# Technical Report: Biofilter System Status Update

## Summary

We initially began development on **Biofilter/LOKI version 3.0.0,** but work was paused upon identifying critical issues with **version 2.4.3** that required resolution before progressing with the new version. Our focus shifted to addressing these issues in version 2.4.3, prioritizing stability and user needs. The following report outlines the actions taken, current progress, challenges, and next steps.

Create the Issue Tests Area and answer the Issues:

* Rasika
* ..

Install the Sphinx in the project to control the documentations

Install and set the Poetry, add a new folder dist with the install package (pip install biofilter)

Instalado o tox.

Criado o branch gh-pages e setting o pages para direcionar para a documentacao tecnica do projeto.   
- configurado o Actions para preparacao do ambiente.

### Key Actions Taken

**1. Refocusing Development on Version 2.4.3**

* The team was instructed to identify and document all problems in version 2.4.3 in the GitHub Issues section.
* Recovery of the 2.4.3 codebase was necessary as the repository had a single branch, Master, containing version 3.0.0 code.
* A new branch structure was created to organize and clarify the development:

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| A screenshot of a computer  Description automatically generated | * **Main**: Contains the restored 2.4.3 code and is designate d as the future default branch. * **Development** (tagged 2.4.4): Branch for active work on version 2.4.3 improvements. * **dev-3.0.0**: Copy of the Master branch to isolate version 3.0.0 development. * Upon team approval, the current default branch (Master) will be removed, and Main will be set as the default to ensure users access the stable version (2.4.3). |

**2. Setup Improvements**

* Adjusted *`setup.py`* to include dependencies, allowing installation via `*pip install`.* This simplified execution by enabling global use of `*biofilter*` and `*loki`* commands.
* Plan to transition from `*setup.py`* to Poetry (https://python-poetry.org/) for better dependency management and a more transparent development process.

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**3. Documentation Enhancements**

* Created a dedicated technical documentation area for developers, providing a quick onboarding guide.
* Added a *README.md* to the **Development** branch to explain its purpose and updates.
* Installed and configured tools like **Black**, **Coverage**, and **Pytest**. Development dependencies were excluded from *setup*.*py* but listed in a *requirements-dev.txt.*
* Plans to incorporate **Sphinx** (https://www.sphinx-doc.org/en/master/) for structured documentation in the coming weeks.

**4. Code Restructuring**

* Separated functionalities into distinct files:
  + Entry point file
  + Argument configuration file
  + Class and methods file with **mixins** for better organization
* Added detailed comments to explain methods, constants, and schemas used for generating temporary databases.
* Maintained current functionality while improving code clarity and maintainability.

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**5. Integration of LOKI**

* Currently, the Biofilter system requires LOKI files to be embedded due to its packaging model.
* Kept this structure to avoid disrupting Biofilter’s functionality but recommend reviewing this approach in future updates to separate LOKI as an independent, installable package.

### Current Development Focus

**1. Unit Tests**

* Developing tests to validate Biofilter methods' functionality (not LOKI-specific data).
* Using **Coverage** to track statement-level testing, currently at **55% coverage with 220 tests**, including LOKI's embedded code (loki\_db.py only).
* Implementing a test-driven development (TDD) approach to ensure integrity before modifying any code.

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In the past few weeks, we have added nearly 10,000 lines of code to the Biofilter project, primarily in the form of detailed comments on methods and unit tests. These additions aim to enhance the project’s manageability and readability, significantly reducing the onboarding time for future collaborators. By providing clear documentation and robust testing frameworks, we ensure that new team members can quickly understand the system’s structure and functionality, fostering smoother project development and maintenance.



The table highlights the progress in modularizing the code, increasing test coverage, and reorganizing functionalities, facilitating the project's maintenance and scalability.

A screenshot of a computer program

Description automatically generated**2. Functional and Issues Tests**

* Addressed two GitHub issues as functional test cases:
* Issue 1: Biofilter Group Annotation <https://github.com/RitchieLab/biofilter/issues/15>
* Issue 2: build 37 LOKI

<https://github.com/RitchieLab/LOKI/issues/16>

* These tests are located under a separate directory (`tests/ISSUE`) for isolation and can be discarded once the problems are fully resolved.
* Collaborating with the team to identify essential functional tests for key features like annotations, filters, models, installation, and LOKI integration.

### Challenges

**1. Codebase Complexity**

* The original Biofilter code was a single file with no comments, making it difficult to debug and enhance.
* While restructuring improved maintainability, further work is needed to ensure long-term project integrity.

