

# The Chemistry Development Kit in the Netherlands

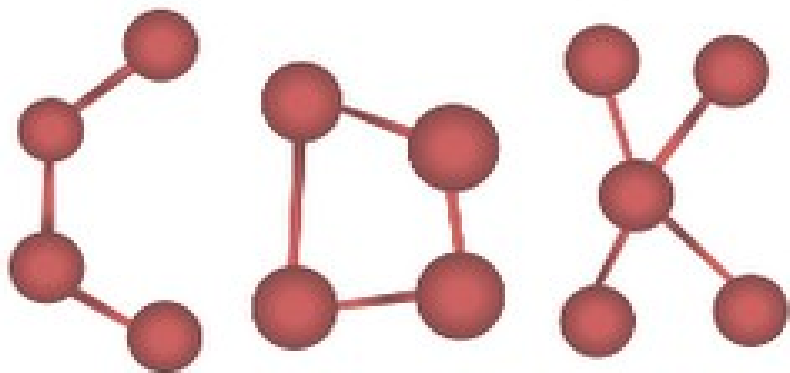
Egon Willighagen & Miguel Rojas-Chertó

BiGCaT - Maastricht University

Div. Anal. Biosciences – Leiden University

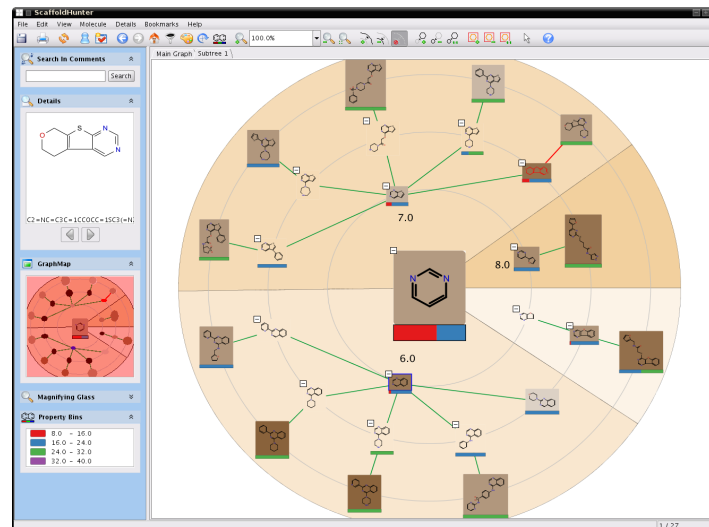
*NBIC 2012, Lunteren*

# Open Source Cheminformatics

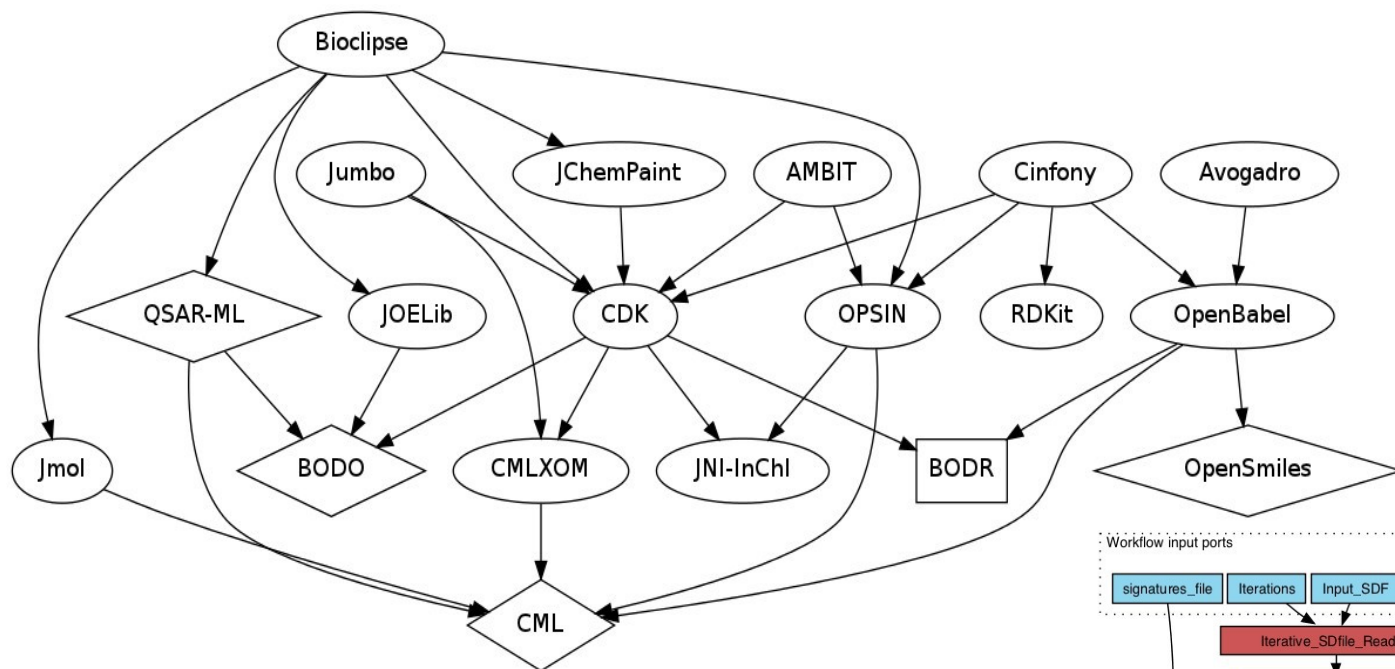


- Metabolomics
- Drug discovery
- Toxicology

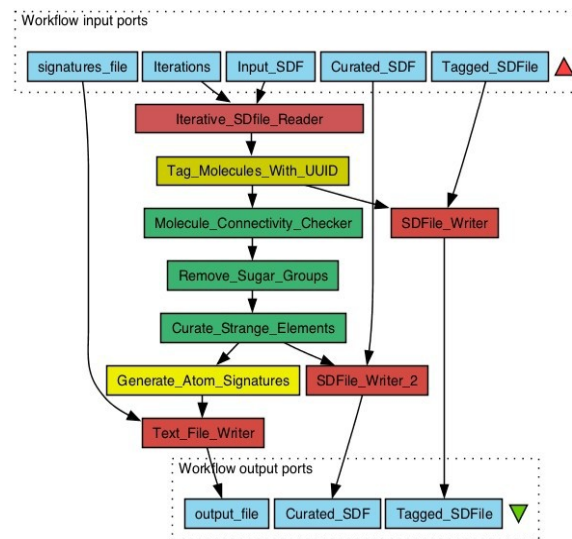
JOURNAL OF CHEMICAL INFORMATION AND MODELING	29	18.710 %	■
JOURNAL OF CHEMINFORMATICS	15	9.677 %	■
BMC BIOINFORMATICS	13	8.387 %	■
BIOINFORMATICS	10	6.452 %	■
JOURNAL OF CHEMICAL INFORMATION AND COMPUTER SCIENCES	6	3.871 %	■
JOURNAL OF COMPUTER AIDED MOLECULAR DESIGN	5	3.226 %	■
JOURNAL OF MOLECULAR BIOLOGY	4	2.581 %	■
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ACTA CHIMICA SINICA	3	1.935 %	■
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EUROPEAN JOURNAL OF MEDICINAL CHEMISTRY	2	1.290 %	■
EXPERT OPINION ON DRUG DISCOVERY	2	1.290 %	■



Wetzel et al., Nature Chemical Biology, 2009



O'Boyle et al., JChemInf, 2011



Truszkowski et al., JChemInf, 2011

# History

- Based on Steinbeck's CompChem, 1997
- CML support added in 1999
- Combining JChemPaint and Jmol in 2000
  - meeting @ Notre Dame University
  - funding by Specs & BioSpecs (D. Wife) !
- Now led by Steinbeck (EBI), Guha (NIH), me (BiGCaT)

