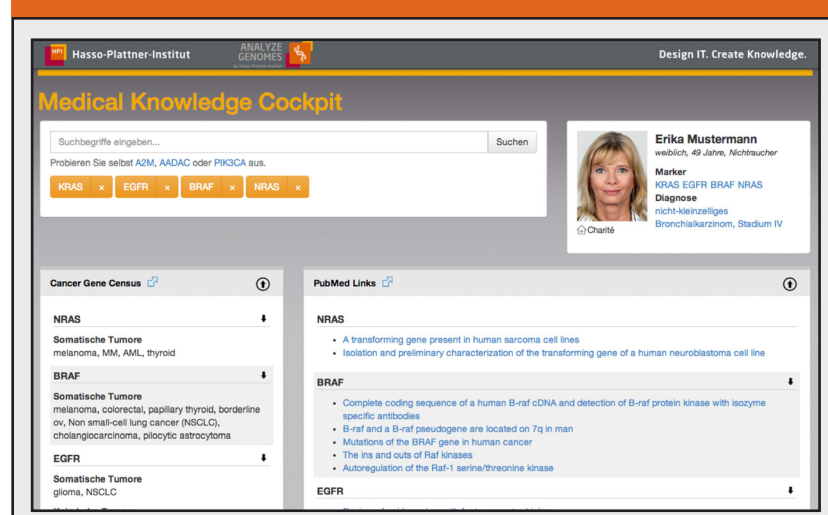


IN-MEMORY TECHNOLOGY ENABLING REAL-TIME ANALYSIS OF BIG MEDICAL DATA

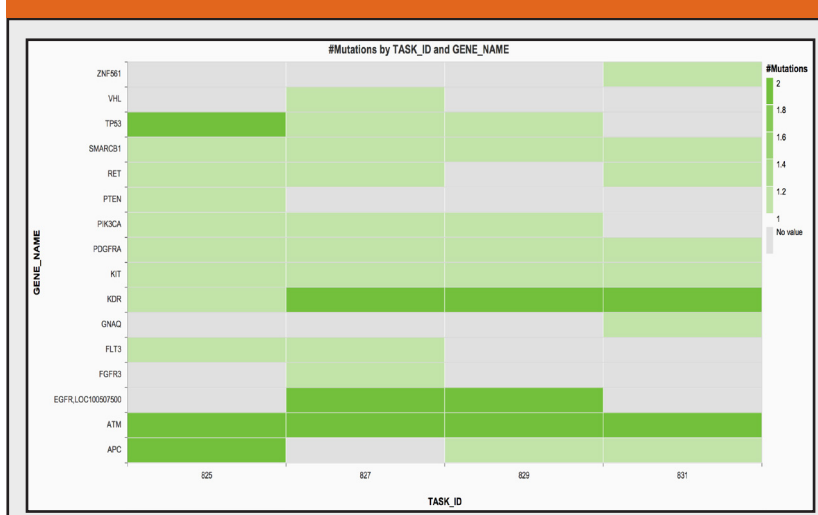
ANALYTICS

MEDICAL KNOWLEDGE COCKPIT



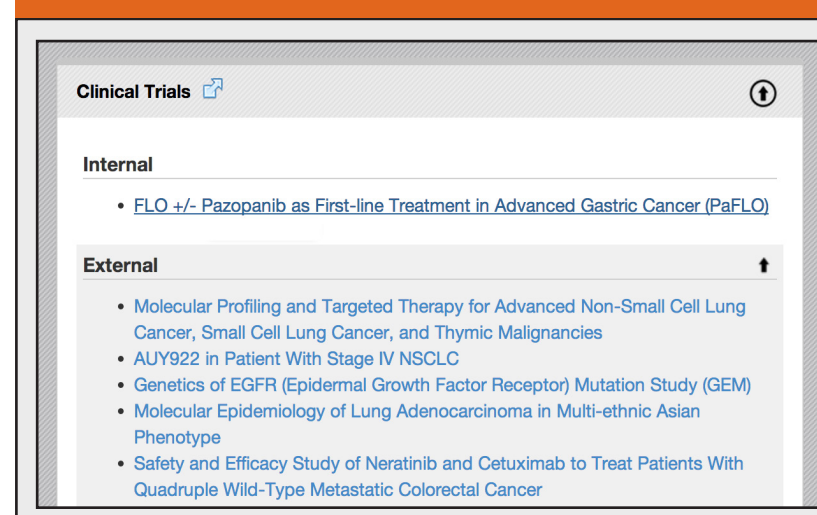
Combination of international research results to enable personalized treatment

