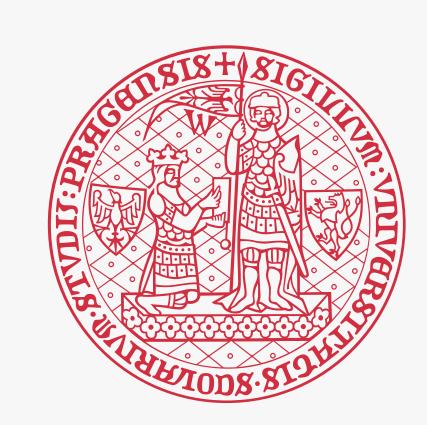


Configurable point rasterization for large scatterplots



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Thesis goals

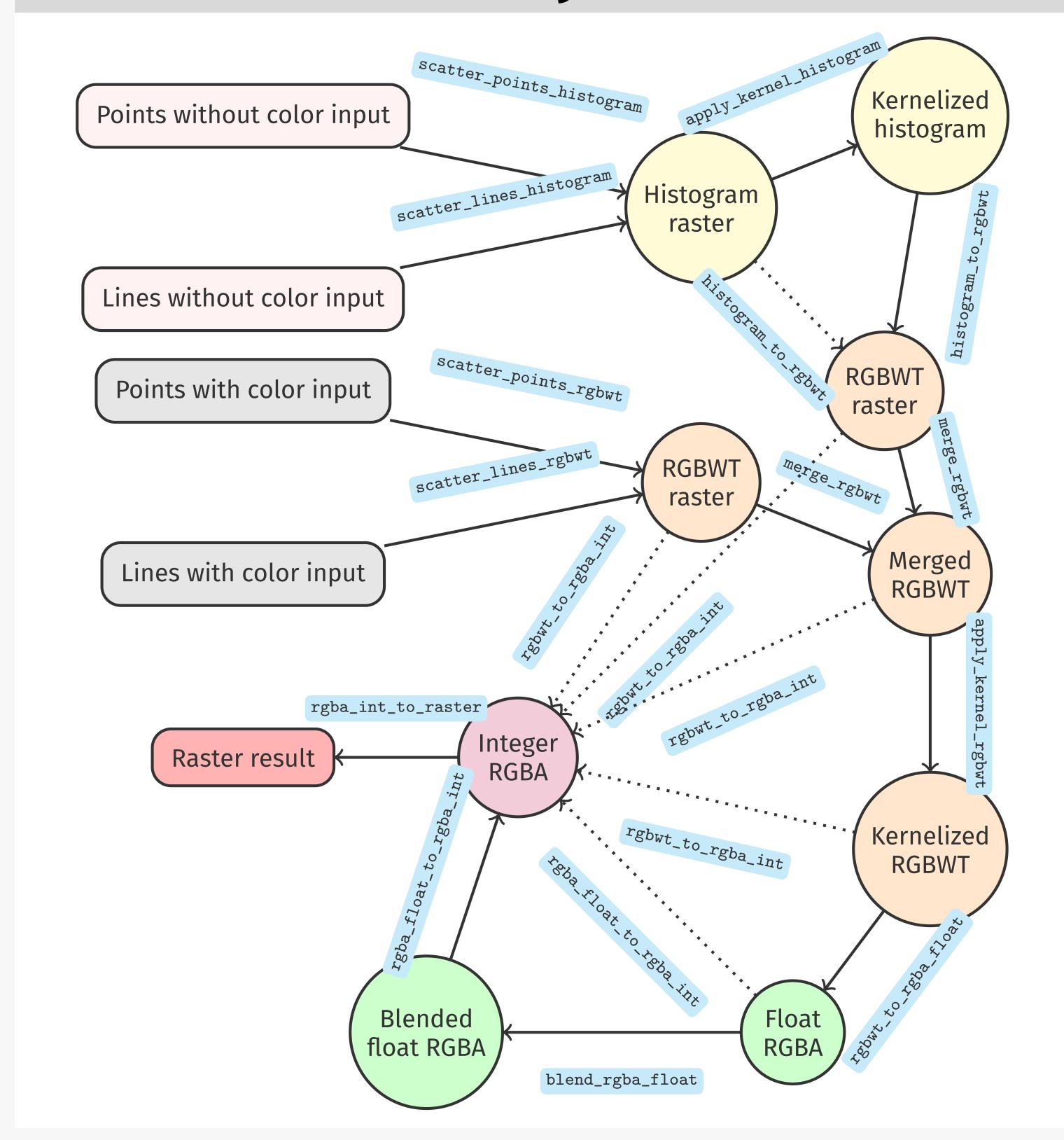
Scattermore is a simple R package used for scatterplot visualizations. The goal of the thesis is to expand scattermore to support new plotting functionality while retaining the performance.

