Improving Metabolite Identification with Chemoinformatics

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June 9th 2011 ICCS 2011

Metabolome all low molecular weight molecules (metabolites) in cells, body fluids, tissues, etc.

Metabolomics

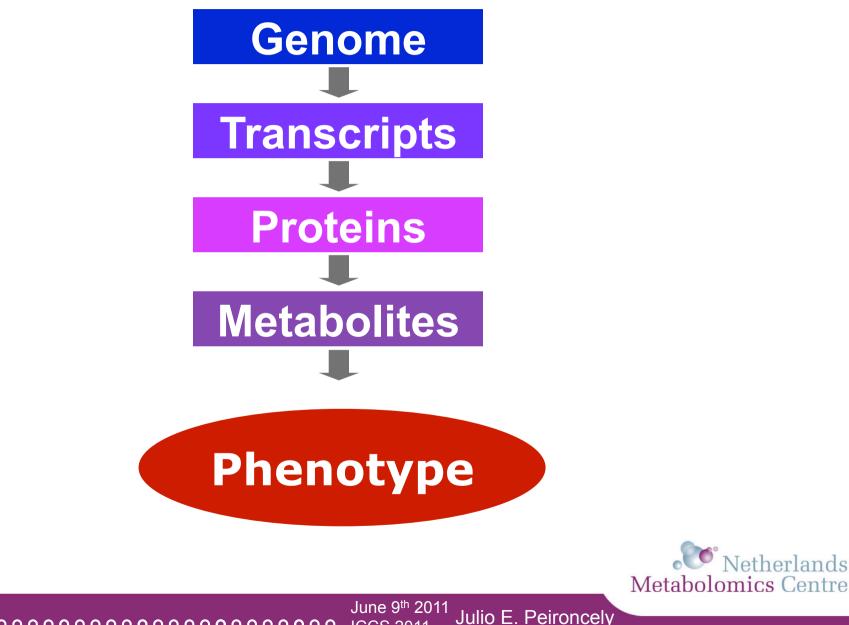
the quantitative and qualitative analysis of all metabolites in samples of cells, body fluids, tissues, etc.

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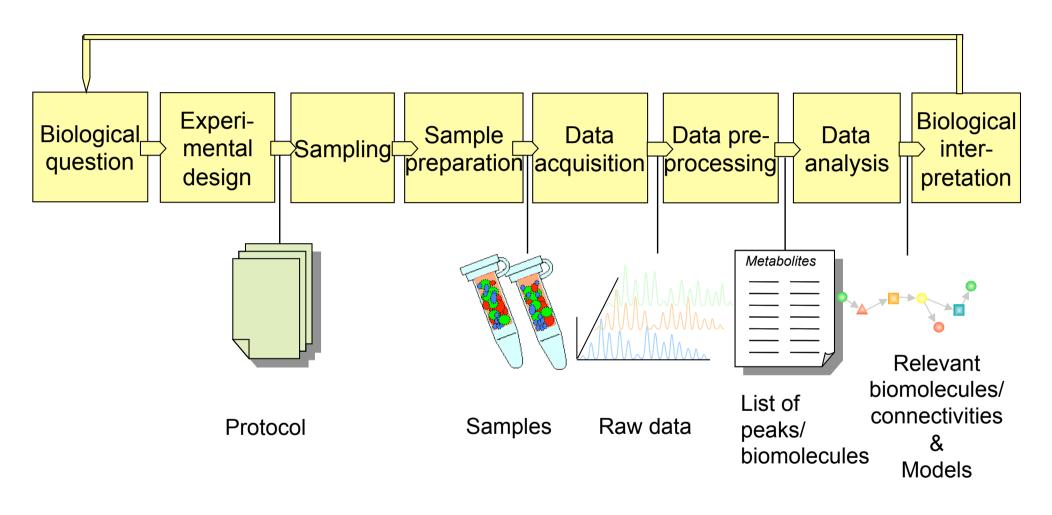
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Metabolomics



Metabolomics



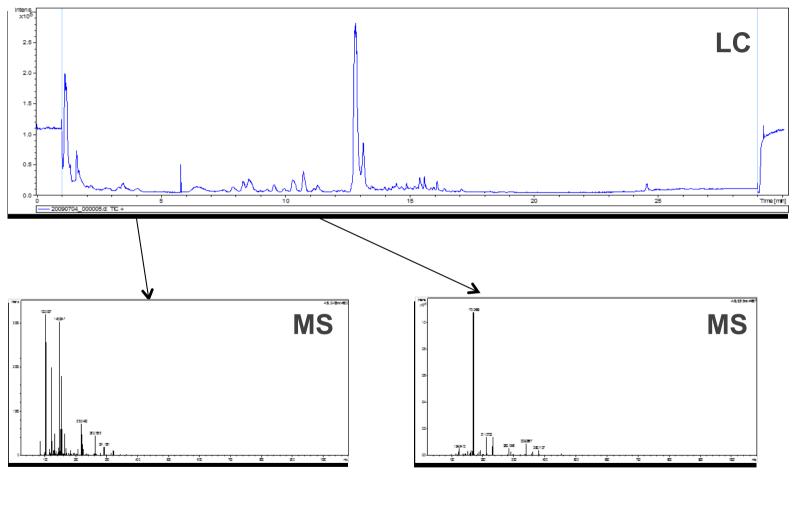


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LC-MS and Metabolite Identification