DRUG RESPONSE ANALYSIS



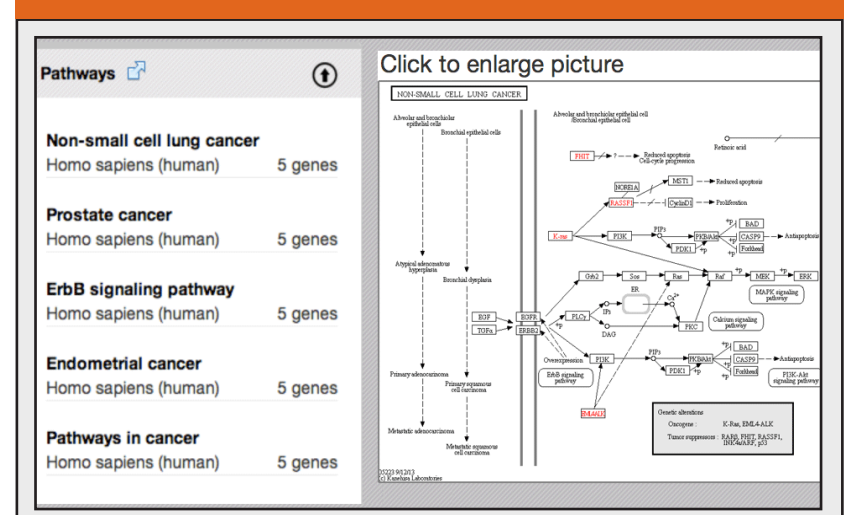
Prediction of drug response based on real-time analysis of similar tumors

IDENTIFICATION OF CLINICAL TRIALS



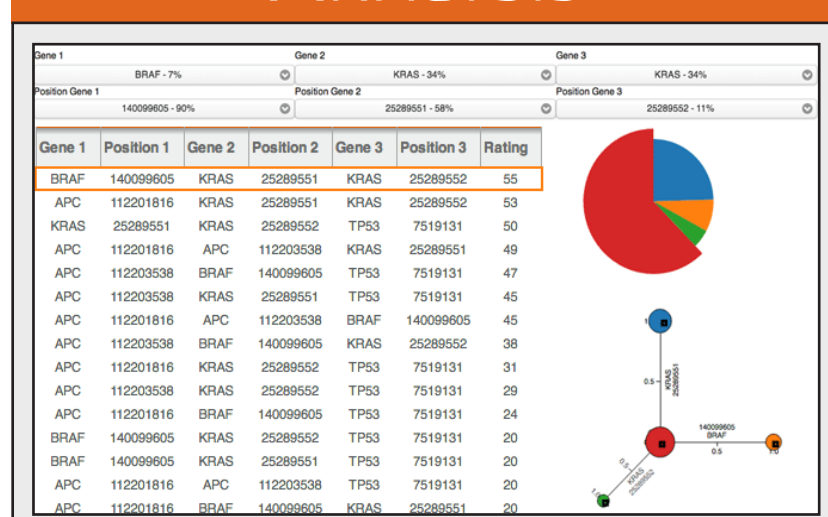
Automatic proposal of clinical trials based on individual patient anamnesis

PATHWAY TOPOLOGY ANALYSIS



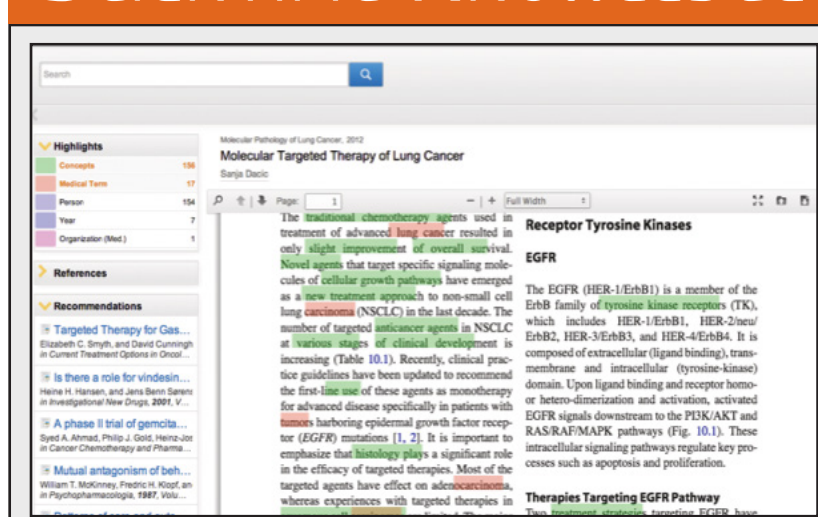
Graph topology analysis based on curated cellular pathways

