

Final conference 24th June 2020



NanoCommons KnowledgeBase: Supporting a lasting legacy from SmartNanoTox data and models









- Remove barriers from nanosafety regulatory and industry processes
- Develop an integrated Knowledgebase to facilitate development and application of regulatory tools such as grouping & read-across
- Create an interconnected community via a FAIR data single market
- Enable full exploitation of EU-funded research data & promotion of data driven innovation leading to positive socioeconomic impact.





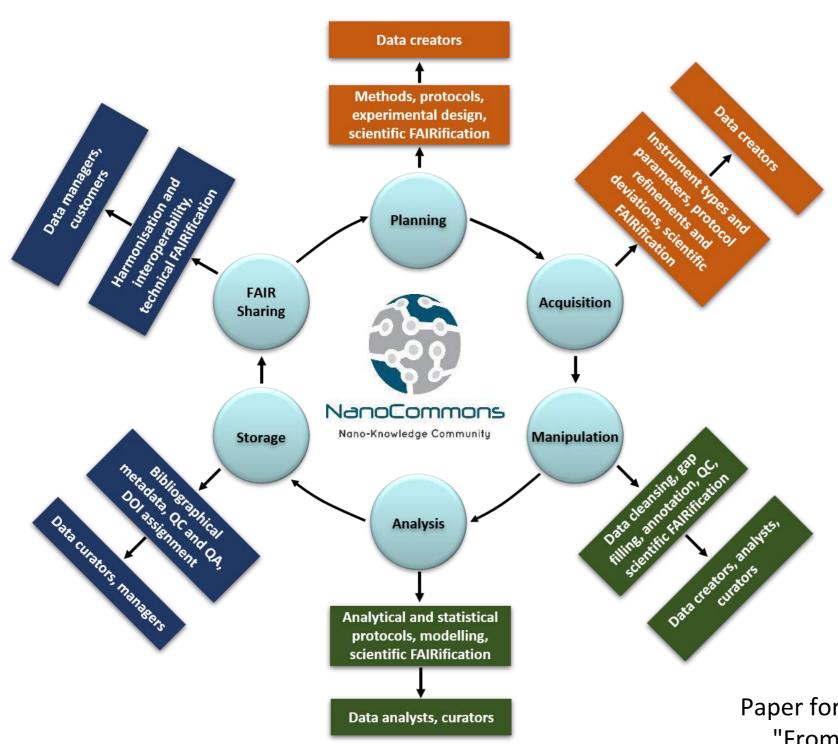
Data Processing & Analysis





Data lifecycle & metadata





Metadata is "data about data."

Descriptive metadata – assay type, cell line used, concentrations, calibrations, instrument and software parameters, etc.

Technical metadata – statistical approach, data cleaning / gap filling approach etc.

Bibliographic metadata – who generated the data, publication information, licence information, etc.

Paper for submission to Nanomaterials special issue "From Nanoinformatics to Nanomaterials Risk Assessment and Governance"