Date: Tue, 04 May 1999 12:35:10 +0200  
From: Christoph Steinbeck  
To: Egon Willighagen

> Egon Willighagen wrote:  
>  
> Dear Christoph Steinbeck,  
>  
> Yesterday I visited your site on JChemPaint. I like to contribute some  
> of my expertise on Java and CML (1).  
>  
> CML is a markup language that is able to contain chemical information.  
> It can contain for example physical properties, for which I use CML in  
> my Dictionary on Organic Chemistry (2).  
> But is also might contain spectra, bibliographic references etc. And  
> of course 2D and 3D structural information.  
>  
> Therefore I propose to write both CML-input and -output procedures for  
> the JChemPaint project.  
>  
> I hope to hear from you soon.  
>  
> Yours sincerely,  
>  
> Egon Willighagen  
>  
> 1. <http://www.xml-cml.org/>  
> 2. <http://www.sci.kun.nl/sigma/Chemisch/Woordenboek/>

Dear Egon,

thanks very much for your mail and your offer to write CML-input and output routines for JChemPaint. That really sounds great to me and I will give you access to our CVS tree as soon as we have discussed the details.

Cheers,

Chris

--C. S.

Dr. Christoph Steinbeck (<http://www.ice.mpg.de/~stein>)



## E. CDK Authors

In acknowledgment to all the work done by the many people who have contributed to the success of the CDK, here is a list of those who wrote smaller or larger parts of the CDK library:

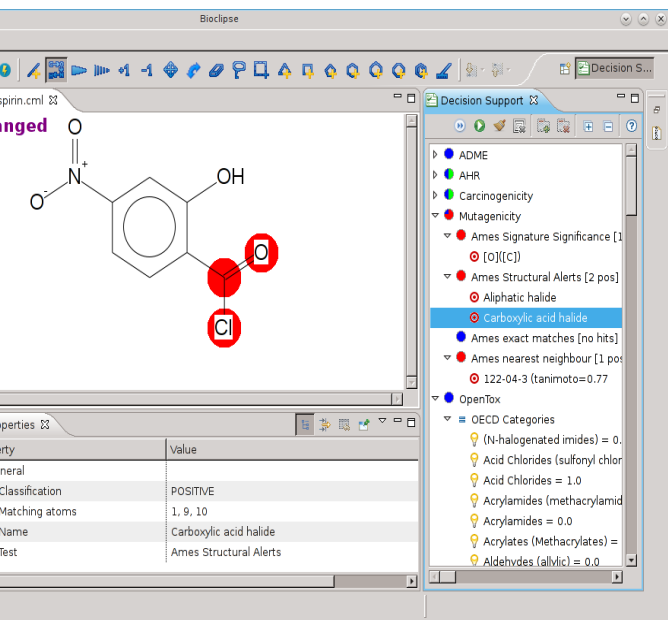
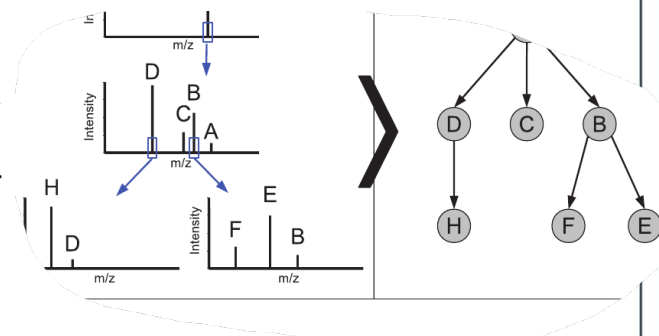
Sam Adams, Jonathan Alvarsson, Rich Apodaca, Saravanaraj N Ayyampalayam, Ulrich Bauer, Stephan Beisken, Arvid Berg, Ed Cannon, Fabian Dortu, Martin Eklund, Matteo Floris, Dan Gezelter, Uli Fechner, Thorsten Flügel, Nimish Gopal, Rajarshi Guha, Yonquan Han, Thierry Hanser, Kai Hartmann, Tobias Helmus, Christian Hoppe, Oliver Horlecher, Miguel Howard, Nina Jeliaskova, Geert Josten, Dmitry Katsubo, Jules Kerssemaekers, Anatoli Krassavine, Stefan Kuhn, Uli Köhler, Violeta Labarta, Jonty Lawson, Daniel Leidert, Edgar Luttmann, Todd Martin, John May, Nathanaël Mazuir, Stephan Michels, Scooter Morris, Peter Murray-Rust, Carl Mäsak, Irilenia Nobeli, Peter Odéus, Niels Out, Jerome Pansand, Julio Peironcely, Chris Pudney, Syed Asad Rahman, Jonathan Rienstra-Kiracofe, Mark Rijnbeek, David Robinson, Miguel Rojas Cherto, Bhupinder Sandhu, Jean-Sebastien Senecal, Onkar Shinde, Sulev Sild, Bradley Smith, Ola Spjuth, Christoph Steinbeck, Aleksey Tarkhov, Stephan Tomkinson, Gillean Torrance, Andrej Trzaskowski, Paul Turner, Jörg Wegner, Yap Chun Wei, Stephane Werner, Egon Willighagen, Lars Willighagen, Yong Zhang, and Daniel Zaharevitz.

