

## **A CLOUD of possibilities: Finding new therapies by combining drugs**

**The CeMM Library of Unique Drugs (CLOUD) is the first condensed set of FDA-approved drugs representing the entire target and chemical space of all clinical compounds. Its potential was shown in a combinatorial high throughput screen at the CeMM chemical screening platform, published in Nature Chemical Biology: by testing all CLOUD compounds in combination with each other, a pair of hitherto unrelated drugs proved to be highly effective against multiple prostate cancer cell lines known for their resistance to therapy. Testing CLOUD combinations in this highly automated procedure could pave the way for a new era of drug repurposing and provide novel strategies for personalized medicine.**

(Vienna, May 22, 2017) Two drugs taken together can sometimes lead to outcomes that largely deviate from the effect of the separated compounds – a fact well known from warnings on patient information leaflets. However, while doctors strongly advise against unsupervised mixing of drugs, the synergy of two combined pharmaceuticals assessed in an experimental setting can reveal completely new therapeutic options. Nevertheless, finding a novel combination of drugs for a given disease within the more than 30,000 drug products approved by the regulatory agencies was hitherto a big challenge for scientists.

To facilitate systematic screening for synergistic interactions of drugs, CeMM PI Stefan Kubicek and his colleagues established a collection of 308 compounds (CeMM Library of Unique Drugs, CLOUD) that effectively represent the diversity of structures and molecular targets of all FDA-approved chemical entities. Moreover, the scientists proved the potential of the CLOUD with CeMM's highly automated chemical screening platform by identifying a novel synergistic effect of two drugs (flutamide and phenprocoumon (PPC)) on prostate cancer cells. The results of Kubicek's team with Marco Licciardello as first author were published in Nature Chemical Biology ([DOI:10.1038/nchembio.2382](https://doi.org/10.1038/nchembio.2382))

For the establishment of the CLOUD, a clever series of condensation steps was necessary: the CeMM scientists first determined and extracted 2171 unique active pharmaceutical ingredients from the database, discarding all products with identical compounds. Next, they removed large macromolecules like antibodies as well as salt fragments, and discarded all molecules that exert their biological effects through mechanisms other than protein-ligand interactions, are not used to treat diseases or are found only in topical products.

With the remaining 954 systemically active small molecules (STEAM collection), the work had just begun: in order to create a comprehensive collection of compounds that fits on a standard 384-well screening plate, the researchers appended biological activities to all drugs with known molecular targets and grouped them into 176 classes of similar structure and activity. With a sophisticated clustering algorithm, 239 representative drugs were selected from those classes. Combined with 34 drugs with unknown target and 35 active forms of prodrugs (that otherwise need to be metabolized to become active), 308 compounds were selected in total for the CLOUD - the world's first library representing all FDA-approved chemical entities including the active form of prodrugs.

To put the combinatorial screen with the CLOUD to the test, Kubicek's group investigated the effect of pairwise combinations of CLOUD compounds on the viability on KBM7 leukemia cells, a cell line well suited for drug experiments. Using a dose chosen for each compound individually based on the clinically relevant maximum plasma concentration, the scientists found a strong synergistic interaction between flutamide, a drug approved for the treatment of prostate cancer, and phenprocoumon (PPC), an anti-thrombosis compound. In combination, flutamide and PPC efficiently killed the cancer cells.

After identifying the androgen receptor (AR) as molecular target of the synergistic interaction, the scientists tried the drug combination on prostate cancer cells known to be hard to treat - and hit the bulls eye. "The combination induced massive cell death in prostate cancer cells. We then went back to the entire approved drug list, and indeed, we could show that all drugs from the clusters that flutamide and phenprocoumon represent synergize. Thereby we validated the reductionist concept underlying the CLOUD library," Stefan Kubicek explains.

With their experiments, Kubicek's team in collaboration with scientists from the Medical University of Vienna, the Uppsala University, Enamine Kiev and the Max Planck Institute for Informatics in Saarbrücken proved that the CLOUD is the ideal set of compounds to develop screening assays and discover new applications for approved active ingredients. At CeMM, a number of key discoveries on new applications for approved drugs have already been made with the CLOUD. Furthermore, as shown in the current issue of Nature Chemical Biology, the CLOUD is ideal for finding new drug combinations. "In view of these successes, I would predict that this set of compounds will become world standard for all screening campaigns", Stefan Kubicek emphasizes.

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**Attached pictures:** 1) Schematic representation of the filtering and clustering procedure leading to the 308 CLOUD drugs (© Nature Chemical Biology / Stefan Kubicek), 2) Immunofluorescence analysis of prostate cancer cells treated with 15mM flutamide, 35 µM PPC or the combination for

24 h. Scale Bar 20  $\mu$ M (© Nature Chemical Biology / Stefan Kubicek) 3) Senior author Stefan Kubicek (© CeMM/Sazel)

**The study** “A combinatorial screen of the CLOUD uncovers a synergy targeting the androgen receptor” was published online in advance in *Nature Chemical Biology* on May 22, 2017. [DOI:10.1038/nchembio.2382](https://doi.org/10.1038/nchembio.2382)

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**Stefan Kubicek** studied organic chemistry in Vienna and Zürich. He received his Ph.D. in Thomas Jenuwein’s group at the Institute for Molecular Pathology (IMP) in Vienna followed by postdoctoral work with Stuart Schreiber at the Broad Institute of Harvard and MIT in the U.S. He joined CeMM in 2010. He is the Head of the Chemical Screening at CeMM and the Platform Austria for Chemical Biology (PLACEBO) and the Christian Doppler Laboratory for Chemical Epigenetics and Anti-Infectives.

The mission of **CeMM Research Center for Molecular Medicine of the Austrian Academy of Sciences** is to achieve maximum scientific innovation in molecular medicine to improve healthcare. At CeMM, an international and creative team of scientists and medical doctors pursues free-minded basic life science research in a large and vibrant hospital environment of outstanding medical tradition and practice. CeMM’s research is based on post-genomic technologies and focuses on societally important diseases, such as immune disorders and infections, cancer and metabolic disorders. CeMM operates in a unique mode of super-cooperation, connecting biology with medicine, experiments with computation, discovery with translation, and science with society and the arts. The goal of CeMM is to pioneer the science that nurtures the precise, personalized, predictive and preventive medicine of the future. CeMM trains a modern blend of biomedical scientists and is located at the campus of the General Hospital and the Medical University of Vienna.

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