RPLC-microTOF (Bruker)

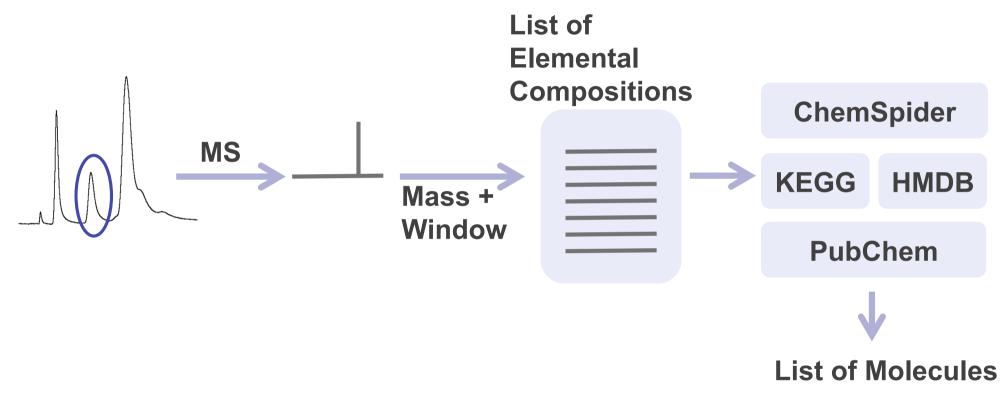


peak = m/z@rt = metabolite



Kloet et al, Metabolomics 2011 June 9th 2011 Julio E. Peironcely

When all you have is a mass and a "window"

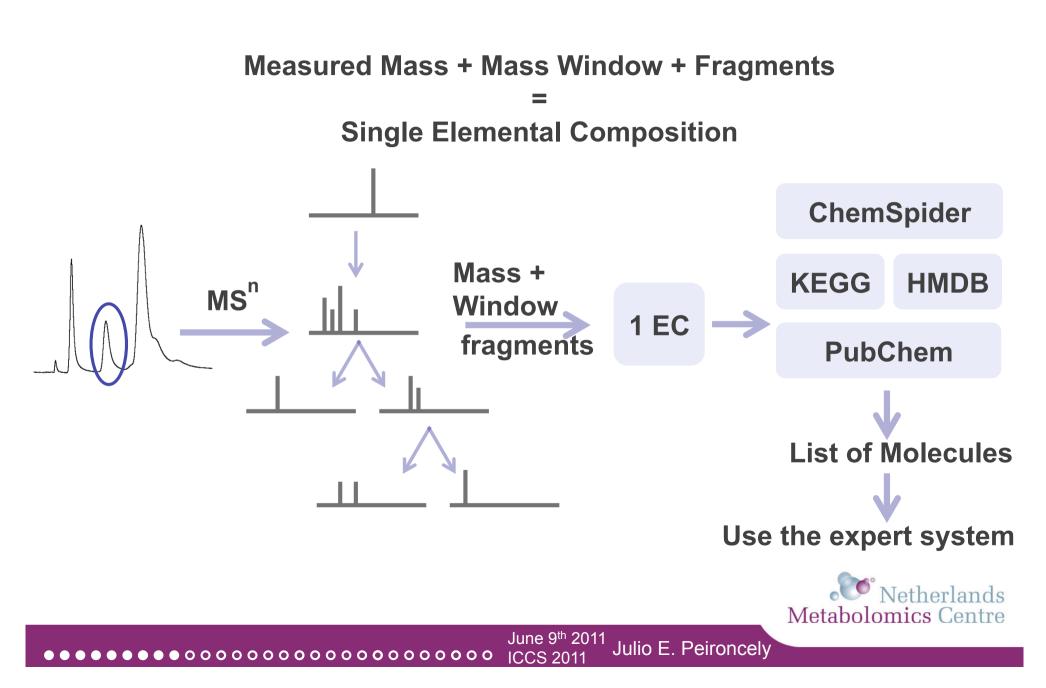


Measured Mass + Mass Window = Multiple Elemental Compositions

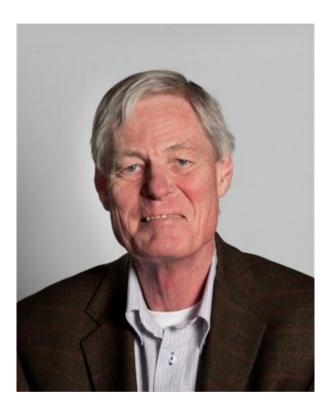


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When you have MSⁿ



The expert system



Dr. Ronnie van Doorn



Dr. Albert Tas



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Bottlenecks in Metabolite Identification

