

Development of an end-to-end Quantitative Model-Informed Drug Development (MIDD) ECOSYSTEM

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Background

Model-Informed Drug Development (MIDD) is essential in pharmaceutical R&D. Growing therapeutic complexity and regulatory expectations for model qualification **necessitate a structured, automated, and scalable MIDD ecosystem**.

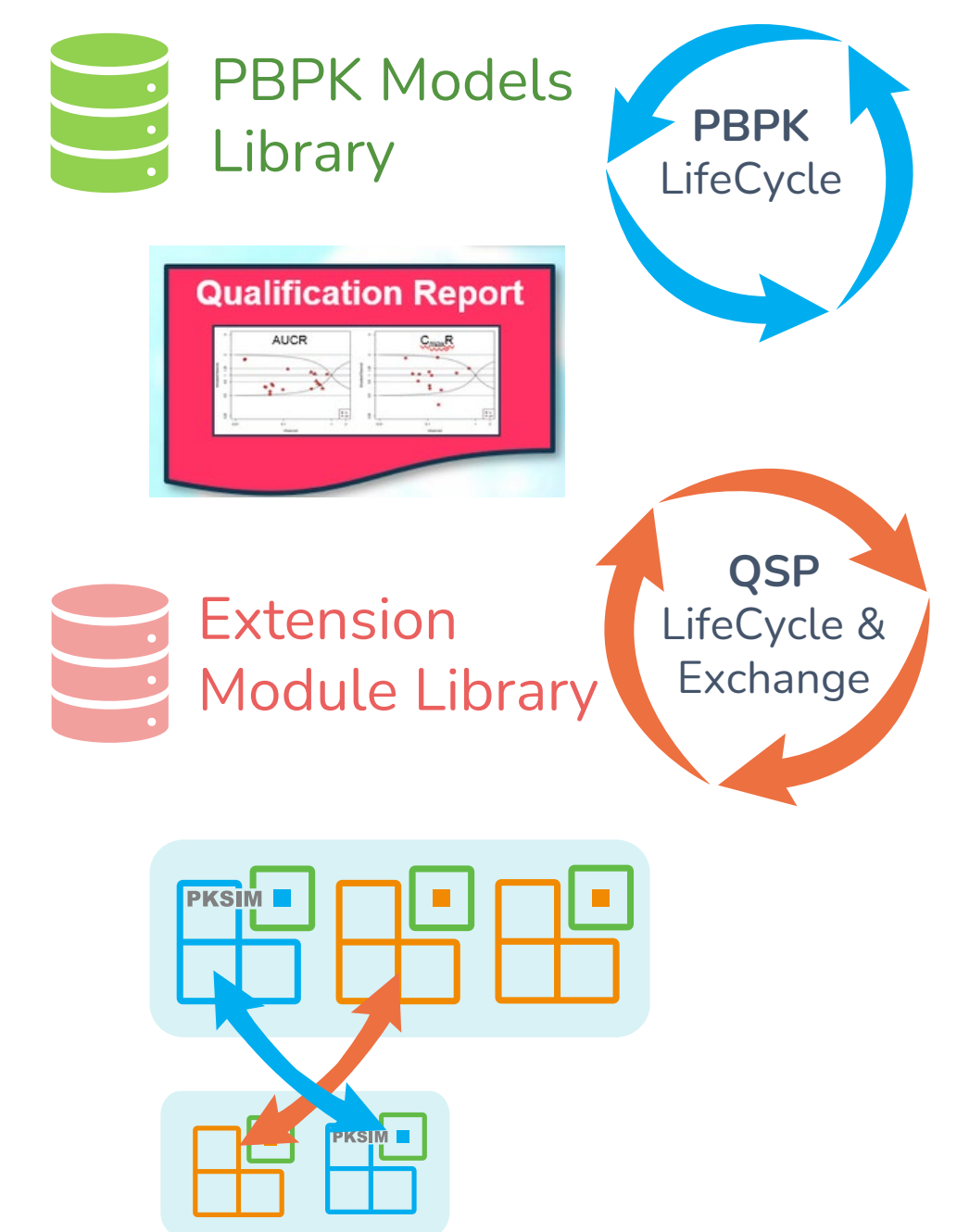
The Open Systems Pharmacology (OSP) Suite [1] Version 12 introduces **modularization capabilities** enabling such an integrated ecosystem. This supports seamless **development, validation, and deployment** of Physiologically Based Pharmacokinetic (PBPK) and Quantitative Systems Pharmacology/Toxicology (QSP/T) models, ensuring **robust, reproducible decision-making**.

