

Kvik: Interactive exploration of multi-omics data from the NOWAC postgenome biobank

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Kvik

Integrates state of the art pathway maps from the KEGG database and gene expression data from the NOWAC postgenome biobank in a single system

Kvik is a web app, so you can access it anywhere

Kvik

Does not require any installation or plug-ins. Runs directly in the web browser.

Uses the expert drawn pathway images from **KEGG**.

Has an extendible backend implemented in **R**, managing datasets on its own

Demo



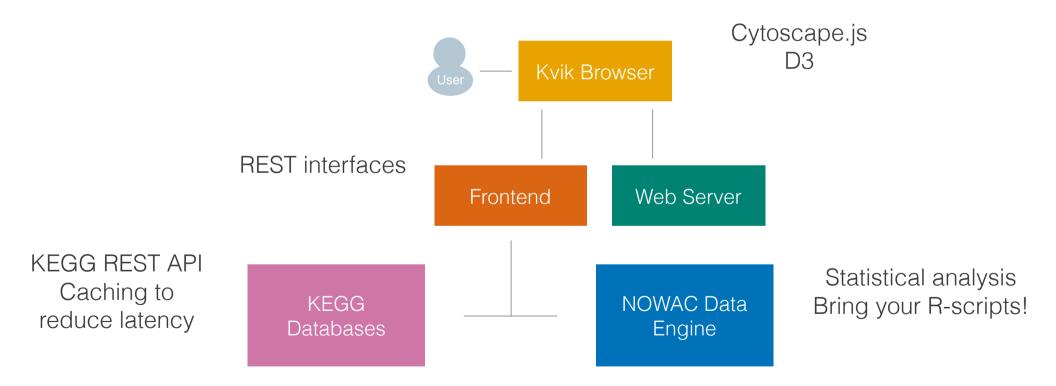
Welcome!

Kvik is an interactive system for exploring the dynamics of carcinogenesis through integrated studies of biological pathways and genomic data. It provides researchers with a lightweight web application for navigating through biological pathways from the KEGG database integrated with genomic data from the Norwegian Women and Cancer (NOWAC) postgenome biobank.

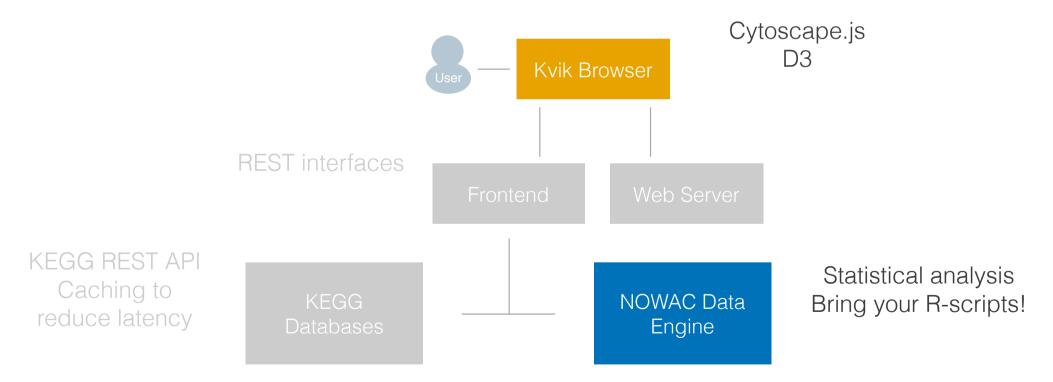
To explore pathways and genomic data, click the start button below.

Start!

System Architecture



System Architecture



Future Work

More advanced statistical analyses through statistical packages like Bioconductor

Integrate **new data sources** and **more data** from the NOWAC biobank.

Making it accessible to you guys!

The Tromsø Large Display Wall

A 22 megapixel backprojected display composed of 28 tiled projectors. Projectors driven by a 28-node display cluster.



Concluding Remarks

Kvik visualizes biological pathways from KEGG and gene expression data from the NOWAC biobank

Open-sourced at github.com/fjukstad/kvik



Thank you!

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Kvik — Lead sled dog on Fridtjof Nansen's Fram expedition

