

THE FUTURE OF DATA INTEGRATION:

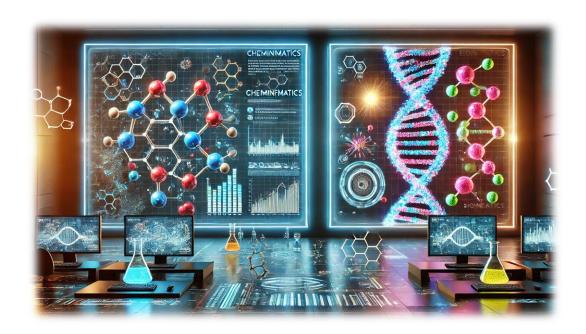
BIOINFORMATICS AND CHEMINFORMATICS

Fabrizio Abbinante, Head of iDATA Unit, EFSA

Stakeholder Forum, 27 Nov 2024 Brussels

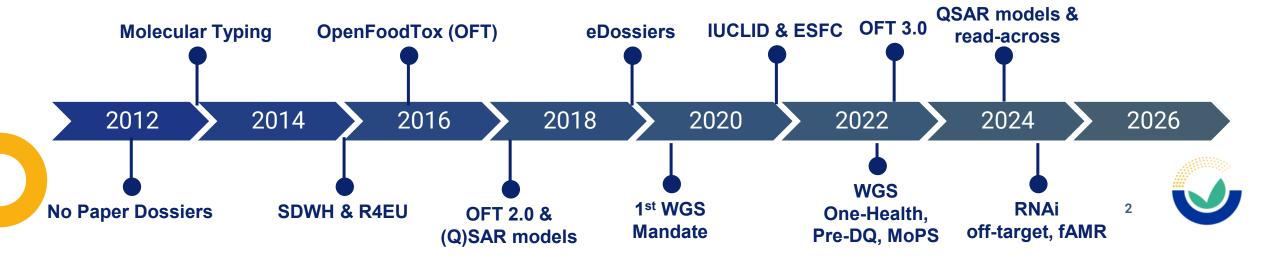


BIOINFORMATICS AND CHEMINFORMATICS @ EFSA



Provide an update on EFSA's progress in developing **bioinformatic** and **cheminformatic** data pipelines and tools (data and code) ensuring interoperability and (re-)use of data at data ingestion, validation and (re-)use stages.

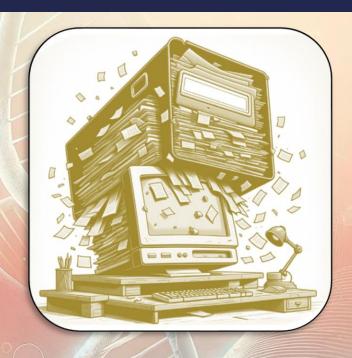
Gather feedback from stakeholders on their requirements and explore the most effective ways to engage them in these initiatives.



BIOINFORMATICS UNTIL 2020: NO WGS STORAGE, NO PIPELINES, NO TOOLS

OFFICERS MANUALLY
SCREEN A GENETIC
SEQUENCE LETTER BY
LETTER





GENOMICS DATA
RECEIVED FROM
APPLICANTS

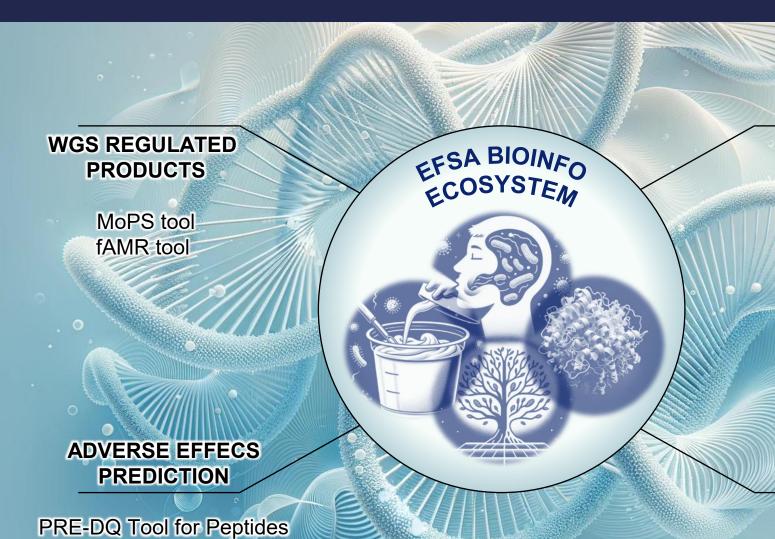
NO PIPELINES
NO TOOLS

MANDATE TO ANALYZE WGS DATA





PRESENT: EFSA INTERNAL BIOINFORMATIC ECOSYSTEM OF TOOLS



RNAi off-targets Tool

WGS FOODBORNE OUTBREKS

in collaboration
with ECDC

OTHERS (under development)

Prediction of Protein Toxicity tool

BENEFITS FROM THE USE OF BIOINFORMATICS



Faster risk assessment• Automate data analysis pipelines to deliver immediate results and promptly flag concerns for risk assessors (e.g., WGS One Health and MoPS sequencing analysis).