It was initially conceived to enhance PBPK-QSP platform management for the Diabetes Disease Platform [2, 3].

Results

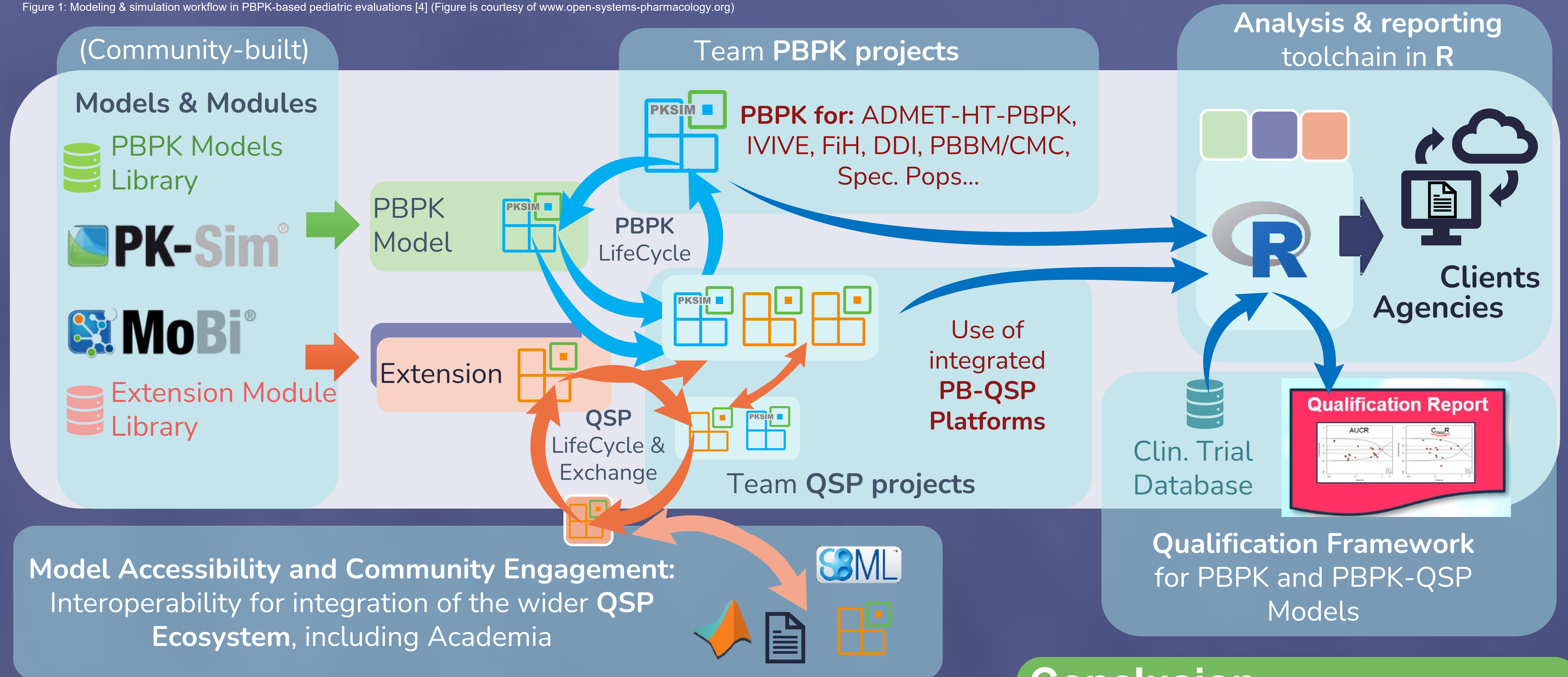
The **end-to-end MIDD ecosystem**, leveraging OSP Suite V12, implements structured **model storage, validation, modularization, and continuous integration**. Key components include:

- A continuously expanding **PK-Sim model library** of qualified PBPK models. This centralized repository provides **verified models** for various drugs, populations (e.g., pediatric, disease-states), and mechanistic scenarios, **reducing duplication and standardizing methodologies**. Automated model **qualification** ensures reliability for applications like drug-drug interaction (DDI) assessments and special population extrapolations.
- A **MoBi module library** of community-driven QSP/T models, encapsulating disease biology and drug mechanisms. It offers **prebuilt, validated modules for disease progression and pharmacodynamic effects**, enabling pathway-based **PBPK-QSP integration**. This fosters **open-source collaboration, model sharing, and version control**, efficiency.
- **Modular interchangeability** of PBPK models within QSP platforms, a key OSP Suite V12 innovation, enabling Continuous Integration/Continuous Deployment (CI/CD). This **allows dynamic PBPK model replacement in QSP frameworks and automated PBPK-QSP platform qualification** against predefined criteria, ensuring models remain current and continuously validated for regulatory compliance.



This integrated approach, bridging PBPK and QSP/T modeling, has been successfully implemented across diverse pipelines, including **in-vitro systems, high-throughput PBPK, custom modules (e.g., ocular absorption, pregnancy), and integrated disease/drug effect models across safety endpoints and therapeutic areas**.

Figure 1: Modeling & simulation workflow in PBPK-based pediatric evaluations [4] (Figure is courtesy of www.open-systems-pharmacology.org)



Conclusion

The **end-to-end MIDD ecosystem**, powered by OSP Suite V12, represents a significant **advancement in PBPK-QSP modeling**, effectively linking model development, qualification, and regulatory delivery. Its core strengths—continuously evolving PK-Sim and MoBi libraries for reusable, validated models, coupled with modular interchangeability and CI/CD for agile, automated PBPK-QSP platform qualification—drive this transformation. By implementing automated qualification, standardized repositories, and modular workflows integrated with OSP-R and ESQlabs-R packages, the **ecosystem markedly enhances transparency, reproducibility, and regulatory credibility**. This ultimately empowers regulatory engagement with robust, continuously validated models, accelerating drug development, reducing predictive uncertainty, and supporting evidence-based approvals. The **synergy of open-source software, modularity, and automation is pivotal for maximizing MIDD's impact in pharmaceutical R&D**.

References

[1] Lippert J, et al. Open Systems Pharmacology Community-An Open Access, Open Source, Open Science Approach to Modeling and Simulation in Pharmaceutical Sciences. CPT Pharmacometrics Syst Pharmacol. 2019; Dec;8(12):878-882. doi: 10.1002/psp4.12473.

[2] Schaller, S., et al., A Generic Integrated Physiologically based Whole-body Model of the Glucose-Insulin-Glucagon Regulatory System. CPT Pharmacometrics Syst Pharmacol. 2013, 2: 1-10 65. https://doi.org/10.1038/psp.2013.40
 [3] Balazki, P., Schaller, S. et al., A Quantitative Systems Pharmacology Kidney Model of Diabetes Associated Renal Hyperfiltration and the Effects of SGLT Inhibitors. CPT Pharmacometrics Syst Pharmacol. 2018, 7: 788-797. https://doi.org/10.1002/psp4.12359



Supporting the open-source development of:
PK-Sim **MoBi** **OSP** OPEN SYSTEMS PHARMACOLOGY Software Suite
 www.Open-Systems-Pharmacology.org