Many metabolites in LC-MS not identified

HighRes MS can obtain 1 EC = many structures

No tools for automatic identification

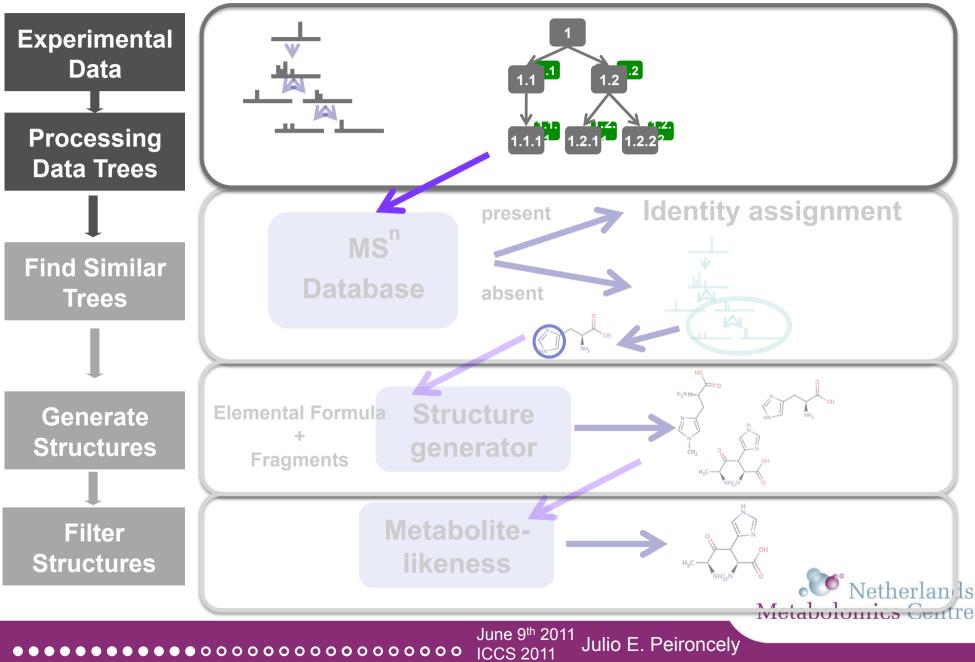
Takes long for the expert to identify metabolites

