

A new molecular toolkit for the de-novo design of bioactive agents

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Taxadien synthase with the substrate geranylgeranyldiphosphat in the active center of the enzyme. The green dots mark the catalytically relevant Mg2+-ions, which are involved in the initial hydrolysis of the phosphate residue. Credit: Max Hirte / TUM



Nature provides mankind with a wide variety of valuable bioactive agents ranging from vitamins over vital fatty acids to cancer inhibiting substances. Many of these substances are difficult to obtain directly from the environment or can not be produced effectively by chemical total synthesis.

Scientists of the Technical University of Munich (TUM) are taking a new approach: Using synthetic biotechnology methodologies they have developed a biochemical strategy to synthesize natural and completely artificial medical agents by a templated enzyme design process. First products, a precursor of the anti-cancer medicament Taxol, anti-inflammatory substances and omega-3 <u>fatty acids</u> prove the successfulness of their strategy.

The bark of the pacific yew tree (*Taxus brevifolia*), for example, contains Taxol, an agent that is used in the treatment of breast, ovary and lung cancers. Unfortunately, this yew species is not widespread and it is protected.

Essential <u>omega-3 fatty acids</u>, a component of infant nutrition, for example, are currently produced predominantly from fish and crustaceans - placing a further pressure on the already strongly affected marine ecosystem.

The aim of the workgroup led by Thomas Brück, professor for Industrial Biocatalysis at the Technical University of Munich, is thus to obtain <u>chemical substances</u> in a sustainable manner on an industrial scale using methods of biochemistry, bioinformatics and biotechnology.

"Gold" from straw - a yeast with great potential

Now Brück and his team have succeeded in genetically altering the hitherto biotechnologically unexploited *Trichosporon oleaginosus* yeast



by coercing it to produce essential omega 3 fatty acids alpha-linoleic acid (ALA), eicosapentaenoic acid (EPA) and the anti-inflammatory conjugated linoleic acid (CLA).



Coautor Felix Bracharz preparing a high-throughput nile red assay. Analysis of the fluorescence (see computer screen in the background) allows the evaluation of the lipid content of the cells. Credit: Andreas Battenberg / TUM

The yeast can thrive cultivation media derived from agricultural waste, including straw, wood chips, wheat bran, and even hitherto unused marine waste materials like crab shells. "This yeast is quite unique



because it can also exploit monomer sugar substances, which are normally difficult to metabolize," explains Brück. "We thus obtain valuable chemical substances from waste - without harming the environment."

When Trichosporon oleaginosus cells become stressed in nature, from a shortage of nitrogen or phosphate, for example, they build up energy reserves in the form of fat. Even though the yeast cells no longer grow optimally, the resulting triglyceride fat reserves make up as much as 70 percent of their dry weight.

In future projects the researchers led by Brück hope to further modify the oil producing yeast so that it produces the desired fats in adequate quantities, even under normal nutrient conditions and without limiting their growth.

From simulations to custom-tailored enzymes

A methodology recently presented by Brück's research group in the highly ranked scientific journal *Proceedings of the National Academy of Sciences (PNAS)* takes the idea one step further: Molecular-mechanical computer simulations allow them to decipher the individual steps in which a specific class of enzymes produces bioactive natural products. These include precursor stages of the cancer medication Taxol.

Employing only computer simulation, Brück and his team correctly predicted, for the first time ever, all intermediate stages in the complex cascade of reactions that take place in the presence of this enzyme. In this manner they elucidated the enzyme's precise mode of action, as well as the relationship between its structure and function. This was previously not possible using classical biochemical methodologies.

"This approach is very promising because using computer simulation we



can change enzymes in a targeted manner and predict which products will be produced as a result," says Brück. "If we then interconnect different enzyme activities with each other, we can even create new molecules that are not found in nature."



Trichosporon oleaginosus yeast cells. Staining with nile red makes the lipid droplets in the cells observable. Credit: Veronika Redai / TUM

43-fold increase of product yield

The production of cyclooctatin, a potent anti-inflammatory agent,



provides a good example: The scientists coupled a diterpene synthase enzyme with a new hydroxylase/reductase enzyme complex in an Escherichia coli-based microbial production system provided for the production of trihydroxylated diterpene cyclooctatin.

It was the in-silico discovery and experimental verification of a new reductase/ferredoxin system derived from the recently published genome of Streptomyces afghaniensis, which enabled the scientists to raise the product yield by a factor of 43, compared with the native producer.

In the future, biotechnologists could adopt approaches similar to those deployed by engineers when they develop production steps for a new automobile. Making use of the synthetic biotechnology insight, they could put together a synthesis method for new active agents using a chain of reactions with modified enzymes. This would significantly reduce the long and arduous process of "puzzling out" new synthesis routes in the laboratory, as commonly applied today.

More information: Christian Görner et al, Identification, characterization and molecular adaptation of class I redox systems for the production of hydroxylated diterpenoids, *Microbial Cell Factories* (2016). DOI: 10.1186/s12934-016-0487-6

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