Samples and

implementation

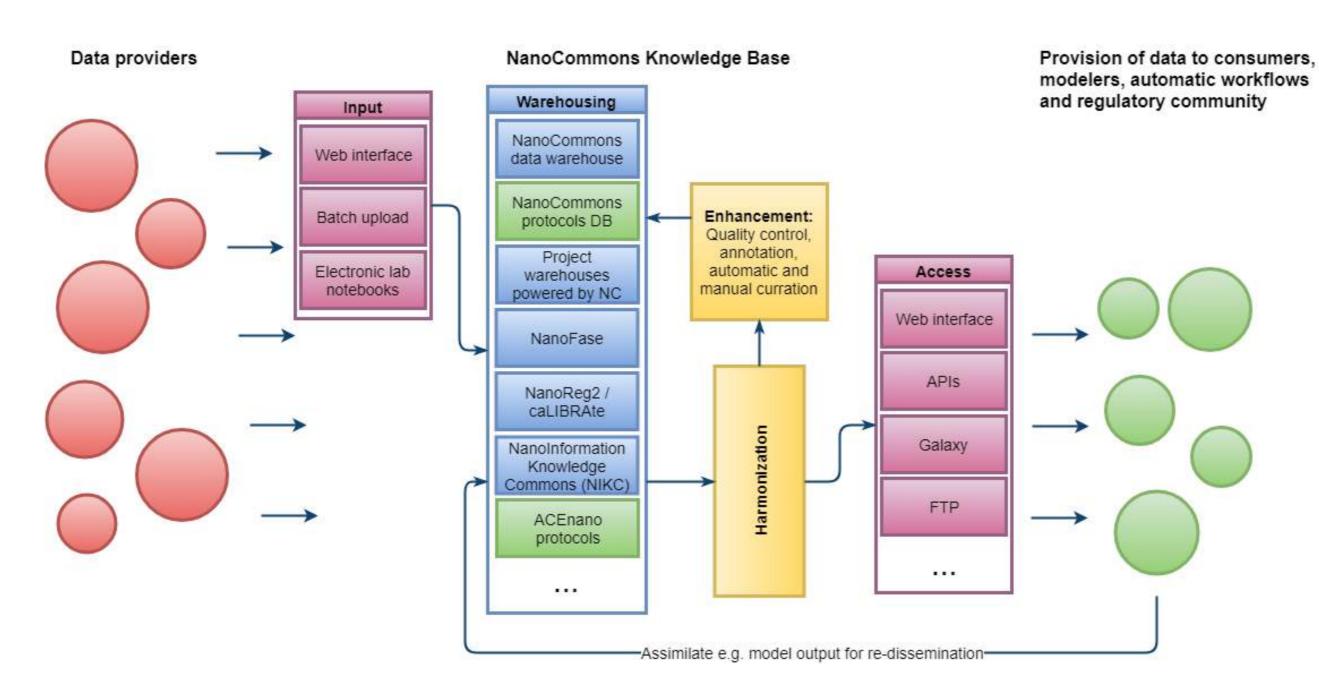
endpoints

Experimental design and protocol implementation





NanoCommons KB Architecture concept

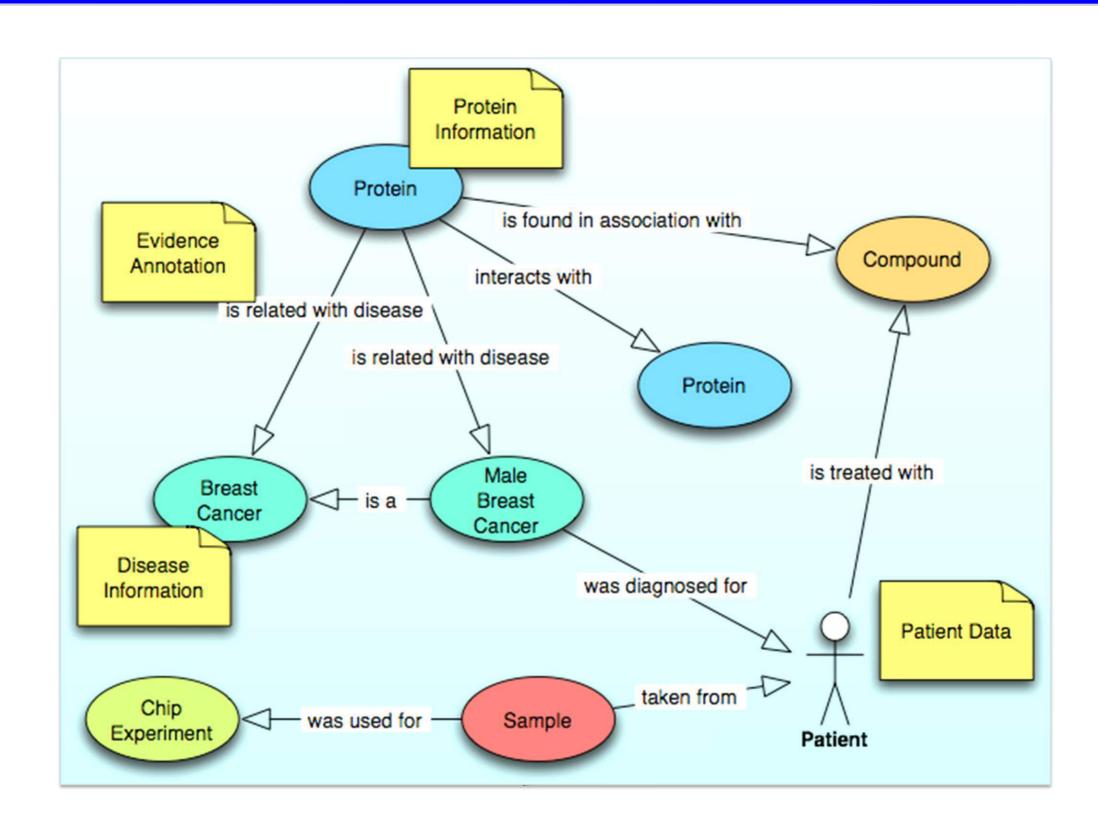






Knowledge management - handling complexity

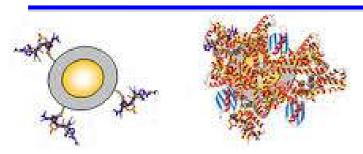