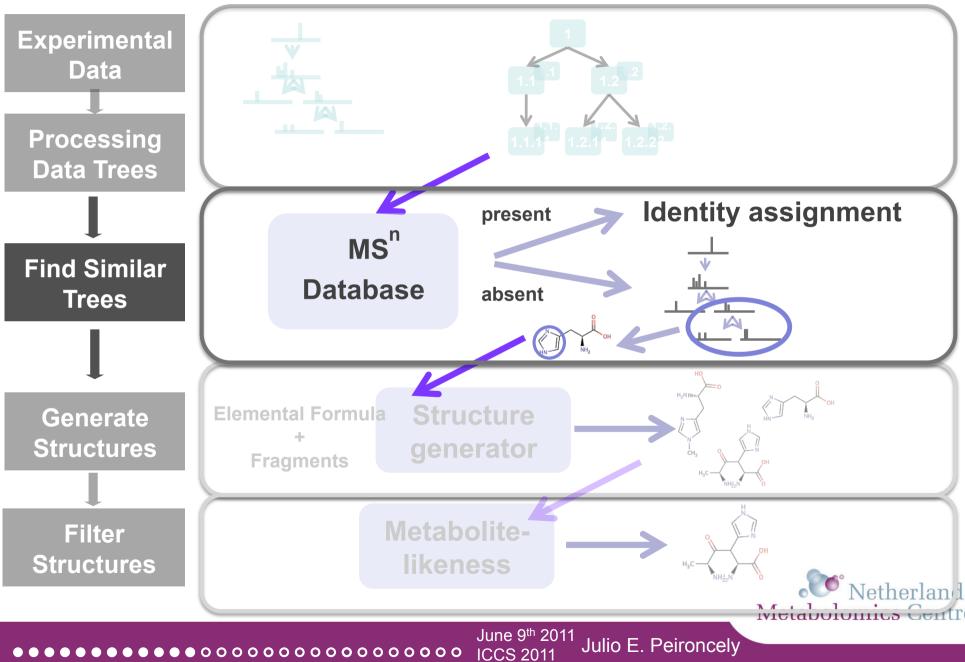
Challenges

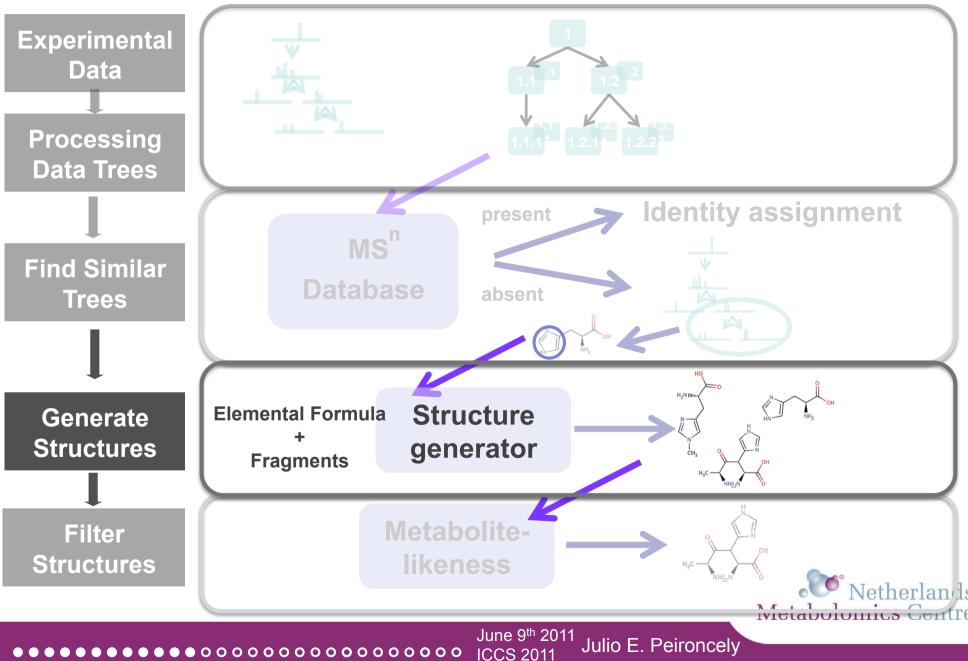
Analytical methods

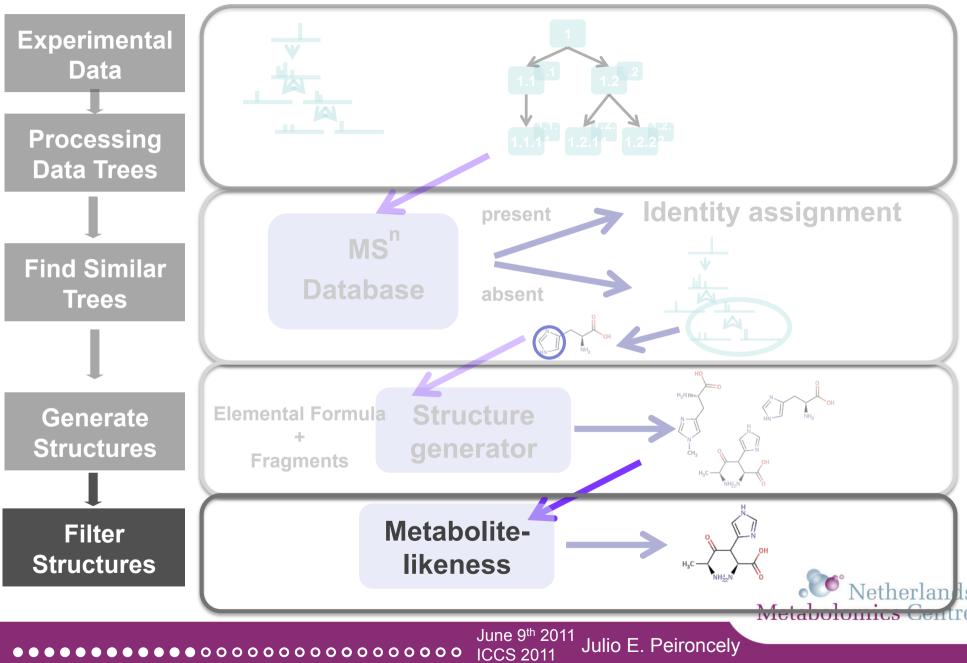
Software

Databases



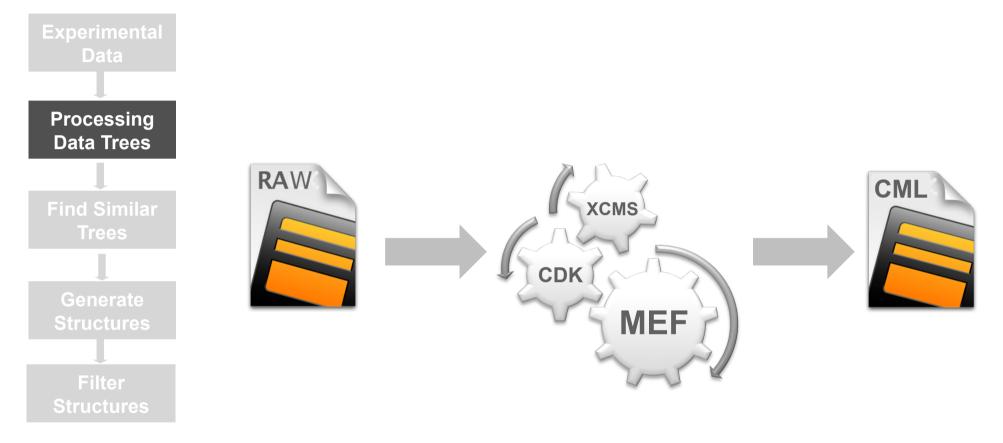






From data to information

MEF: spectral to fragmentation tree



