iJRASET
International Journal For Research in Applied Science and Engineering Technology



INTERNATIONAL JOURNAL FOR RESEARCH

IN APPLIED SCIENCE & ENGINEERING TECHNOLOGY

Volume: 14 **Issue:** VI **Month of publication:** June 2026

DOI: https://doi.org/10.22214/ijraset.2026.83443

www.ijraset.com

Call: 08813907089 | E-mail ID: ijraset@gmail.com

International Journal for Research in Applied Science & Engineering Technology (IJRASET)
ISSN: 2321-9653; IC Value: 45.98; SJ Impact Factor: 7.538
Volume 14 Issue VI June 2026- Available at www.ijraset.com

Review On: Computational Aided Drug Design for Polyherbal Formulations

Miss .Asmita D Tengle[1], Miss. Ketakee R. More[2], Dr. Nitin M Gawai[3], Miss.Sayali R Bhalsen[4], Miss.Vaidehi V Tukurul[5]
B . Pharmacy Department, Mahadev Kanchan College Of Pharmaceutical Education And Research
Ajinkya Charitable Foundation , Uruli Kanchan , Pin Code – 412202, Pune , India

Abstract: ***Polyherbal formulations are widely used in traditional medicine because they combine multiple botanicals to improve therapeutic efficacy, broaden biological coverage, and potentially reduce toxicity compared with single-herb preparations . However, their scientific development is constrained by multicomponent complexity, uncertain mechanisms, variable phytochemical profiles, and the difficulty of selecting optimal herb ratios. Computational aided drug design (CADD) offers a practical way to address these limitations by enabling virtual screening, molecular docking, pharmacophore modeling, network pharmacology, and synergy prediction before experimental validation . In recent years, machine learning-based approaches have expanded rapidly and are increasingly used to predict synergistic drug combinations from high-throughput data . For polyherbal systems, these tools can prioritize bioactive compounds, identify multi-target interactions, and support formulation optimization. This review discusses the conceptual basis of synergy in polyherbal medicine, outlines major CADD strategies applicable to herbal formulations, and examines computational methods for synergy prediction. It also highlights the role of systems biology and artificial intelligence in advancing evidence-based herbal drug development. Finally, the review addresses limitations such as data standardization, model interpretability, and translational barriers, while proposing future directions for precision herbal medicine.***

Keywords: ***Polyherbal formulations, computer-aided drug design, synergy prediction, molecular docking, network pharmacology, machine learning, phytochemicals***

I. INTRODUCTION

Polyherbal formulation refers to the use of two or more herbs in a single preparation to achieve a desired therapeutic effect . In traditional medicine systems, especially Ayurveda, the combination of herbs is not accidental but is often based on the idea that multiple ingredients can work together through complementary mechanisms . Some components may act directly on the disease target, while others may improve absorption, stability, or metabolism of co-administered constituents. This makes polyherbal medicine an attractive concept for complex diseases that involve multiple pathways.

At the same time, polyherbal formulations present major scientific challenges. They contain many phytochemicals, many of which may be active, inactive, or only weakly active in isolation. Their biological effects may depend on interactions among constituents, and these interactions may be synergistic, additive, or antagonistic. Conventional trial-and-error methods are not sufficient to analyze such systems efficiently. For this reason, computational aided drug design has become increasingly important in herbal research . CADD allows researchers to model targets, screen compounds, and predict likely interactions before performing laboratory experiments.

This review focuses on the use of CADD in polyherbal formulation development. It explains the role of synergy, summarizes major computational strategies, and reviews synergy prediction models and emerging approaches. The goal is to provide a rational framework for designing evidence-based polyherbal products using modern computational methods.

II. CONCEPT OF SYNERGY

Synergy is the core concept underlying polyherbal formulation design. In simple terms, synergy means that the combined effect of two or more agents is greater than the sum of their individual effects . In herbal medicine, synergy can arise in several ways. One herb may enhance the activity of another at the receptor or pathway level, while another may improve pharmacokinetic behavior by increasing absorption or reducing degradation .

©IJRASET: All Rights are Reserved

ISSN: 2321-9653; IC Value: 45.98; SJ Impact Factor: 7.538
Volume 14 Issue VI June 2026- Available at www.ijraset.com

Synergy is usually divided into pharmacodynamic and pharmacokinetic forms. Pharmacodynamic synergy occurs when different ingredients affect the same disease process through distinct targets or pathways. Pharmacokinetic synergy occurs when one ingredient influences the absorption, distribution, metabolism, or elimination of another . Both forms are highly relevant to herbal formulations and can explain why whole mixtures sometimes outperform isolated compounds.