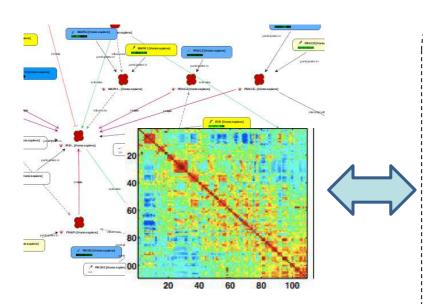
Semantic Model

Coupling data and applications

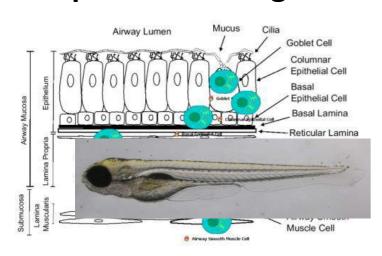




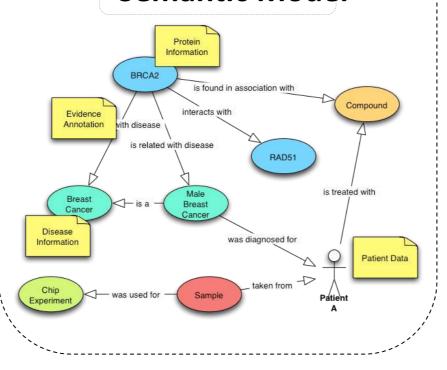
Particle characterisation



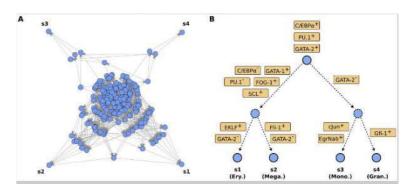
Molecular "omics" data and prior knowledge