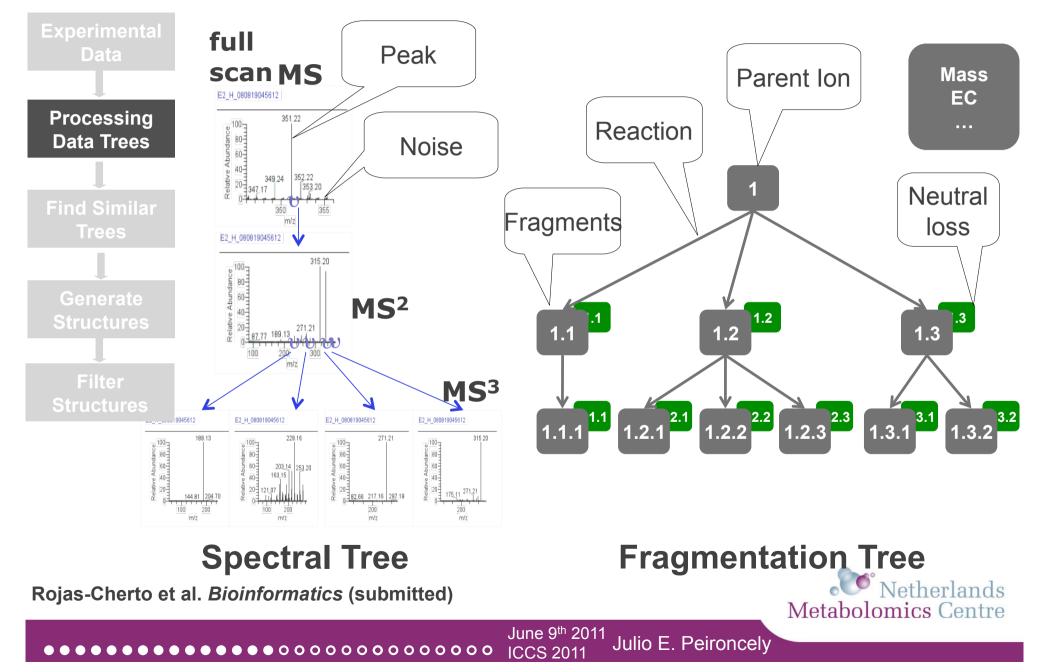
Rojas-Cherto et al. *Bioinformatics* (submitted)



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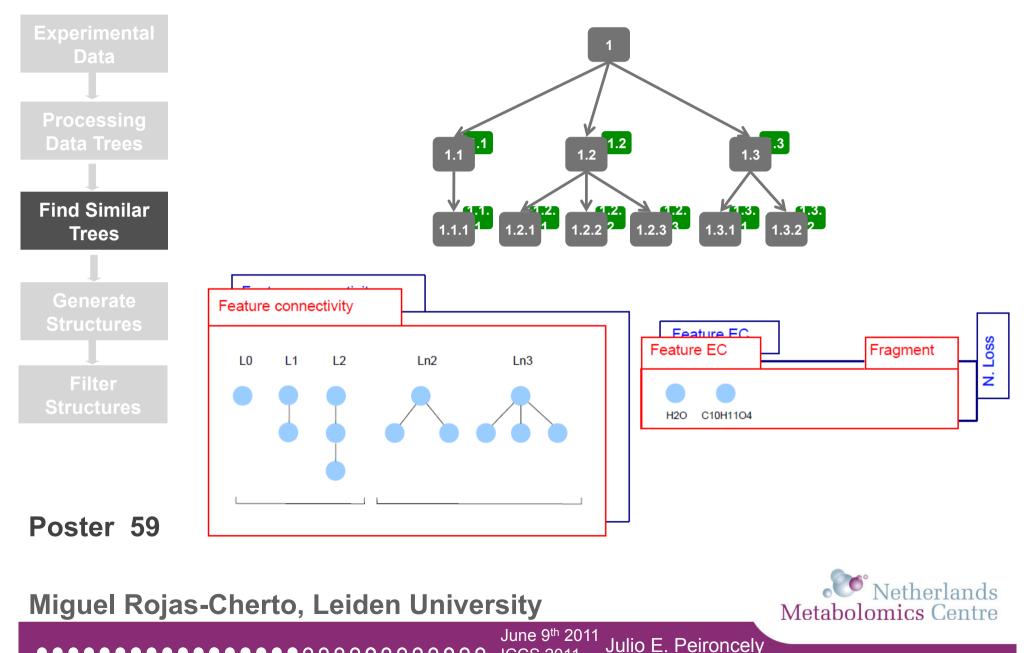
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MEF: spectral to fragmentation tree