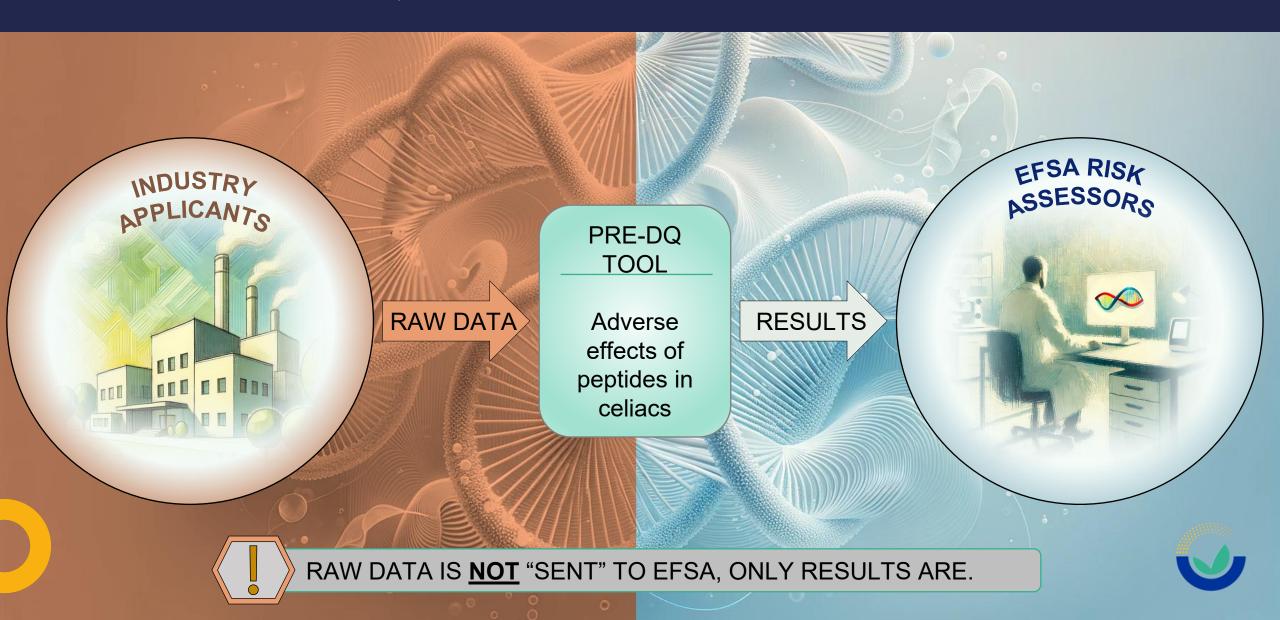
More consistent risk assessment• Enable standardized approaches for risk assessment by sharing pipelines with industry, ensuring consistent methods of analysis for all stakeholders (e.g., common bioinformatics workflows).



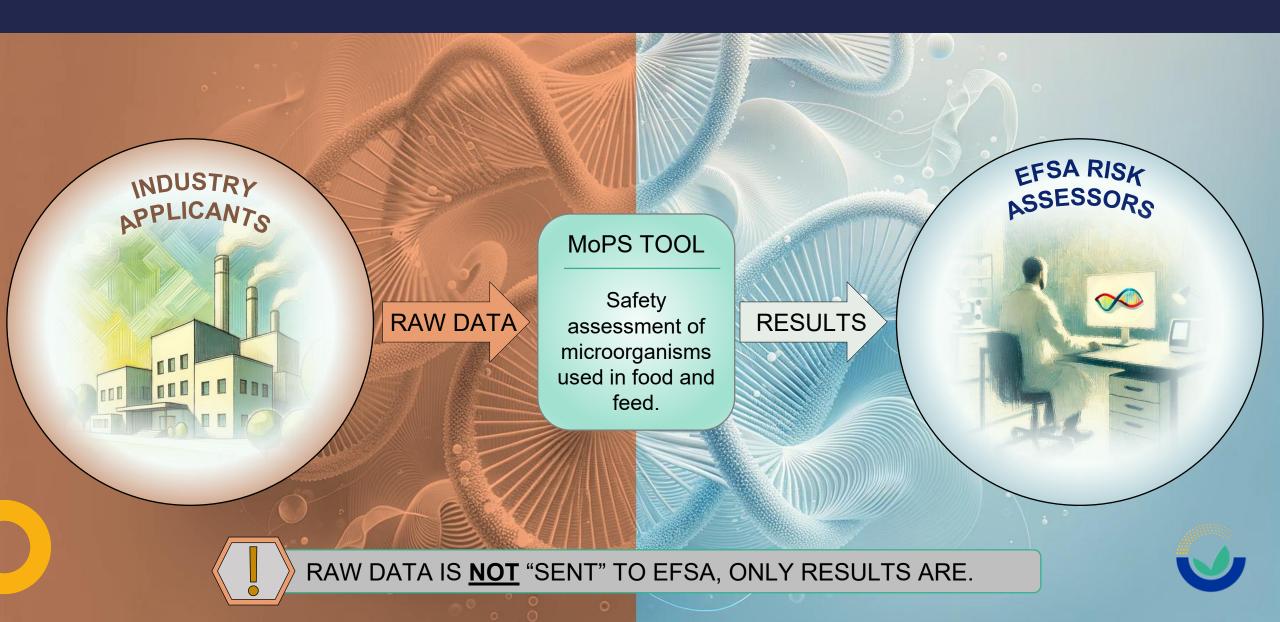
More efficient risk assessment• Utilize predictive tools to forecast adverse effects, enabling a tiered approach to hazard evaluation and improving the efficiency of risk assessment (e.g., in silico toxicity models).