From a modern research perspective, synergy should be measured rather than assumed. Mathematical models such as Bliss independence, Loewe additivity, Highest Single Agent, and ZIP are widely used in drug-combination research to distinguish synergy from additivity . These models can be adapted to polyherbal studies when experimental dose-response data are available. This is important because a mixture that appears effective may simply reflect the action of one strong constituent rather than true combination synergy.

III. CADD STRATEGIES

Computer-aided drug design includes a family of methods that use computational tools to identify, analyze, and optimize biologically active compounds . In herbal research, CADD is valuable because it can reduce the large search space created by multicomponent mixtures. Rather than testing every possible combination in the laboratory, researchers can first use in silico methods to identify promising compounds and targets.

IV. STRUCTURE-BASED DESIGN

Structure-based drug design relies on the three-dimensional structure of the target protein or receptor. Candidate phytochemicals are docked into the target binding site to estimate binding affinity and interaction patterns . In polyherbal research, this can reveal whether compounds from different herbs bind the same target or complementary targets. Molecular docking is often the first step because it is fast and cost-effective.

V. LIGAND-BASED DESIGN

Ligand-based design is useful when the target structure is unknown or not sufficiently reliable. It uses known active compounds to build pharmacophore models, similarity searches, or QSAR models. This approach is especially helpful in natural product research because many phytochemicals have known biological effects but incomplete structural biology data. Ligand-based methods can prioritize analogs or help identify key chemical features responsible for activity.

VI. VIRTUAL SCREENING

Virtual screening allows researchers to test large libraries of compounds against selected targets. In polyherbal systems, the compound library may include phytochemicals from all herbs in the formula. Screening can identify molecules that contribute most strongly to the observed therapeutic effect. It can also help remove compounds with poor developability by integrating ADMET filtering and toxicity prediction.

VII. NETWORK PHARMACOLOGY

Network pharmacology is highly suitable for polyherbal research because it matches the multi-component and multi-target nature of herbal medicine . Instead of focusing on a single receptor, network pharmacology maps the relationships among compounds, targets, pathways, and diseases. This makes it possible to understand how a formula may act across several biological levels.

The typical workflow begins with collecting phytochemical data from the herbs in the formulation. Targets are then predicted using databases or docking results. Disease-related genes are gathered from biomedical databases, and overlapping targets are identified. These targets are then connected in a network to highlight hub genes and key pathways. Such analyses are especially useful in cancer, inflammation, diabetes, and other complex disorders where single-target intervention is often insufficient .

In polyherbal formulation research, network pharmacology can serve as a bridge between traditional knowledge and molecular mechanism. It helps explain why multiple herbs may work better together than separately. However, the method depends heavily on data quality. Errors in phytochemical annotation, incomplete target databases, or overly broad pathway lists can affect reliability. Therefore, network pharmacology should be combined with docking, simulation, and experimental validation.

©IJRASET: All Rights are Reserved | SJ Impact Factor 7.538 | ISRA Journal Impact Factor 7.894 |

VIII. SYNERGY PREDICTION

Synergy prediction is one of the most important emerging areas in computational drug combination research. Recent reviews show strong growth in machine learning methods for predicting synergistic combinations, especially using high-throughput screens . Although most studies focus on synthetic drugs, the same principles can be used for herbal combinations and polyherbal formulas.

Classical synergy models remain important because they provide interpretable benchmarks. Bliss independence assumes that two agents act independently. Loewe additivity compares the observed effect with the expected effect under dose equivalence. Highest Single Agent compares the combination with the most effective single ingredient. ZIP estimates the deviation from a non-interaction baseline . These models can help determine whether a pair of herbs or phytochemicals truly shows synergy.

Machine learning adds a predictive layer. Deep learning models can integrate chemical structure, gene expression, and response data to estimate likely synergy before laboratory testing . For polyherbal formulations, this could involve phytochemical fingerprints, target networks, and pathway signatures. Such models may eventually support formula design by identifying the best herb combinations and ratios.

IX. AI AND MACHINE LEARNING

Artificial intelligence is rapidly changing the drug discovery landscape, and herbal medicine research is beginning to benefit from these tools. AI can process large and complex datasets, identify hidden relationships, and support prediction in systems where human interpretation is difficult. In ethnopharmacology and natural product research, AI has been described as a useful tool for combining molecular docking, data mining, and other discovery methods .