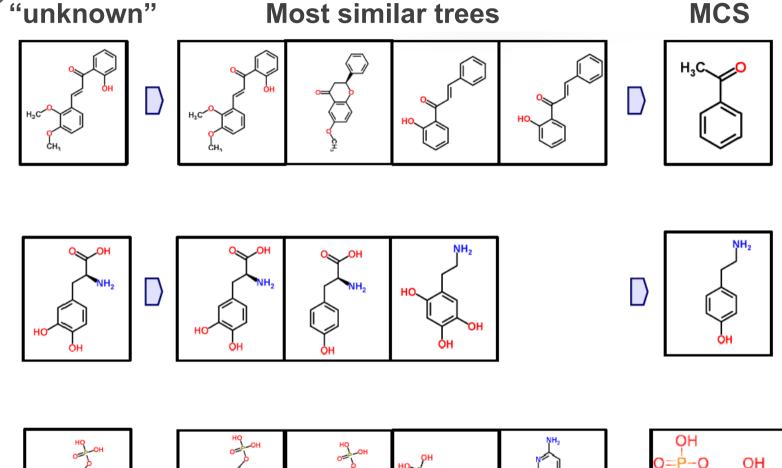
Extracting more information from your data

Fragmentation tree fingerprints



ICCS 2011

Fragmentation tree fingerprints results "unknown" Most similar trees



Poster 59

Data

Find Similar Trees

Structures

Miguel Rojas-Cherto, Leiden University

Netherlands Metabolomics Centre

ÓΗ

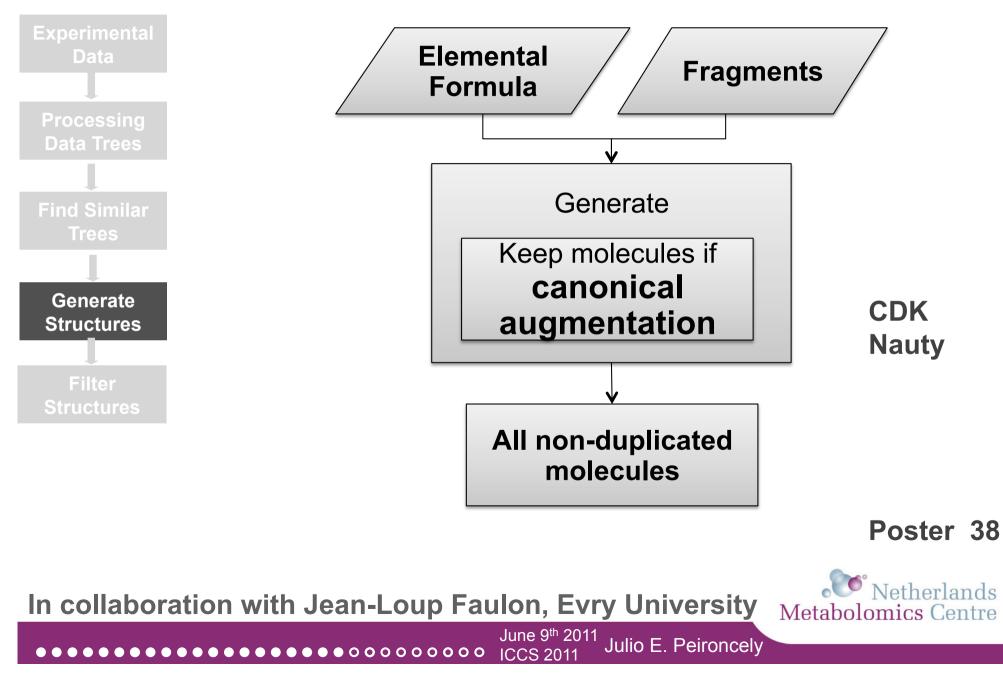
-OH

 $\left[\right)$

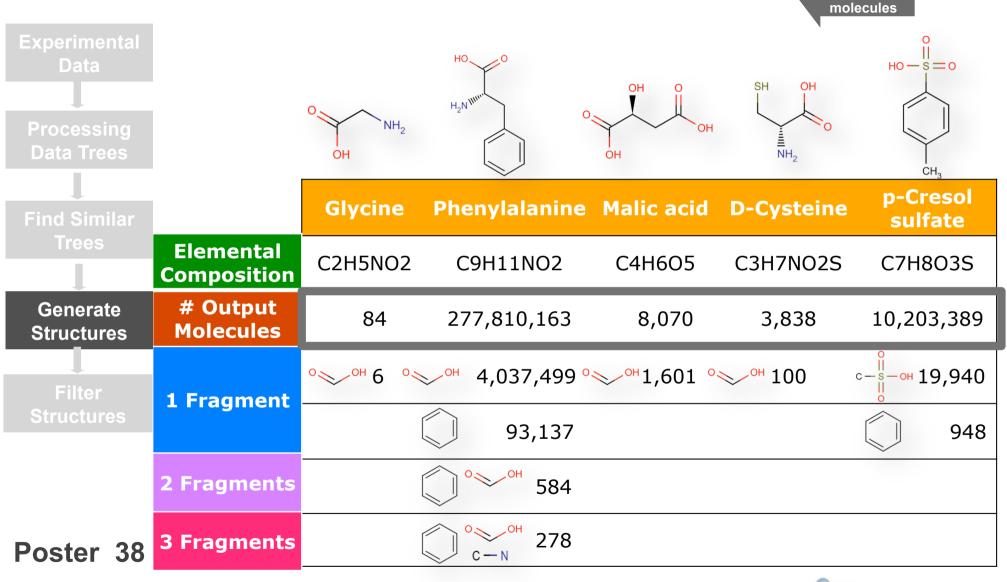
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De-novo identification

