

# Coffee Lectures: The 25<sup>th</sup> Series

Tue, Wed, Thu 13:00–13:10, live at HCI G2

and online via Zoom <https://ethz.zoom.us/j/63020946312>



**Tuesday, 22.3.22**

## Full text search in Google Books (No. 2)

You do not have time to read books to find out if a particular topic has been covered? Google Books can help you out – even in cases where no full text view is provided.

*Presenter: Andrej Kilian*

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[www.infozentrum.ethz.ch](http://www.infozentrum.ethz.ch)**

**Wednesday, 23.3.22**

## ChemSpider (No. 26)

There is more than just Reaxys and SciFinder<sup>®</sup>: ChemSpider is a freely available database with over 114 million structures, provided by the British Royal Society of Chemistry. Via an API, automated queries are possible.

*Presenter: Dr. Leo Betschart*

**Thursday, 24.3.22**

## Navigating your way through EMBL-EBI (No. 52)

New initiatives, data repositories, on-demand training – discover a slice of what the EMBL's European Bioinformatics Institute has to offer.

*Presenter: Dr. Gina Cannarozzi*



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Collector's Cards!**

**Tuesday, 29.3.22**

## Markush structure searching (No. 48)

How do you make sure that your new chemical compound is not already covered by an existing patent? Markush structures are generic descriptions of chemical structures and used in chemical patents worldwide. We will show how to search for them in Reaxys and SciFinder.

*Presenter: Dr. Leo Betschart*

**Wednesday, 30.3.22**

## WordPress blog software for information management (No. 15)

Using the WordPress service of ETH Zurich, you can set up websites or journal clubs easily. The Information Center presents its own practical application examples.

*Presenter: Dr. Oliver Renn*

**Thursday, 31.3.22**

## LaTeX made simple with Overleaf (No. 68)

Overleaf is the most popular online LaTeX collaborative editor. It comes with a ready to go LaTeX environment and all you need to start writing is a web browser

*Presenter: Dr. Jozica Dolenc*

**Tuesday, 5.4.22**

## KnowItAll Informatics System Spectroscopy Software (No. 6)

Get to know the benefits of the KnowItAll software solution for spectral analysis, identification and search over the commonly used KnowItAll AnyWare web interface.

*Presenter: Dr. Jozica Dolenc*

**Wednesday, 6.4.22**

## Use the full power of Scopus (No. 17)

Scopus offers many options to expand literature searches, to make them more specific and to analyze the results. We will show you when it is useful to do the extra click.

*Presenter: Andrej Kilian*

**Thursday, 7.4.22**

## Alpha Fold (No. 61)

AlphaFold, an AI system for predicting protein structure, has had unprecedented success. We present an overview of the software and the database of structures it has predicted, both of which are open access.

*Presenter: Dr. Gina Cannarozzi*