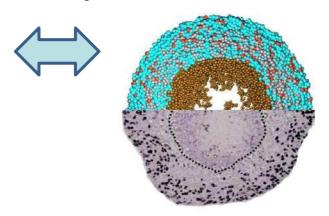
Knowledge management Semantic Model



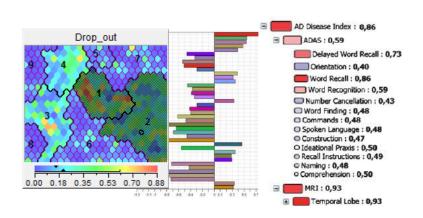
Network inference and data analysis



Computational Simulation



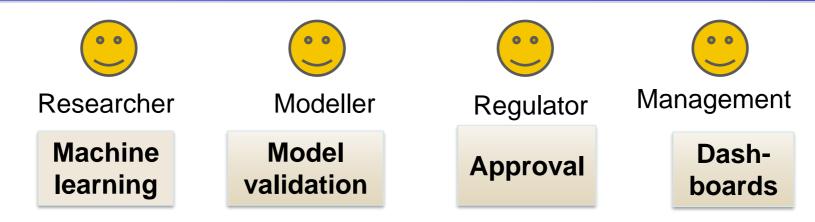
Biomarker profile

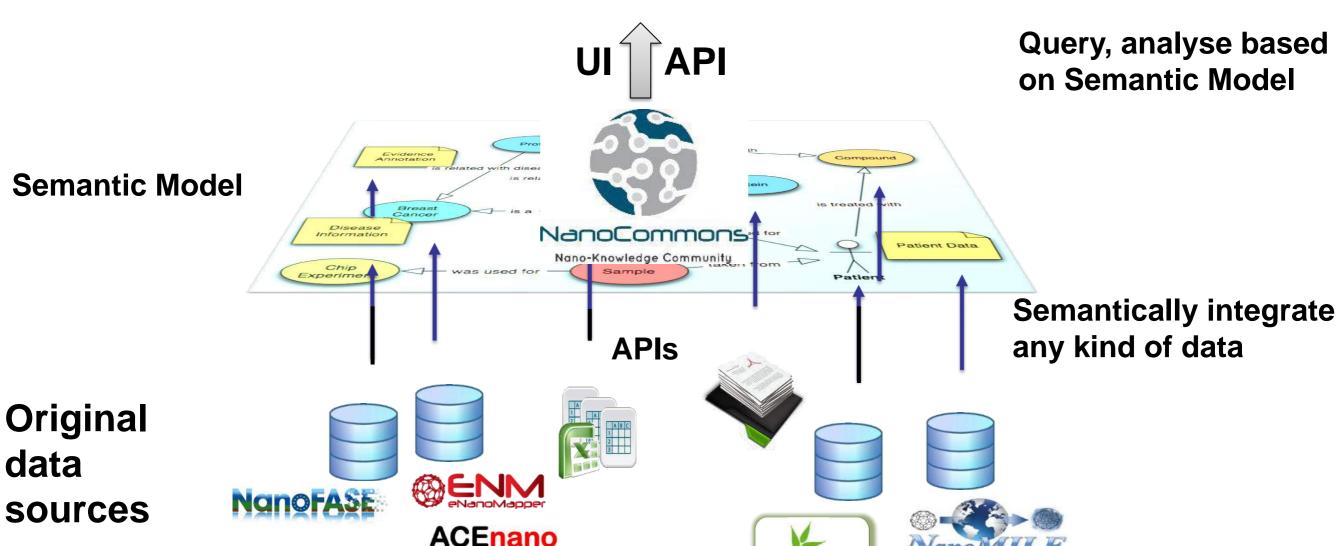


Classical Toxicology

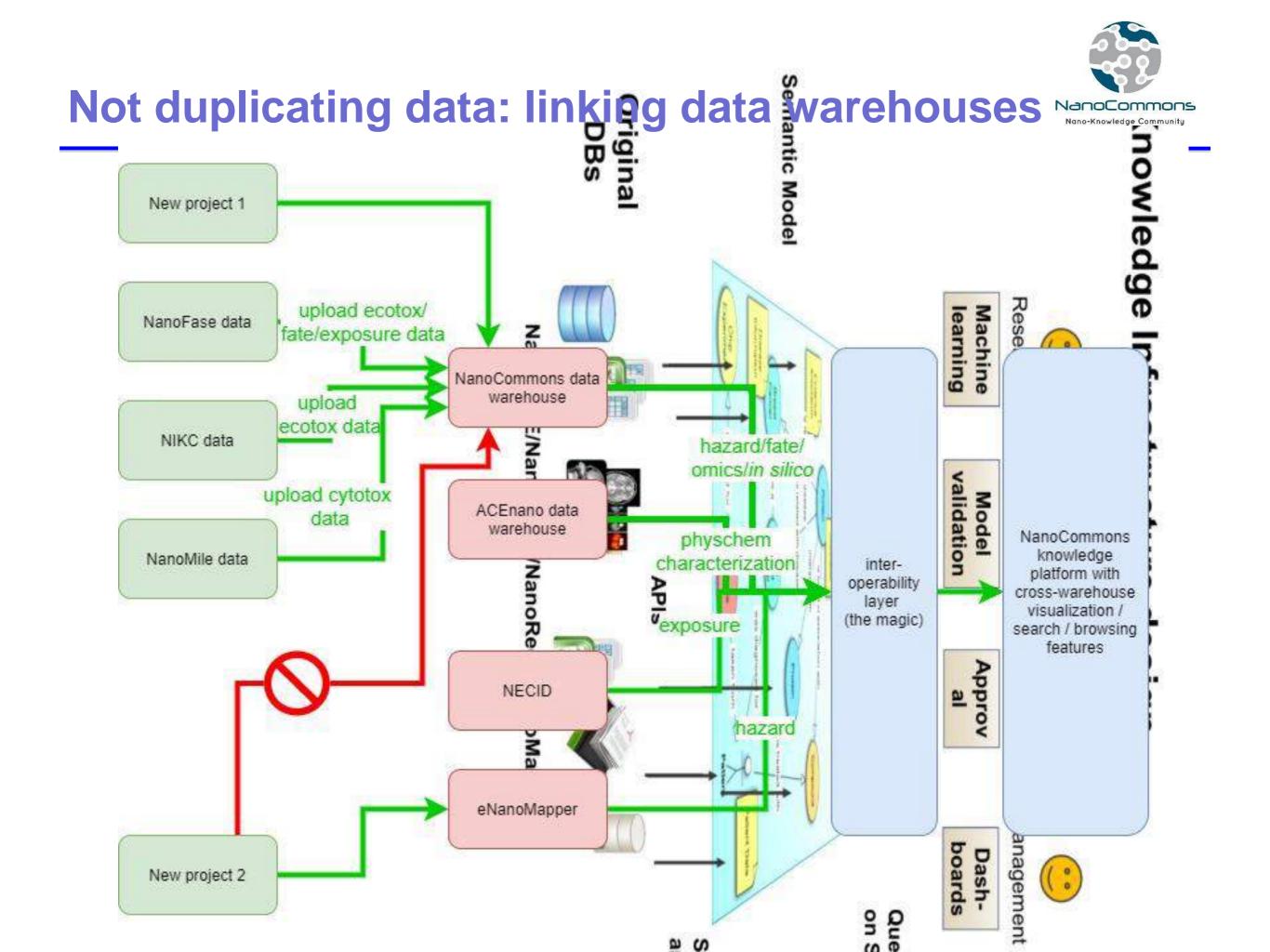


Semantic Model for integration and interoperability





NIKC



NanoCommons Knowledge Base - Content



