# The AETIONOMY Knowledge Base

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innovative

medicines initiative

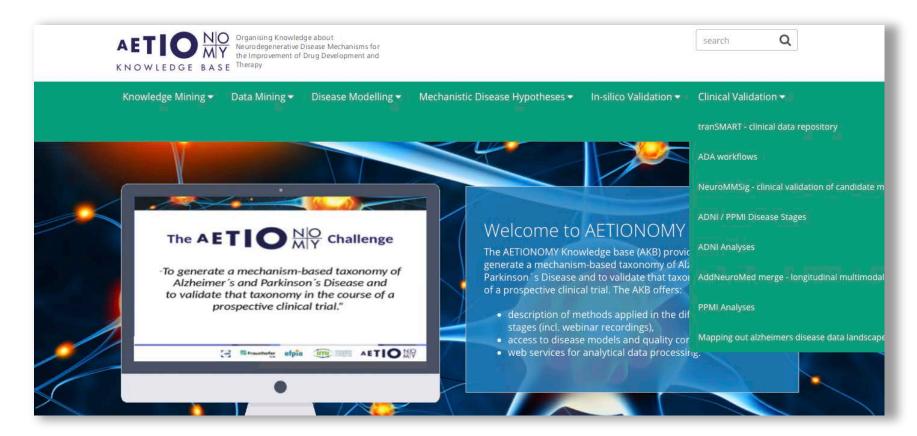


#### Mission

To increase knowledge of the causes of Alzheimer's and Parkinson's Disease by generating a mechanism-based taxonomy; to validate the taxonomy in a prospective clinical study that demonstrates its suitability for identifying patient subgroups (based on discrete disease mechanisms); to support future drug development and lay the foundation for improved identification and treatment of patient subgroups currently classified as having AD or PD.



## **AETIONOMY Knowledge Base (AKB)**



https://data.aetionomy.scai.fraunhofer.de/













#### Integrated Storage: Bring heterogeneous data together





## **AETIONOMY Knowledge base**

- One point stop for AETIONOMY knowledge and services
- platform to generate a mechanism-based taxonomy of Alzheimer's and Parkinson's disease
- description of methods applied (incl. webinar recordings, publications and use cases)
- access to disease models and quality controlled data
- web services for analytical data processing













### **AETIONOMY Knowledge Base: who can benefit?**



#### Users who can benefit

AETIONOMY intends to disseminate the approaches and methodologies and how they can be adap can benefit differently from these information:

- General public and Patient Organizations with new information for both the diseases.
- · Policy makers with insights into new disease models,
- · Bioinformaticians and students evaluating new data mining approaches,
- . Medical experts validating new mechanistic disease hypotheses, and

## Addresses targeted Usergroups with linked information:

- 1. General public and Patient Organizations
- 2. Policy makers
- 3. Bioinformaticians and students
- 4. Medical experts







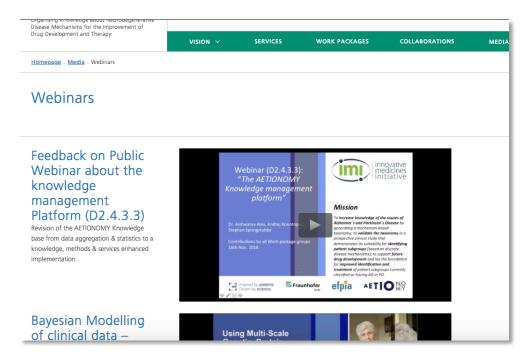






#### **AETIONOMY Knowledge Base: Awareness and training**

#### Offers training on bioinformatic methods and services



https://data.aetionomy.scai.fraunhofer.de/

https://www.aetionomy.eu/en/media/webinars.html

## Links to services, tutorials and webinars:

- 6. How to get access to services?
- Information retrieval (literature mining): SCAIView Neuro
- BEL disease model viewer (collected knowledge): BEL-Commons
- Mechanistic candidates repository (graph mining): NeuroMMSig
- Data features:
   Bayesian Network Viewer
- Trajectories:

  EPAD longitudinal viewer Images Google Drive
- Clinical data management: tranSMART
- Statistical and analysis workflows:
   ADA













### **AETIONOMY Knowledge Base: Linked to project phases**



## Project phases and applied approaches & methods:

- 1. Knowledge mining
- 2. Data Mining
- 3. Disease modelling
- Mechanistic disease
   hypotheses
- 5. In-silico validation
- 6. Clinical validation