PRESENT: PRE-DQ TOOL OPEN TO EXTERNALS

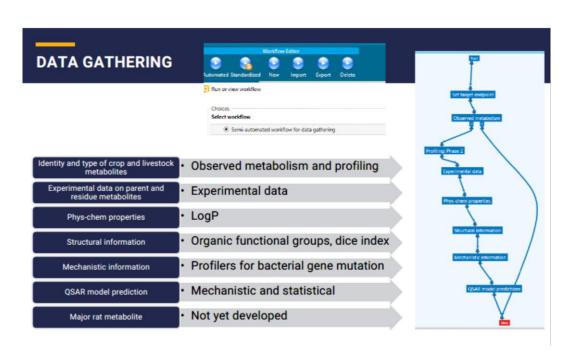


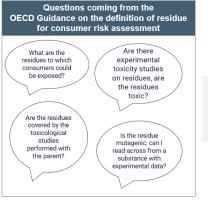
FUTURE: MOPS TOOL OPEN TO INDUSTRY

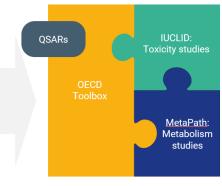


EXAMPLE OF CHEMINFORMATICS APPLICATION AT EFSA

- The OECD QSAR Toolbox is a freely available cheminformatics tool
- EFSA is developing a workflow that will be made publicly available to predict the mutagenicity of pesticide residues using IUCLID data on as input according to a new OECD Guideline
 - ✓ Identification of existing data
 - √ Toxicological predictions









OPENFOODTOX 3.0 AND IN SILICO TOOLS - PUBLICLY AVAILABLE

- 20+ QSAR models available in VEGA HUB in silico platform to predict chemical properties/toxicity for:
 - Human health
 - Animal health
 - Environment



- Physico-chemical
- New software VERA (Virtual Extensive Read-Across)
 publicly available to support read-across and grouping of
 chemicals



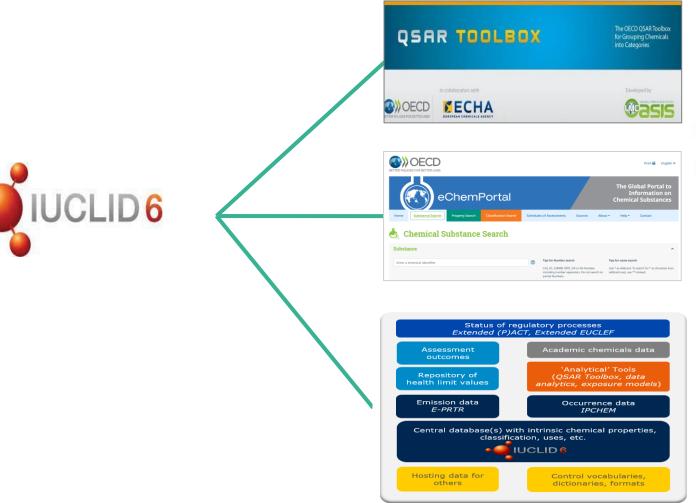
https://www.vegahub.eu/portfolio-types/in-silico-models/





EXTENDING THE USE OF IUCLID FORMAT

Structured data in **IUCLID** enables interoperability with **cheminformatics** and **dissemination** tools













EU Common Data Platform on Chemicals

BENEFITS FROM THE USE OF CHEMINFORMATICS



Faster risk assessment Facilitate the identification and use of chemicals data (identification of common pesticides metabolites) with similar structure and toxicological profile that could be assessed and managed in groups improving efficiency of risk assessment



More consistent risk assessment EFSA will develop and make available models and tools that follow OECD Guidelines and increase the regulatory certainty of the validity of the approach (e.g. assessment of pesticide residues using a workflow developed by EFSA in the QSAR Toolbox).



More efficient risk assessment Using computational tools to predict chemical properties when experimental data are not available. Predict toxicity with Quantitative Structure Activities Relationships (QSAR) models.



ASSESSMENT OF PESTICIDE RESIDUES USING QSAR TOOLBOX



OECD QSAR Toolbox will read data on metabolites in IUCLID format and generate predictions for risk assessment



CHEMINFORMATICS AND BIOINFORMATICS @ EFSA

By augmenting *Chemistry* and *Biology* with *Informatics* we have enabled **ChemInformatics** and **BioInformatics** pipelines, models and tools with concrete benefits for regulators, industry and academia.

With the advent of AI we can further augment what we have created so far in BioInformatics and ChemInformatics



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