One important use of AI is property prediction. Models can estimate ADMET behavior, toxicity, and bioactivity directly from molecular structure. This helps researchers prioritize compounds with greater development potential. AI can also assist in target prediction, where algorithms infer likely protein targets from chemical features. For polyherbal formulations, this is useful because a mixture can contain many compounds with overlapping or complementary activities.

Another emerging application is formulation optimization. Genetic algorithms and neural networks can be used to search for the best ratios of herbs or phytochemicals under defined constraints. In principle, such systems could balance efficacy, toxicity, and stability at the same time. However, AI models in herbal research are limited by the quality and size of available datasets. Because herbal databases remain incomplete, AI should be treated as a decision-support tool rather than a replacement for experimental science .

X. VALIDATION AND STANDARDIZATION

Computational predictions must always be validated experimentally. A strong research workflow should move from in silico analysis to in vitro testing, then to in vivo studies, and finally to clinical evaluation where appropriate. Docking scores and network predictions are useful for hypothesis generation, but they are not sufficient evidence on their own. This is especially important in herbal research because complex mixtures can easily produce misleading results.

Standardization is another major issue. Polyherbal formulations may vary depending on plant origin, harvesting season, extraction process, and storage conditions. These differences can change the phytochemical profile and affect computational predictions. For reproducible research, standardized compound libraries, reliable target databases, and transparent reporting of model parameters are essential. Without these controls, CADD outputs may not be comparable across studies.Regulatory translation is also challenging. Even when a polyherbal formula appears promising in silico, it still requires evidence of quality, safety, and consistency. Computational tools can support regulatory submission by identifying likely active markers and explaining mechanism of action. However, they cannot replace toxicity testing, quality control, or human studies. The best approach is therefore a combined pipeline in which computation guides experiment, and experiment confirms computation.

XI. FUTURE DIRECTIONS

The future of polyherbal formulation design will likely depend on integrating multiple computational approaches. Structure-based methods, ligand-based modeling, network pharmacology, and machine learning each contribute different strengths. Together, they can create a more rational and efficient development pipeline for herbal medicine. This is especially valuable for researchers working on topical formulations, syrups, and other multicomponent dosage forms.

One important future direction is multi-omics integration. Transcriptomics, proteomics, metabolomics, and microbiome data can help explain the broader biological response to a polyherbal formula.

©IJRASET: All Rights are Reserved | SJ Impact Factor 7.538 | ISRA Journal Impact Factor 7.894 |

Another direction is personalized herbal medicine, where formulas are selected or optimized according to the patient's biological profile. These approaches may be particularly useful for complex disorders such as PCOS, inflammatory skin disease, and metabolic syndrome.

More advanced approaches, such as quantum-inspired modeling and hybrid AI workflows, may also become relevant . These methods could improve scoring accuracy and help deal with the complexity of multicomponent mixtures. At the same time, the field needs better open databases, standardized workflows, and reproducible validation studies. Without high-quality data, even the best computational tools will remain limited.

XII. CONCLUSION

Computational aided drug design provides a strong framework for rational polyherbal formulation development . It can help identify active compounds, map targets, predict synergy, and prioritize combinations for experimental testing. This is especially useful in traditional medicine, where formulas contain many ingredients and the mechanism of action is often unclear .

The central value of CADD is that it transforms polyherbal research from an empirical process into a more evidence-based one. Synergy prediction, network pharmacology, and machine learning can improve formulation design, but only when supported by good data and experimental validation . For researchers developing herbal syrups, creams, or other dosage forms, this approach offers a practical and modern path toward scientifically validated products.

