

The MetWare developers meeting in Halle



Published April 9, 2008

Citation

Willighagen, E. (2008, April 9). The MetWare developers meeting in Halle. *Chem-bla-ics*. <https://doi.org/10.59350/7tfeg-dvs86>

Keywords

Metware, Metabolomics

Abstract

Today starts the MetWare developers meeting, hosted by Steffen Neumann, at the Leibniz-Institut für Pflanzenbiochemie. Steffen's group and the Applied Bioinformatics group where I now work, are co-developing an opensource platform for metabolomics data management.

Copyright

Copyright © None 2008. Distributed under the terms of the [Creative Commons Attribution 4.0 International License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Today starts the [MetWare](#) developers meeting, hosted by [Steffen Neumann](#), at the [Leibniz-Institut für Pflanzenbiochemie](#). Steffen's group and the [Applied Bioinformatics](#) group where I now work, are co-developing an opensource platform for metabolomics data management. Not really a full LIMS system, but a system to keep track of all the facts about the experiments and samples we would use when analyzing the data in order to find new chemistry, biomarkers, etc (see [this earlier blog](#) too). Good news is, that BioAssist is developing a support platform for the [NMC](#), and plans to use MetWare as a main component.

OK, off to catch my train now. See you online ([#metware](#) @ [irc.freenode.net](#)); the wiki has an [agenda for the meeting](#).