Structure Generator



Structure Generator Results



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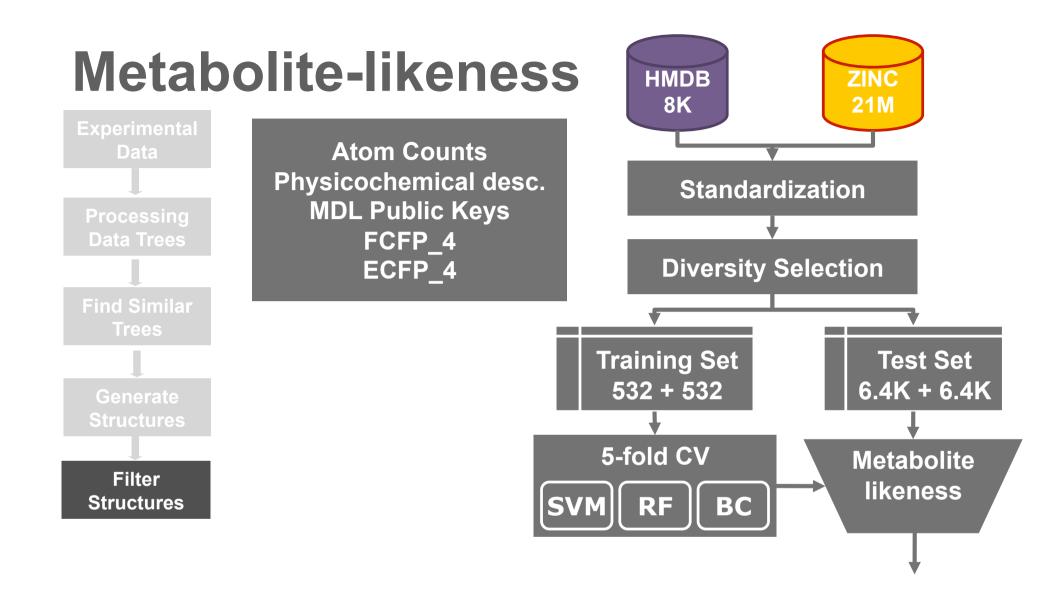
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In collaboration with Jean-Loup Faulon, Evry University