- **High-customizable API.** We assign each subtask to a single method to ensure more variation in output for the user.
- Reduction of overplotting. We introduce the new RGBWT data format, which combines images equally.
- Fast plotting. We implement the computationally expensive operation in C language.

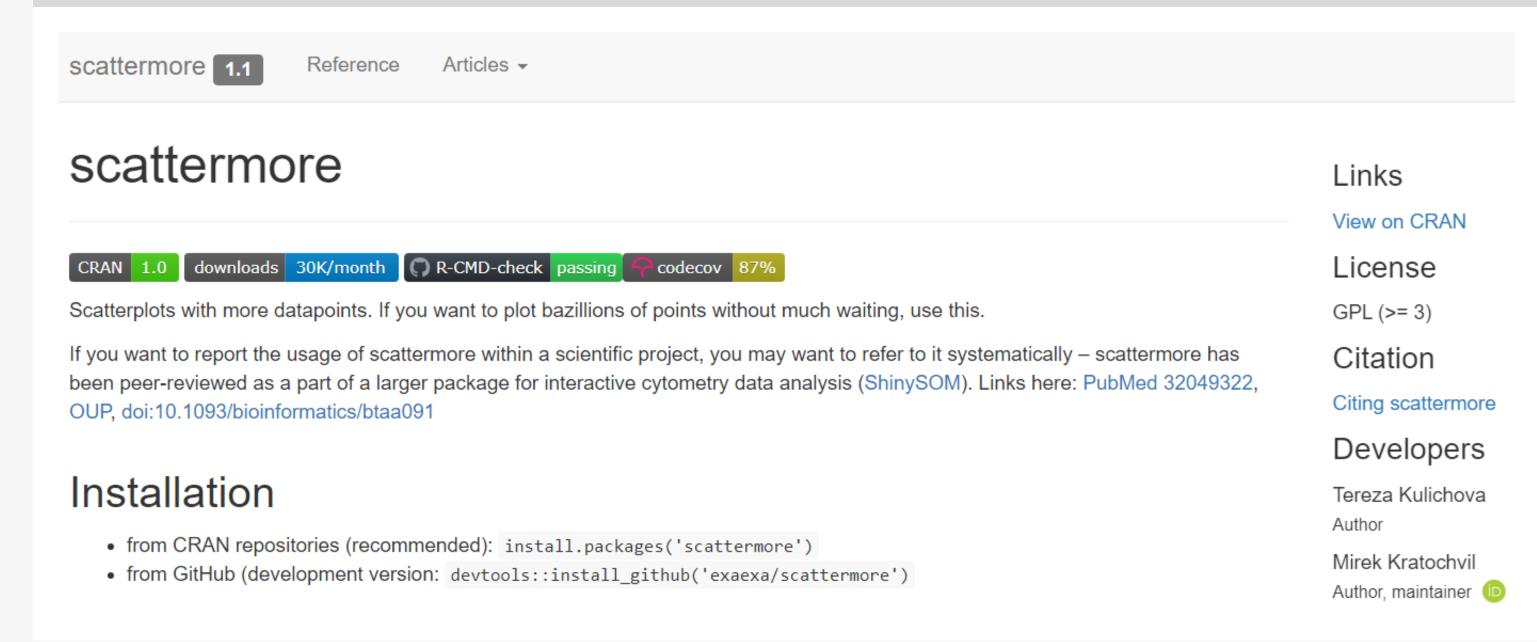
Approach

- Design a specific pixel format that supports orderindependent alpha-blending.
- Try some acceleration possibilities (SIMD, multithreading).
- Package and document everything properly to make scattermore easier to use.