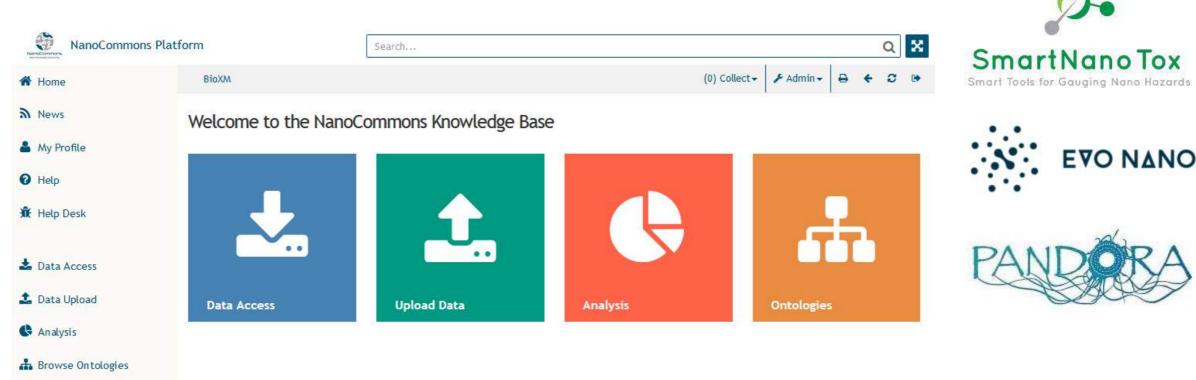








https://ssl.biomax.de/nanocommons



~500 Particles, Phys-Chem Characterisations, Protocols, Toxicology Experiments, Omics, Mesocosm from NanoMILE/NanoFASE (NanoMILE data freely available, NanoFASE data embargoed 2 years (DTA available))

In-silico descriptors, Corona predictions, groupings, alerts from NanoCommons / NanoSolveIT

Literature mined datasets (>250 studies) relating to > 350 particles

Expanding rapidly via ongoing TA projects.....