**2. Embedded LOKI Files**

* LOKI files are hardcoded into the Biofilter system, limiting modularity and scalability. Separating LOKI into an independent package would resolve this issue.

### Next Steps

**1. Finalize Version 2.4.4**

* Complete unit and functional tests to validate stability.
* Resolve remaining GitHub issues reported by the team.
* Prepare for release using the LOKI database on the LPC environment.

**2. Improve Documentation**

* Implement Sphinx for developer and user documentation.
* Link functional tests to documentation as examples for users.

**3. Plan for Version 3.0.0**

* Review identified improvements for the LOKI database and Biofilter structure.
* Transition to modern development practices, including the use of ORM and modular package management.

This comprehensive approach ensures immediate issues with version 2.4.3 are resolved while laying the foundation for a robust version 3.0.0.

Log do tox:  
biofilter on  development [✘!?] via 🐍 3.11.9 using ☁️ default/terra-7de85c67 via biofilter

➜ poetry run tox

py310: install\_deps> python -I -m pip install poetry

.pkg: \_optional\_hooks> python /Users/andrerico/Works/Sys/biofilter/venv/lib/python3.11/site-packages/pyproject\_api/\_backend.py True poetry.core.masonry.api

.pkg: get\_requires\_for\_build\_sdist> python /Users/andrerico/Works/Sys/biofilter/venv/lib/python3.11/site-packages/pyproject\_api/\_backend.py True poetry.core.masonry.api

.pkg: get\_requires\_for\_build\_wheel> python /Users/andrerico/Works/Sys/biofilter/venv/lib/python3.11/site-packages/pyproject\_api/\_backend.py True poetry.core.masonry.api

.pkg: prepare\_metadata\_for\_build\_wheel> python /Users/andrerico/Works/Sys/biofilter/venv/lib/python3.11/site-packages/pyproject\_api/\_backend.py True poetry.core.masonry.api

.pkg: build\_sdist> python /Users/andrerico/Works/Sys/biofilter/venv/lib/python3.11/site-packages/pyproject\_api/\_backend.py True poetry.core.masonry.api

py310: install\_package\_deps> python -I -m pip install apsw==3.46.1.0 click==8.1.7 iniconfig==2.0.0 packaging==24.1 platformdirs==4.3.6 pluggy==1.5.0

py310: install\_package> python -I -m pip install --force-reinstall --no-deps /Users/andrerico/Works/Sys/biofilter/.tox/.tmp/package/3/biofilter-2.4.4.tar.gz

py310: commands[0]> poetry install --no-root

Installing dependencies from lock file

Package operations: 29 installs, 0 updates, 0 removals

- Installing markupsafe (3.0.2)

- Installing alabaster (1.0.0)

- Installing babel (2.16.0)

- Installing docutils (0.21.2)

- Installing imagesize (1.4.1)

- Installing jinja2 (3.1.4)

- Installing pygments (2.18.0)

- Installing snowballstemmer (2.2.0)

- Installing sphinxcontrib-applehelp (2.0.0)

- Installing sphinxcontrib-devhelp (2.0.0)

- Installing sphinxcontrib-htmlhelp (2.1.0)

- Installing sphinxcontrib-jsmath (1.0.1)

- Installing sphinxcontrib-qthelp (2.0.0)

- Installing sphinxcontrib-serializinghtml (2.0.0)

- Installing sphinx (8.1.3)

- Installing cachetools (5.5.0)

- Installing chardet (5.2.0)

- Installing colorama (0.4.6)

- Installing exceptiongroup (1.2.2)

- Installing mypy-extensions (1.0.0)

- Installing pathspec (0.12.1)

- Installing pyproject-api (1.8.0)

- Installing sphinxcontrib-jquery (4.1)

- Installing typing-extensions (4.12.2)

- Installing black (24.10.0)

- Installing coverage (7.6.8)

- Installing pytest (8.3.4)

- Installing sphinx-rtd-theme (3.0.2)

- Installing tox (4.23.2)

py310: commands[1]> pytest tests/

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============================================================ warnings summary ============================================================

tests/units/test\_logger\_mixin.py:8

/Users/andrerico/Works/Sys/biofilter/tests/units/test\_logger\_mixin.py:8: PytestCollectionWarning: cannot collect test class 'TestLoggerMixin' because it has a \_\_init\_\_ constructor (from: tests/units/test\_logger\_mixin.py)

class TestLoggerMixin(LoggerMixin):

tests/units/test\_user\_knoeledge\_retrievel\_mixin.py:29

/Users/andrerico/Works/Sys/biofilter/tests/units/test\_user\_knoeledge\_retrievel\_mixin.py:29: PytestCollectionWarning: cannot collect test class 'TestUserKnowledgeRetrievalMixin' because it has a \_\_init\_\_ constructor (from: tests/units/test\_user\_knoeledge\_retrievel\_mixin.py)