PATIENT COHORT ANALYSIS



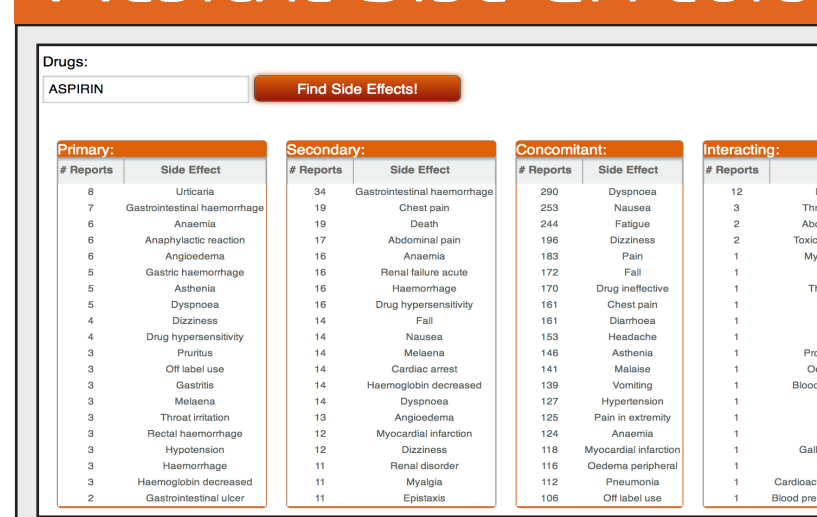
Exploration of genotypic and phenotypic correlations in patient cohorts

IDENTIFICATION OF SCIENTIFIC KNOWLEDGE



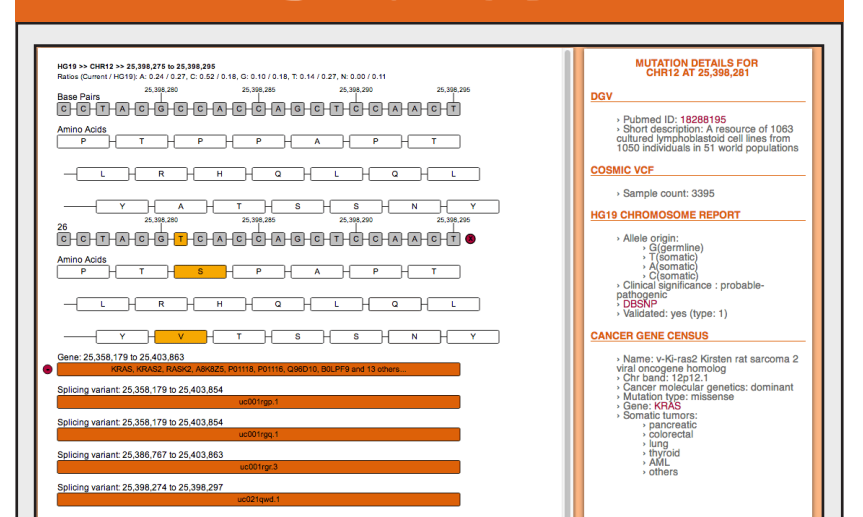
Screening of latest scientific publications relevant for personalized treatment

ASSESSMENT OF MEDICAL SIDE EFFECTS



Identification of drug interactions to ensure pharmacotherapy safety

INTERACTIVE GENOME BROWSER



Interactive exploration of genomes at any site and any level of detail

PROCESSING

Prioritized Task Scheduling
Prioritized resource management across departments and individual users

Annotation and Updater Framework
Permanent monitoring and integration of international annotation databases

User-Defined Analysis Workflows
Graphical modeling of analysis workflows integrating third-party tools

Combining Structured and Unstructured Data
Application of text analysis features to identify medical entities

Parallel Execution Framework
Automatic allocation of resources enabling concurrent workflow execution

Resource Monitoring and Accounting
Fair and equal use of computational resources

DATA

> 160k Medical Terms and Pharmaceutical Ingredients

>> 135k Worldwide Clinical Trials

>> 130M Annotations from International Databases

DNA Sequences, e.g. 1,000 Genomes Project

>> 4M Research Publications

>> 1.5k Cellular Pathways

Contact

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ANALYZE GENOMES
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Future SOC Lab
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