Services - Data Sustainability beyond Archives

Archive:

- Data store
- Metadata (Description)





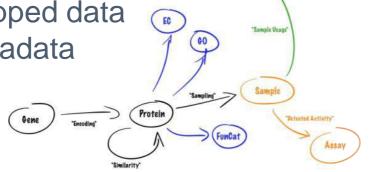
Functions:

- Browse
- Search
- Retrieve
- Analyse



Semantic integration:

- Harmonized and mapped data
- Metadata



Functions:

- Navigate along connections
- Combine molecular, phenotype and environmental Information
- Visualize



Enrichment with e.g. Biological knowledge or Environmental Data

- Genomes
- Pathways
- Protein-protein-interactions
- Transcription factor network
- Water/Soil chemistry (pH, ionic strength, NOM)
- Co-pollutants
- Species distributions / sensitivities
- Long range trends / future predictions





Archive:

- Data store
- Metadata (Description)



Functions:

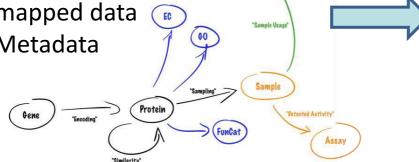
- **Browse**
- Search
- Retrieve
- Analyse



Semantic integration:

Harmonized and mapped data





Functions:

- Navigate along connections
- Combine molecular, phenotype and environmental Information
- Visualize

Example NanoMILE/NanoFASE knowledge base content:

- Genomes
- **Pathways**
- Protein-protein-interactions
- Transcription factor network
- Particles
- Orders/procurement
- Material safety sheets
- Ecotoxicological data
- 'Omics



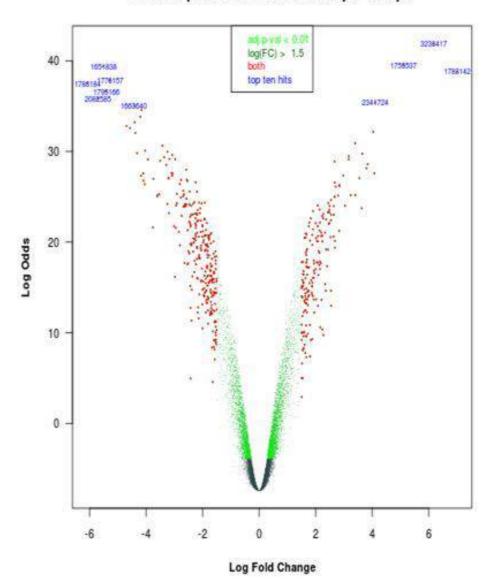
Integrated Services -**Differential Expression analysis**



Analysis RNA-Seq Differential Expression ■ Analysis RNA-Seq Differential Expression 2.2.1 Limit 50 😠 LIMIT