class TestUserKnowledgeRetrievalMixin(UserKnowledgeRetrievalMixin):

tests/units/test\_user\_knowledge\_input\_mixin.py:42

/Users/andrerico/Works/Sys/biofilter/tests/units/test\_user\_knowledge\_input\_mixin.py:42: PytestCollectionWarning: cannot collect test class 'TestUserKnowledgeInputMixin' because it has a \_\_init\_\_ constructor (from: tests/units/test\_user\_knowledge\_input\_mixin.py)

class TestUserKnowledgeInputMixin(UserKnowledgeInputMixin):

-- Docs: https://docs.pytest.org/en/stable/how-to/capture-warnings.html

213 passed, 3 warnings in 5.95s

py310: OK ✔ in 30.9 seconds

py311: install\_package> python -I -m pip install --force-reinstall --no-deps /Users/andrerico/Works/Sys/biofilter/.tox/.tmp/package/4/biofilter-2.4.4.tar.gz

py311: commands[0]> poetry install --no-root

Installing dependencies from lock file

No dependencies to install or update

py311: commands[1]> pytest tests/

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class TestUserKnowledgeRetrievalMixin(UserKnowledgeRetrievalMixin):

tests/units/test\_user\_knowledge\_input\_mixin.py:42

/Users/andrerico/Works/Sys/biofilter/tests/units/test\_user\_knowledge\_input\_mixin.py:42: PytestCollectionWarning: cannot collect test class 'TestUserKnowledgeInputMixin' because it has a \_\_init\_\_ constructor (from: tests/units/test\_user\_knowledge\_input\_mixin.py)

class TestUserKnowledgeInputMixin(UserKnowledgeInputMixin):

-- Docs: https://docs.pytest.org/en/stable/how-to/capture-warnings.html

213 passed, 3 warnings in 4.59s

py311: OK ✔ in 10.66 seconds

py312: install\_deps> python -I -m pip install poetry

py312: install\_package\_deps> python -I -m pip install apsw==3.46.1.0 click==8.1.7 iniconfig==2.0.0 packaging==24.1 platformdirs==4.3.6 pluggy==1.5.0

py312: install\_package> python -I -m pip install --force-reinstall --no-deps /Users/andrerico/Works/Sys/biofilter/.tox/.tmp/package/5/biofilter-2.4.4.tar.gz

py312: commands[0]> poetry install --no-root

Installing dependencies from lock file

Package operations: 27 installs, 0 updates, 0 removals

- Installing markupsafe (3.0.2)

- Installing alabaster (1.0.0)

- Installing babel (2.16.0)

- Installing docutils (0.21.2)

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- Installing black (24.10.0)

- Installing coverage (7.6.8)

- Installing pytest (8.3.4)

- Installing sphinx-rtd-theme (3.0.2)

- Installing tox (4.23.2)

py312: commands[1]> pytest tests/

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class TestUserKnowledgeInputMixin(UserKnowledgeInputMixin):

tests/units/test\_user\_knowledge\_input\_mixin.py::test\_add\_user\_group\_biopolymers

tests/units/test\_user\_knowledge\_input\_mixin.py::test\_add\_user\_group\_biopolymers

tests/units/test\_user\_knowledge\_input\_mixin.py::test\_add\_user\_group\_biopolymers\_tally

tests/units/test\_user\_knowledge\_input\_mixin.py::test\_add\_user\_group\_biopolymers\_tally

/Users/andrerico/Works/Sys/biofilter/biofilter\_modules/mixins/user\_knowledge\_input\_mixin.py:165: DeprecationWarning: Binding 4 ('?4') is a named parameter, but you supplied a sequence which requires nameless (qmark) placeholders. Starting with Python 3.14 an sqlite3.ProgrammingError will be raised.

cursor.executemany(

-- Docs: https://docs.pytest.org/en/stable/how-to/capture-warnings.html

213 passed, 7 warnings in 5.11s

.pkg: \_exit> python /Users/andrerico/Works/Sys/biofilter/venv/lib/python3.11/site-packages/pyproject\_api/\_backend.py True poetry.core.masonry.api

py310: OK (30.90=setup[19.12]+cmd[5.25,6.53] seconds)

py311: OK (10.66=setup[5.02]+cmd[0.87,4.77] seconds)

py312: OK (27.21=setup[16.91]+cmd[4.40,5.89] seconds)

congratulations :) (68.88 seconds)