant advancements in various fields, including materials science, drug discovery, and environmental chemistry.

**Table 1. Types of online simulation platforms in the field of chemistry**

Online Simulation Platforms	Purpose	Example
Virtual Lab	A simulated environment for conducting laboratory experiments digitally	Phet and Labster
Molecular Docking	To predict the ligand-receptor binding for drug discovery	Glide, Autodock Vina, GOLD, Ledock
Spectroscopy and Reaction Simulation	Digital tools that model spectroscopic analysis and chemical reactions to predict outcomes, interpret data	Color Meter - RGB HSL CMYK RYB
Cloud-based computational chemistry tools	Online platforms for simulating molecular structures, reactions, and properties using advanced computational methods	Gaussian OnDemand and Schrödinger LiveDesign

### *2.1 Significance of Cloud-based Spectroscopy and Reaction Modeling tools in Chemistry*

Cloud-based spectroscopy and reaction modeling tools have significantly enhanced the efficiency and collaboration capabilities in the field of chemistry. Platforms like ChemSpectra provide web-based solutions for visualizing and analyzing spectroscopic data, supporting various techniques such as infrared spectroscopy (IR), mass spectrometry (MS), and nuclear magnetic resonance (NMR) spectroscopy. In reaction modeling, tools like Reaction Lab™ leverage mechanistic modeling to predict reaction behaviors based on fundamental physical chemistry principles. This approach allows chemists to optimize reactions with limited experimental data, facilitating a deeper understanding of chemical kinetics and aiding in process development.

Additionally, platforms such as ChemReaX offer free web applications for modeling and simulating basic chemical reactions, making them valuable resources for educational purposes and preliminary research. These cloud-based tools not only streamline data analysis and storage but also promote collaboration among researchers by providing accessible and scalable solutions for complex chemical analyses.<sup>[19,20,21]</sup>

### *2.2 Impact of machine Learning in Chemical Simulation*

Machine learning (ML) impacts chemical simulations by accelerating computations, enabling the analysis of complex phenomena, and facilitating the discovery of new materials and reactions, ultimately leading to more efficient and insightful research. It can accelerate computations or calculations by enabling machine learning algorithms to learn from large datasets and simulation data,

allowing faster and more efficient calculations. It can also bridge the gap between different length and time scales in chemical simulations, allowing for the study of complex systems that are otherwise intractable.<sup>[22,23]</sup>

### 3. Conclusion

This study presents a comprehensive review of web-based simulation platforms in chemistry, highlighting their transformative impact on research, education, and industrial applications. These platforms have significantly improved accessibility to advanced tools for modeling molecular interactions and chemical processes. Various types of simulation platforms are examined, including virtual laboratories, molecular docking tools, online databases, spectroscopy and reaction simulators, and cloud-based computational chemistry resources. The integration of these technologies has enhanced the efficiency and accuracy of chemical studies while democratizing access to high-performance computing capabilities.

Furthermore, the incorporation of artificial intelligence and machine learning into these web-based tools has expanded their capabilities, enabling more sophisticated predictive modeling and data analysis in chemistry. Traditional experimental and analytical approaches have been reshaped by these advancements, fostering more sustainable, collaborative, and cost-effective methodologies. As outlined in the study, cloud-based spectroscopy tools and computational chemistry platforms have played a pivotal role in streamlining workflows, improving data accessibility, and increasing the precision of chemical simulations. These developments have ultimately contributed to significant advancements in drug discovery, materials science, and environmental chemistry.

### Reference:

- [1] Rapin, N., Lund, O., & Castiglione, F. (2011). Immune system simulation online. *Bioinformatics*, 27(14), 2013-2014.
- [2] Ören, T., Mittal, S., & Durak, U. (2019). Modeling and simulation: the essence and increasing importance. *Modeling and simulation of complex communication networks*, 3-26.
- [3] Jiang, D., Zhao, H., Du, H., Deng, Y., Wu, Z., Wang, J., ... & Hou, T. (2023). How Good Are Current Docking Programs at Nucleic Acid–Ligand Docking? A Comprehensive Evaluation. *Journal of Chemical Theory and Computation*, 19(16), 5633-5647.
- [4] Stanzione, F., Giangreco, I., & Cole, J. C. (2021). Use of molecular docking computational tools in drug discovery. *Progress in medicinal chemistry*, 60, 273-343.
- [5] Kadri, M., Boubakri, F. E., Teo, T., Kaghat, F. Z., Azough, A., & Zidani, K. A. (2024). Virtual reality in medical education: Effectiveness of Immersive Virtual Anatomy Laboratory (IVAL) compared to traditional learning approaches. *Displays*, 85, 102870.
- [6] Smith, C. L., & Coleman, S. K. (2017, May). Using Labster to improve Bioscience student learning and engagement in practical classes. In *Heads of Biological Sciences, Royal Society of Biology. Spring 2017 meeting*.
- [7] Perkins, K., Adams, W., Dubson, M., Finkelstein, N., Reid, S., Wieman, C., & LeMaster, R. (2006). PhET: Interactive simulations for teaching and learning physics. *The physics teacher*, 44(1), 18-23.
- [8] Tripepi, M. (2022). Microbiology laboratory simulations: from a last-minute resource during the Covid-19 Pandemic to a valuable learning tool to retain—a semester microbiology laboratory