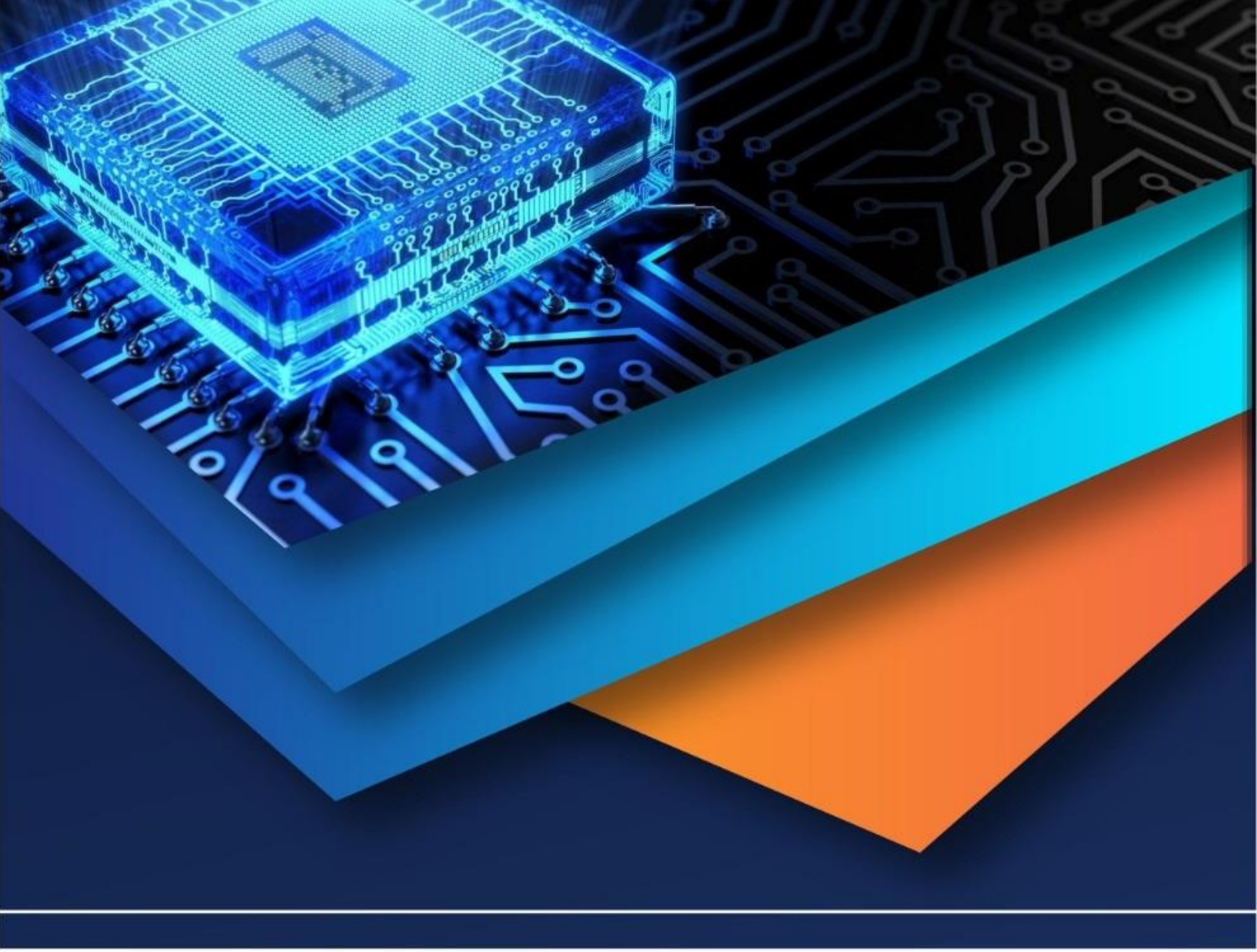
REFERENCES

[1] Aladejana, E. B. (2023). Biological properties of polyherbal formulations: A review of their antimicrobial, anti-inflammatory, antioxidant, and toxicological activities. Pharmacognosy Journal, 15(5), 933–963.
[2] Abbasi, F., & Rousu, J. (2025). New methods for drug synergy prediction: A mini-review. Trends in Pharmacological Sciences.
[3] Current perspectives and trend of computer-aided drug design: A review and bibliometric analysis. (2024). PMC. https://pmc.ncbi.nlm.nih.gov/articles/PMC11175770/
[4] Computer-Aided Drug Design and Drug Discovery. (2023). PMC NIH. https://pmc.ncbi.nlm.nih.gov/articles/PMC10819513/
[5] Computer Aided Drug Design Methodologies with Natural Products. (2022). PMC NIH. https://pmc.ncbi.nlm.nih.gov/articles/PMC9881095/
[6] Computer-Aided Drug Design Methods. (2016). PMC NIH. https://pmc.ncbi.nlm.nih.gov/articles/PMC5248982/
[7] Madhuri, L. G., Vaishnavi, P. A., Anekar, M., & Patil, M. S. (2023). Computer aided drug design: An overview. International Journal of Pharmaceutical and Preventive Research, 14(2), 69–78.
[8] Naik, Y., & Goudanavar, S. (2025). Quantum-inspired computational drug design for phytopharmaceuticals: A herbal holography analysis. Journal of Molecular Modeling, 31(7), 188. https://doi.org/10.1007/s00894-025-06412-w
[9] Navami, U., Divya Jyothi, Nireeeksha Shetty, & Nisha N Shetty. (2024). Design and optimization of polyherbal gels for topical drug delivery by central composite statistical design. Research Journal of Pharmacy and Technology, 17(12), 5702–5706. https://doi.org/10.52711/0974-360X.2024.00868
[10] Saini, J., & Parashar, M. (2025). Development and evaluation of polyherbal dispersible tablets using aqueous leaf extracts. Journal of Pharmaceutical Research International, 37(4), 79–86. https://doi.org/10.9734/jpri/2025/v37i47680
[11] Zhang, X., Liu, Y., & Wang, H. (2024). Integrative network pharmacology, molecular docking, and dynamic simulation analysis of a polyherbal formulation for potential therapeutic impact on prostate cancer. Heliyon, 10(14), e34531. https://doi.org/10.1016/j.heliyon.2024.e34531
[12] Polyherbal formulation concept for synergic action: A review. (2019). Journal of Drug Delivery and Therapeutics, 9(1-s), 123–130. https://doi.org/10.22270/jddt.v9i1-s.2339
[13] Synergistic effect of polyherbal extract containing indigenous medicinal plants. (2025). Applied Journal of Alternative and Integrative Medicine.
[14] Synergistic effects of phytochemicals in polyherbal formulations. (2025). International Journal of Pharmaceutical Sciences Research.
[15] Drug synergy prediction – TDC. (2024). Therapeutics Data Commons. https://tdcommons.ai/multi_pred_tasks/drugsyn/