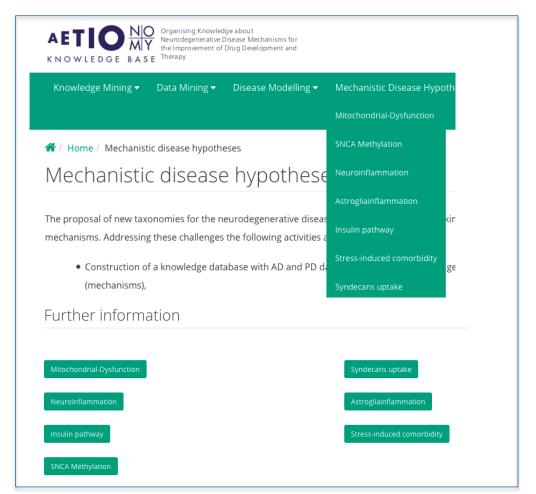


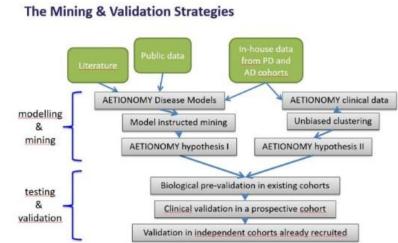






#### **AETIONOMY Knowledge Base: Linked to Mechanistic Disease Hypothesis**





https://data.aetionomy.scai.fraunhofer.de/mechanistic-disease-hypotheses







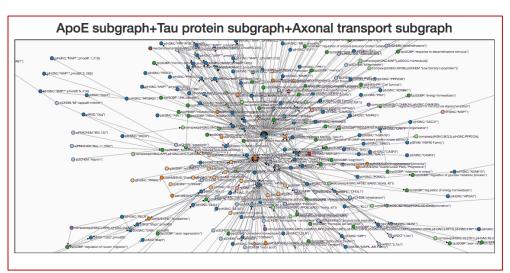




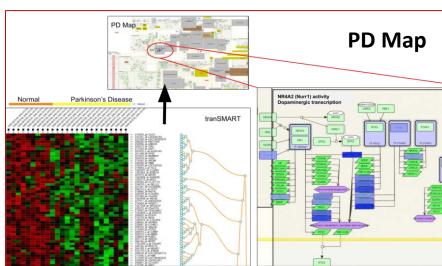


#### **AETIONOMY Knowledge Base: Disease Models**

#### PD and AD Disease maps and models



AD BEL model visualized via the NeuroMMSIG <a href="https://neurommsig.scai.fraunhofer.de/">https://neurommsig.scai.fraunhofer.de/</a>
Hosted and updated by Fraunhofer



PD Map visualized via the MINERVA platform <a href="https://pdmap.uni.lu/minerva/">https://pdmap.uni.lu/minerva/</a>
Hosted and updated by UL







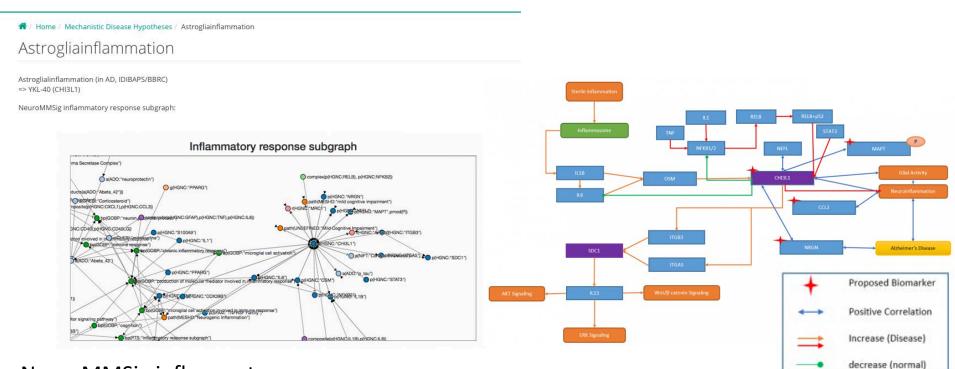






#### **AETIONOMY Knowledge Base: BEL Disease Models**

#### Disease maps as support to hypothesis generation



NeuroMMSig inflammatory response subgraph (AD BEL model)

Cartoon of the candidate mechanism













#### **AETIONOMY Knowledge Base: SBML Disease Maps**

Disease maps as support to hypothesis generation

Overlayed on disease maps (PD Map)

NR4A2 (Nurr1) activity Normal Parkinson's Disease Dopaminergic transcription tranSMART Identifying relevant pathways and upstream or downstream players

Differentially expressed genes from analysis (tranSMART)











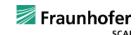


#### **AETIONOMY Knowledge Base: Data harmonisation and curation**

Systematic Inclusion of clinical, EFPIA and public datasets in the **tranSMART clinical repository** 