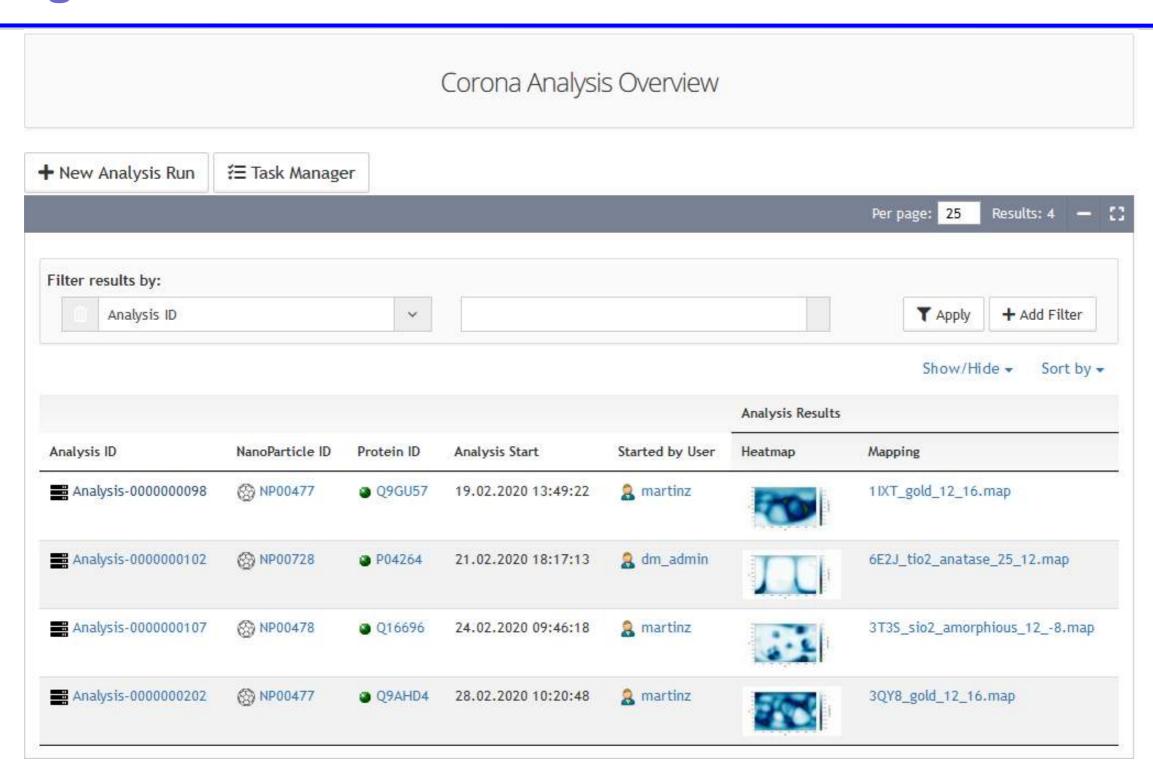
x 🗹 2.3.1 p-value 0.05 😠 **PVAL** 2.4.1 Foldchange 1.5 😮 LogFC x 🗹 Adjust 2.5.1 Adjust p-value for multiple testing by Benjamini Hochberg 😵 x 🗹 Cutoff A549_AgNO3_01hrs_3Rep 😵 A549_AgNO3_01hrs_4Rep 😣 Group1 × 🗹 A549_AgNO3_01hrs_1Rep 😵 A549_AgNO3_01hrs_2Rep (8) A549_AgNO3_06hrs_1Rep 😵 A549_AgNO3_06hrs_2Rep 😵 Group2 × 🗹 A549_AgNO3_06hrs_3Rep 😣 A549_AgNO3_06hrs_4Rep 🗴 A549_AgNO3_24hrs_1Rep 😵 A549_AgNO3_24hrs_2Rep 😵 Group3 × 🗹 A549_AgNO3_24hrs_3Rep 😵 A549_AgNO3_24hrs_4Rep 😮