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MOLGEN same # <u>of</u>

Lots of candidates structures

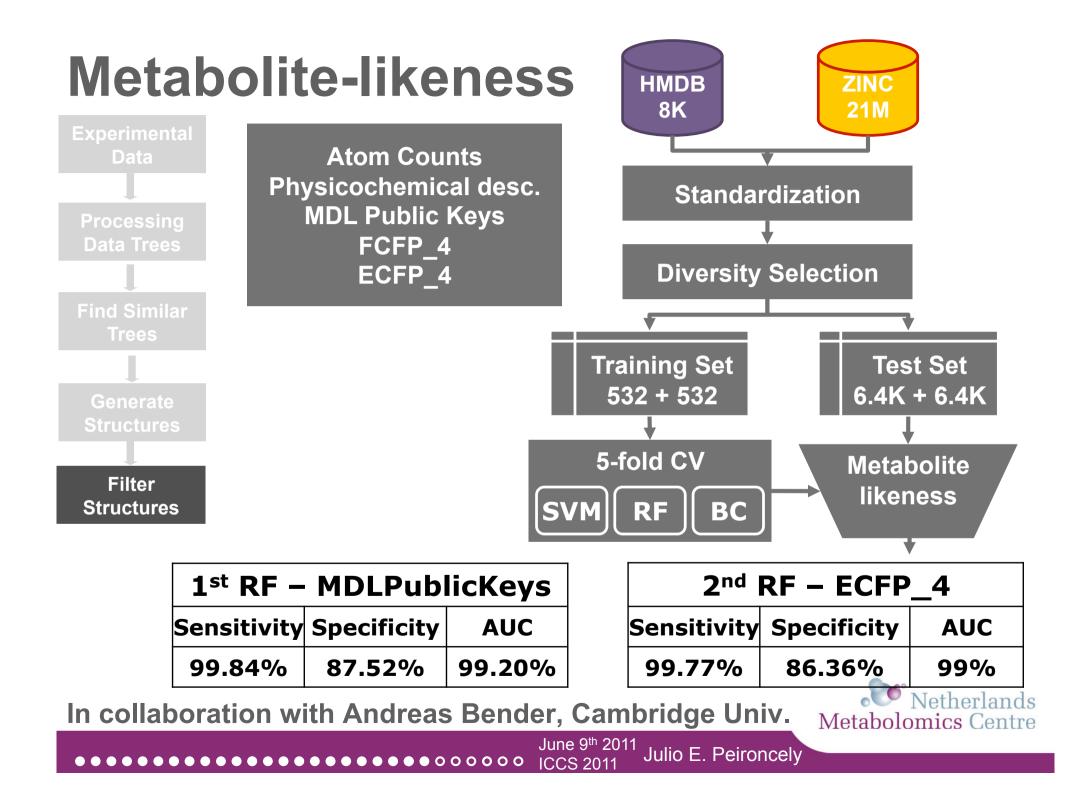


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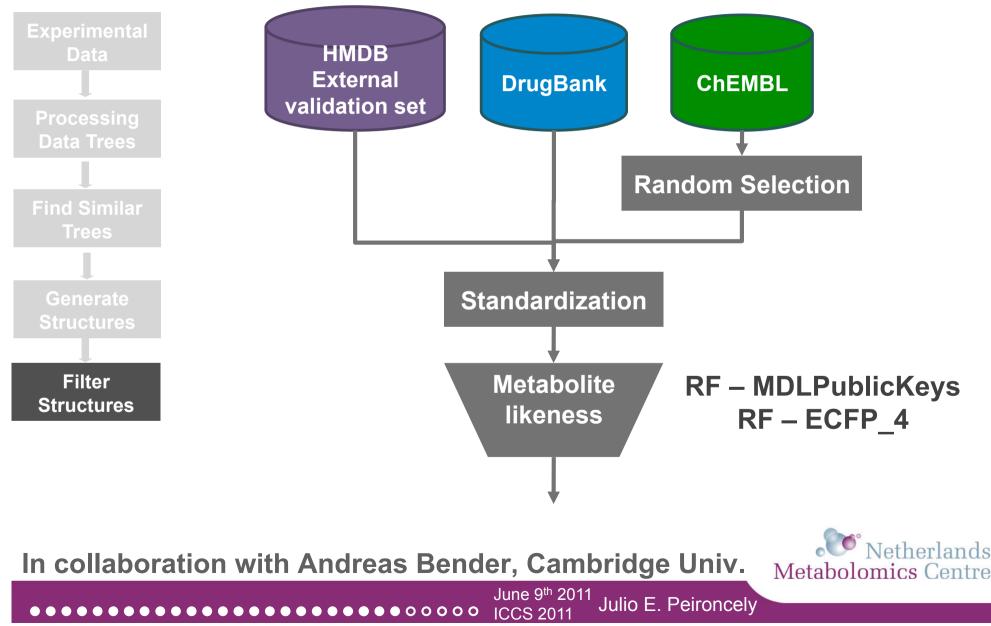
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In collaboration with Andreas Bender, Cambridge Univ.





Metabolite-likeness, external validation



Metabolite-likeness, external validation 3 ChEMBL DrugBank HMDB unofficial Data **Data Trees** 2 Density 1 **Structures** Filter **Structures** 0 0.2 0.4 0.6 0.8 Metabolite-likeness • Netherlands In collaboration with Andreas Bender, Cambridge Univ. **Metabolomics** Centre June 9th 2011 Julio E. Peironcely $\bullet \bullet \circ \circ \circ \circ$ **ICCS 2011**

From information to knowledge

www.MetiTree.nl

still in beta. not suited for production work.

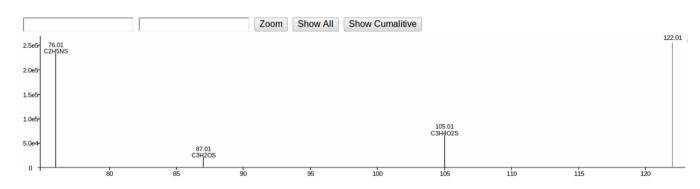


menu

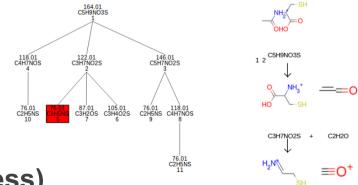
msn data process database(s) query by file query by text

configuration usergroup(s) member(s) api (test page) database viewer home > viewer > index

<<< BACK: Post JSON formatted Tree Structure



Group Database Upload and organize your own trees Visualize trees Find similar trees (to be added: Struct. Gen, Met-likeness)



C2H5NS + CO H2O

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welcome ICCS 2011 :: NMC (Netherlands Metabolomics Centre)

Conclusions Chemoinformatics plays a crucial role in the metabolite identification pipeline

Now it is the time to challenge this pipeline with real cases

Expert is still needed

Julio E. Peironcelv



Acknowledgements



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Leon Coulier Albert Tas



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HMP University of Alberta David Wishart Ying (Edison) Dong

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