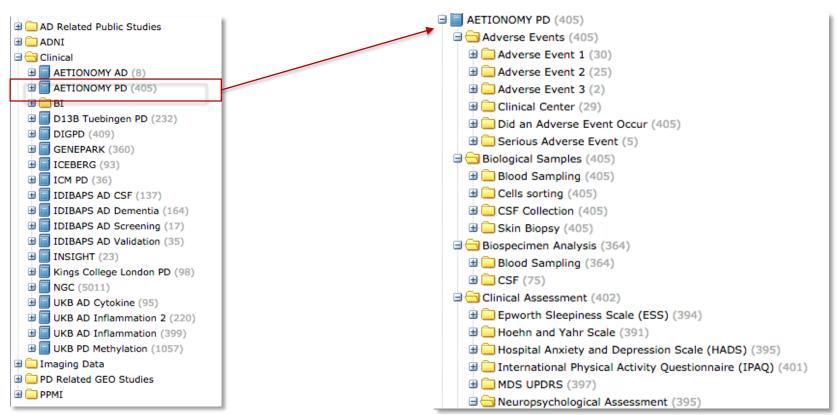






#### **AETIONOMY Knowledge Base: Datasets available through tranSMART**

Systematic Inclusion of clinical, EFPIA and public datasets in the tranSMART clinical repository



https://aetionomy.uni.lu/transmart/login/auth







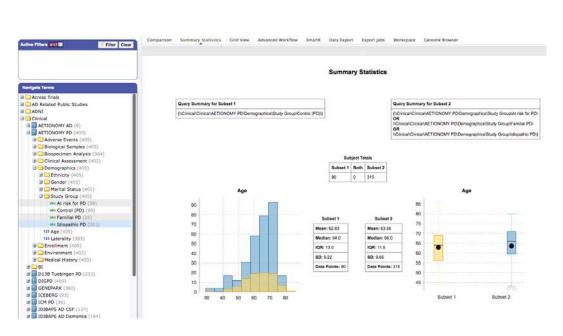


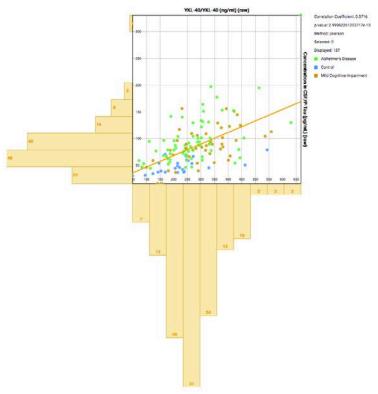




### **AETIONOMY Knowledge Base: Datasets available through tranSMART**

Advanced analytics through smartR analytics in tranSMART

















### AKB Component: Ada - advanced statistical and analysis workflows



https://aetionomy-ada.lcsb.uni.lu/







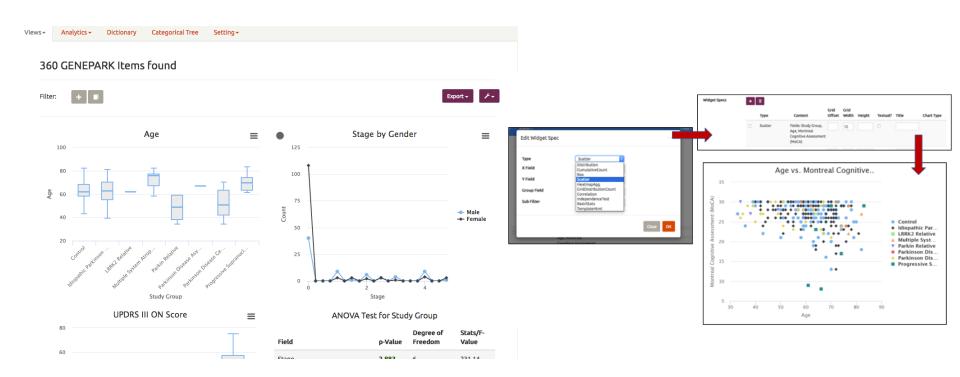






### AKB Component: Ada - advanced statistical and analysis workflows

Customised user views and dynamic and interactive analytics















#### Users who can benefit

AETIONOMY intends to disseminate the approaches and methodologies and how they can be adapted to other projects and research areas. User groups can benefit differently from these information:

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#### **After AETIONOMY ...**

- The AETIONOMY Knowledge Base will continue to be maintained for another 5 years
- All clinical and public datasets will be available for access through a Data Access Committee
- The disease models and services will continue to be developed and updated regularly













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Questions?