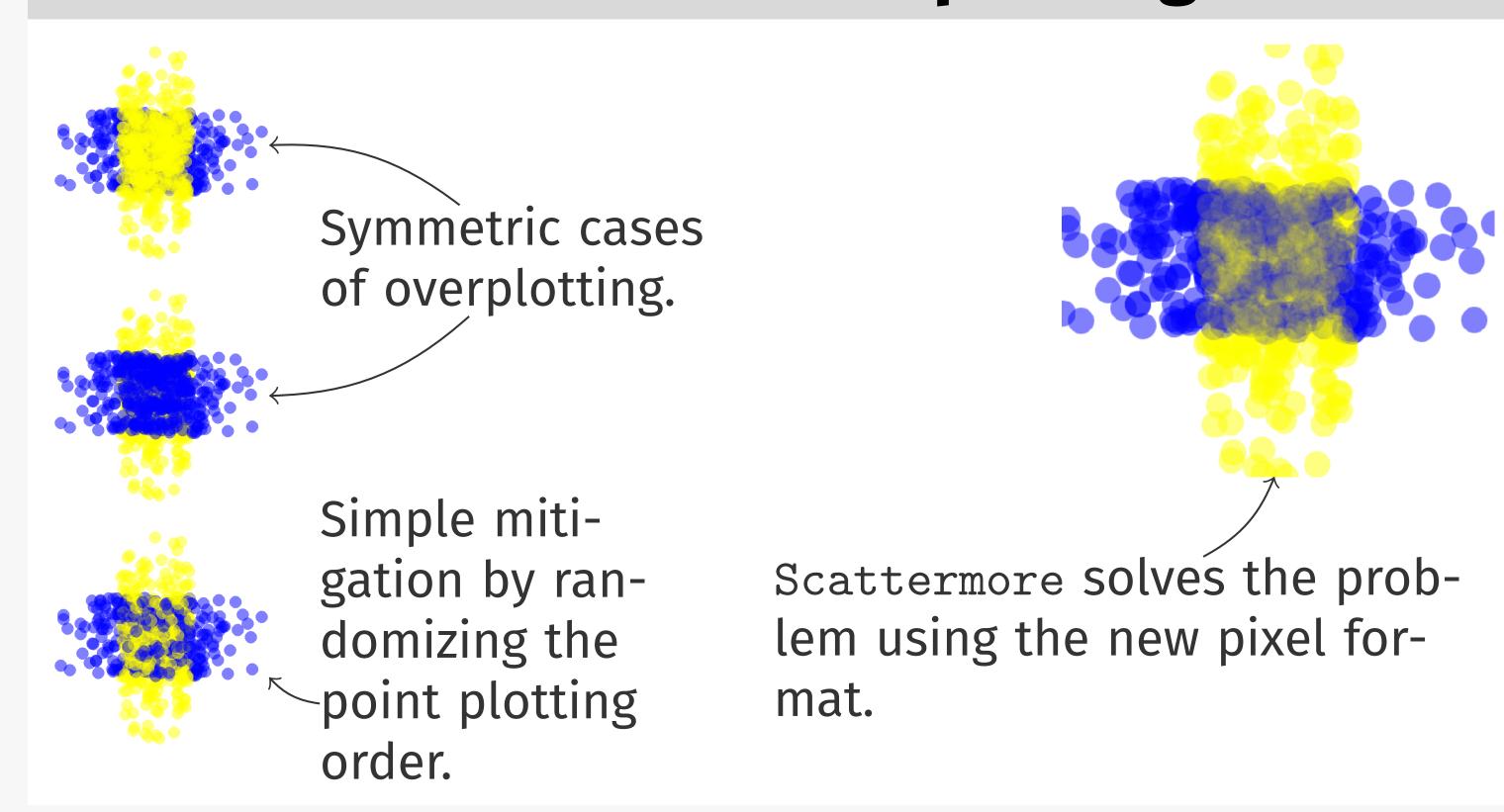
New API & functionality overview



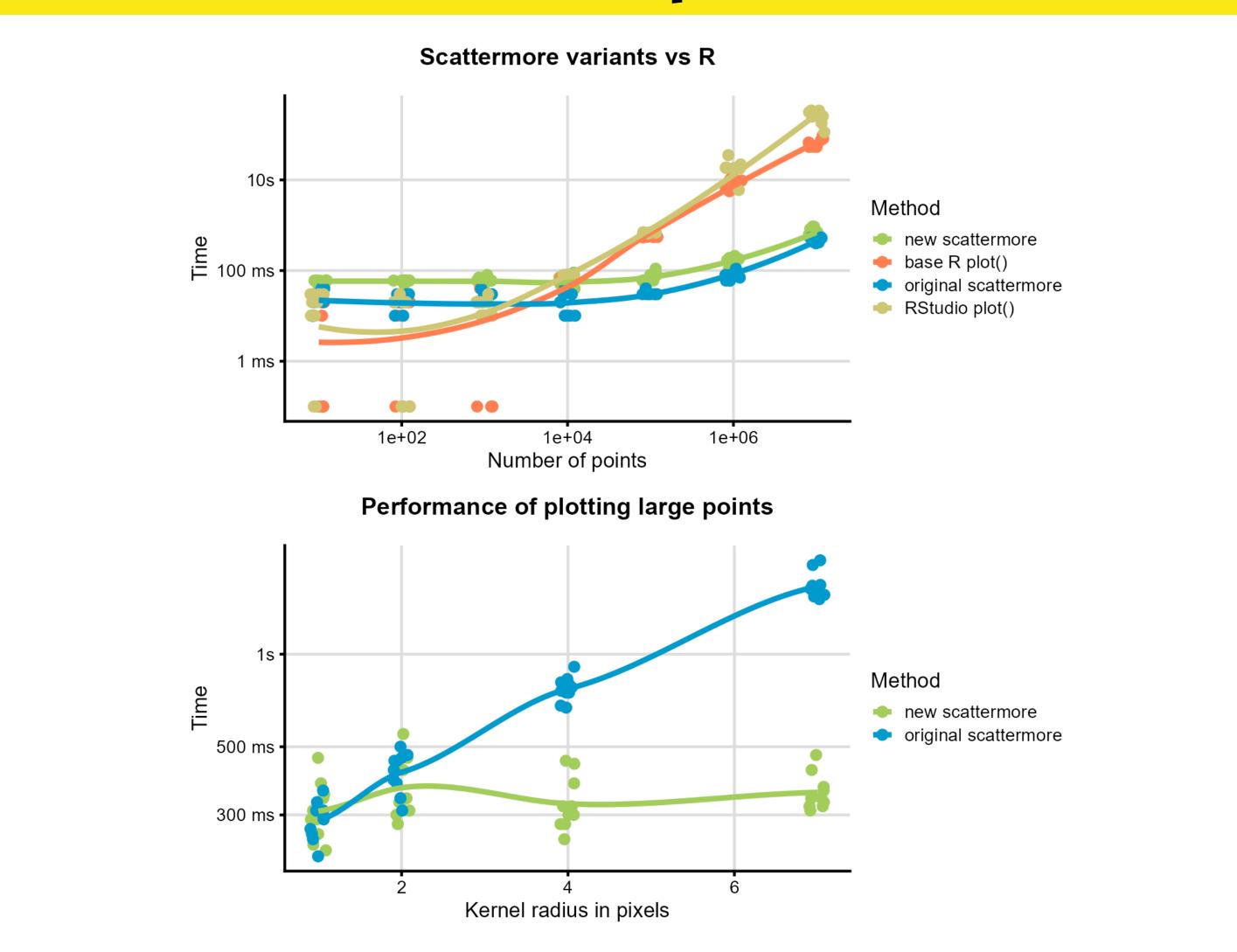
Deployed user-friendly documentation



Result: RGBWT avoids overplotting artifacts



Result: Performance improvements



Result: Complex plotting



Combined plotting of more complicated primitives utilizing functionalities in the redesigned API.

Conclusions

- Scattermore enables users to blur images, blend or merge them according to their preferences and, at the same time, sufficiently fast.
- It is especially useful in the cytometric community for interactive analysis of large biological data.
- Scattermore is available on GitHub as a development version and as R's official CRAN package version.