## Elemental composition determination based on MS<sup>n</sup>

Miguel Rojas-Chertó<sup>1,2,\*</sup>, Piotr T. Kasper<sup>1,2</sup>, Egon L. Willighagen<sup>1,3,4</sup>, Rob J. Vreeken<sup>1,2</sup>, Thomas Hankemeier<sup>1,2</sup> and Theo H. Reijmers<sup>1,2,\*</sup>

<sup>1</sup>Netherlands Metabolomics Centre, <sup>2</sup>Division of Analytical Biosciences, Leiden/Amsterdam Center for Drug Research, Leiden, The Netherlands, <sup>3</sup>Division of Molecular Toxicology, Institute of Environmental Medicine, Karolinska Institutet, Stockholm, Sweden and <sup>4</sup>Plant Research International, Wageningen UR, Wageningen, The Netherlands

Associate Editor: Anna Tramontano



Willighagen *et al.* *BMC Research Notes* 2011, **4**:487  
<http://www.biomedcentral.com/1756-0500/4/487>

### SHORT REPORT

Open Access

## Computational toxicology using the OpenTox application programming interface and Bioclipse

Egon L Willighagen<sup>1,2\*</sup>, Nina Jeliakova<sup>3</sup>, Barry Hardy<sup>4</sup>, Roland C Grafström<sup>2,5</sup> and Ola Spjuth<sup>1</sup>

# Take home tweet

World-class  
Open Source  
Cheminformatics  
here in NL

<http://cdk.sf.net/>

The screenshot shows a Google+ profile for 'Chemistry Development Kit'. The profile header includes the name, a search bar, and options to 'View as...' and 'Edit profile'. The profile picture is a molecular structure diagram. The main content area shows a post from 'Chemistry Development Kit' dated 19 Apr 2012, which is a public post. The post text reads: 'The editor functionality is with CDK and CDK-JChemPaint.' Below the text is a photo of a software interface. A quote from Ola Spjuth is included: 'Ola Spjuth originally shared this post: In the recent days I have implemented a Bioclipse Decision Support module for ChemSpider integrated with ChEMBL-RDF. See blog post below for more info.' To the right of the post, there is a link to 'cdk.sf.net/' and a '+104' reaction count. Below the post, there are sections for '13 IN YOUR CIRCLES' and '97 HAVE YOU IN CIRCLES', each with a row of profile pictures.