[16] Computational methodology for drug discovery. (2024). Nature Collections. https://www.nature.com/collections/dhcgiaigac
[17] AI-driven drug discovery: A comprehensive review. (2025). ACS Omega. https://doi.org/10.1021/acsomega.5c00549
[18] AI in ethnopharmacology, the pharmaceutical industry, and its applications. (2025). Annales Pharmaceutiques Françaises. https://doi.org/10.1016/j.pharma.2025.11.011
[19] Artificial intelligence (AI) applications in drug discovery and development. (2024). PMC NIH. https://pmc.ncbi.nlm.nih.gov/articles/PMC11510778/
[20] Computational frameworks transform antagonism to synergy in optimizing drug combinations. (2025). Nature Digital Medicine. https://doi.org/10.1038/s41746-025-01435-2
[21] MD-Syn: Synergistic drug combination prediction based on a multidimensional feature fusion method and attention mechanisms. (2025). Frontiers in Pharmacology. https://doi.org/10.3389/fphar.2025.1564339
[22] A systematic evaluation of deep learning methods for the prediction of drug synergy. (2023). PLoS Computational Biology, 19(3), e1010200. https://doi.org/10.1371/journal.pcbi.1010200
[23] Systematic review of computational methods for drug combination prediction. (2022). Computational and Structural Biotechnology Journal, 20, 2807–2814. https://doi.org/10.1016/j.csbj.2022.05.055
[24] Network pharmacology and molecular docking approaches for herbal formulations: Mechanistic insights, therapeutic applications, and translational perspectives. (2026). International Journal of Pharma and Bio Sciences, 17(1), 24–33. https://doi.org/10.22376/ijpbs.v17i1.142
[25] Quantum computing in molecular design and drug discovery. (2026). Journal of ISI, 8(1). https://doi.org/10.63158/journalisi.v8i1.1380

©IJRASET: All Rights are Reserved | SJ Impact Factor 7.538 | ISRA Journal Impact Factor 7.894 |

[26] AI-enabled computational strategies in herbal drug discovery. (2026). International Journal of Ayurveda, Herbal Medicine and Integrative Medicine.
[27] Advances in drug design: A review of recent trends, challenges and future scope. (2025). Quarterly Journal of Medicine and Health Sciences.
[28] A review on drug design: Strategies, advances, and future directions. (2025). International Journal of Applied Science and Engineering Technology.
[29] Drug design—Past, present, future. (2022). PMC NIH. https://pmc.ncbi.nlm.nih.gov/articles/PMC8911833/
[30] Research in the field of drug design and development. (2023). PMC NIH. https://pmc.ncbi.nlm.nih.gov/articles/PMC10536713/

©IJRASET: All Rights are Reserved | SJ Impact Factor 7.538 | ISRA Journal Impact Factor 7.894 |



10.22214/IJRASET

45.98

IMPACT FACTOR: 7.129

IMPACT FACTOR: 7.429

INTERNATIONAL JOURNAL FOR RESEARCH

IN APPLIED SCIENCE & ENGINEERING TECHNOLOGY

Call : 08813907089 (24*7 Support on Whatsapp)