Volcanoplot for contrasts: Group1-Group4



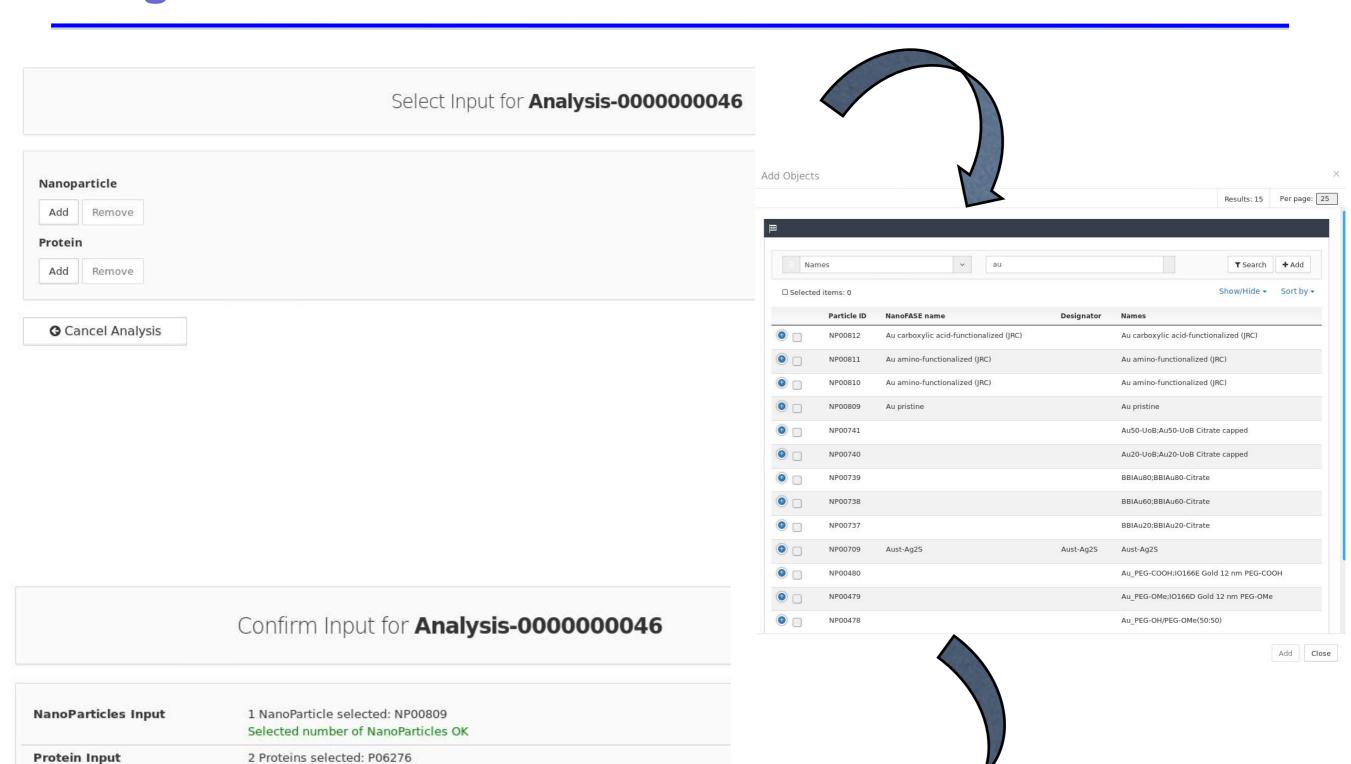


Integrated Services – Corona model





Integrated Services – Corona model





Q9BV73

Please select exactly one Protein



Integrated Services – Corona model

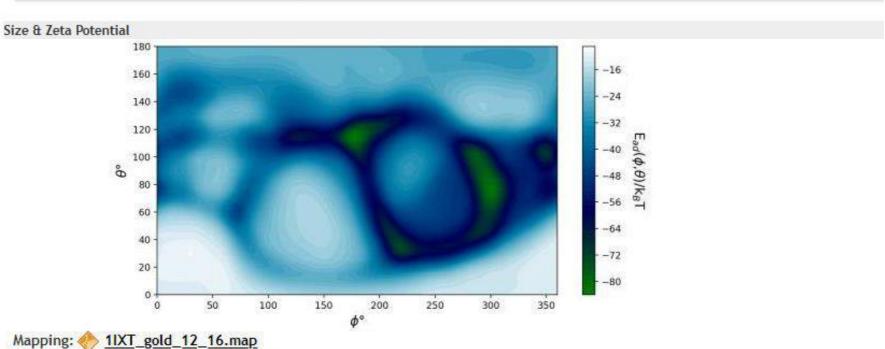
LogFile:

Analysis-0000000098.log



Conotoxin Gm9.1	
	Conotoxin Gm9.1

1IXT C
structure of a novel p-superfamily spasmodic conotoxin reveals an inhibitory cystine knot motif



Next steps in supporting SmartNanoTox



TA project already underway for SmartNanoTox Experimental data

TA in development for computational data and models:

1. Missing a lot of the ontology terms needed for modelling:



Wiki to define the terms (with nanoinformatics project experts / EMMC)

On the fly annotation using agreed terms (while they are being formalized into ontologies, e.g. ENM)

2. Developing tool for harmonization of MODA templates:



Questionnaire format with drop-down menus (as already done for SOPs)

3. Inputting the computational materials / data into KnowledgeBase

Preserving the



legacy through open & FAIR data

Acknowledgements & Thanks































Knowledge Base with FAIR data

- Metadata fully searchable (including Ontologies e.g. NCBI Taxon, GO, eNMP)
- API based data access (Jaqpot)
- Generation of ontological descriptors for data
- Data upload & download in a format suitable for modelling
- Tool integration (Expression analysis, Modelling)