curriculum that uses Labster as prelaboratory activity. *Journal of Microbiology & Biology Education*, 23(1), e00269-21.

[9] Meza Menchaca, T., Juárez-Portilla, C., & C. Zepeda, R. (2020). Past, Present, and Future of Molecular Docking. *Drug Discovery and Development - New Advances*. <https://doi.org/10.5772/intechopen.90921>

[10] Ivanova, L., & Karelson, M. (2022). The impact of software used and the type of target protein on molecular docking accuracy. *Molecules*, 27(24), 9041.

[11] Valdés-Tresanco, M. S., Valdés-Tresanco, M. E., Valiente, P. A., & Moreno, E. (2020). AMDock: a versatile graphical tool for assisting molecular docking with Autodock Vina and Autodock4. *Biology direct*, 15, 1-12.

[12] Kamal, I. M., & Chakrabarti, S. (2023). MetaDOCK: a combinatorial molecular docking approach. *ACS omega*, 8(6), 5850-5860.

[13] Hanwell, M. D., Curtis, D. E., Lonie, D. C., Vandermeersch, T., Zurek, E., & Hutchison, G. R. (2012). Avogadro: an advanced semantic chemical editor, visualization, and analysis platform. *Journal of cheminformatics*, 4, 1-17.

[14] Jansen, T. L. C. (2021). Computational spectroscopy of complex systems. *The Journal of Chemical Physics*, 155(17), 170901. <https://doi.org/10.1063/5.0064092>

[15] Kang, P.-L., & Liu, Z.-P. (2021). Reaction prediction via atomistic simulation: From quantum mechanics to machine learning. *iScience*, 24(1), 102013. <https://doi.org/10.1016/j.isci.2020.102013>

[16] Soroush, M., & Grady, M. C. (2019). Polymers, polymerization reactions, and computational quantum chemistry. In M. Soroush (Ed.), *Computational quantum chemistry* (pp. 1-16). Elsevier. <https://doi.org/10.1016/B978-0-12-815983-5.00001-5>

[17] Hassan, E., Mustafa, Y., & Merkhani, M. (2022). Computation in chemistry: Representative software and resources. 6, 1-22.

[18] Cox, P. B., & Gupta, R. (2022). Contemporary Computational Applications and Tools in Drug Discovery. *ACS medicinal chemistry letters*, 13(7), 1016–1029. <https://doi.org/10.1021/acsmmedchemlett.1c00662>

[19] Huang, Y. C., Tremouilhac, P., Nguyen, A., Jung, N., & Bräse, S. (2021). ChemSpectra: a web-based spectra editor for analytical data. *Journal of Cheminformatics*, 13(1), 8.

[20] all, I. (2025, February 7). *Reaction Lab*. Mt.com. [https://www.mt.com/ca/en/home/products/L1\\_AutochemProducts/scale-up-systems/reaction-lab.html](https://www.mt.com/ca/en/home/products/L1_AutochemProducts/scale-up-systems/reaction-lab.html)

[21] Banks, P. (2017, November 30). *Science by Simulation: ChemReaX*. RSC Education. <https://edu.rsc.org/review/science-by-simulation-chemreax/3008346.article>

[22] Killingsworth, N. J., McNenly, M. J., Whitesides, R. A., & Wagnon, S. W. (2020). Cloud based tool for analysis of chemical kinetic mechanisms. *Combustion and Flame*, 221, 170–179.

[23] Ravinder Dogra, Diwaker, Praveen Kumar, Samjeet Singh Thakur. Role of Cloud Computing in Chemistry. *International Journal of Chemical Engineering and Processing* 2024; 10(2): 1–15p