

4. Protein Engineering

Outline

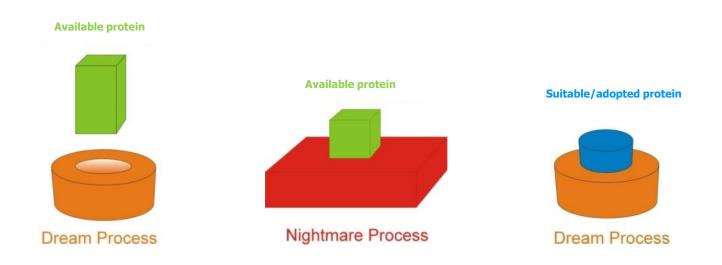
- ☐ Limitations of proteins in biotechnology processes
- ☐ Definition and aim of protein engineering
- Targeted properties of proteins
- Basic approaches in protein engineering
 - DIRECTED EVOLUTION
 - RATIONAL DESIGN
 - SEMI-RATIONAL DESIGN
- Examples

Proteins in biotechnology

availability of optimal protein for specific process

HOW TO OBTAIN OPTIMAL PROTEIN?

- traditional biotechnology adapt process
- modern biotechnology adapt protein



Proteins in biotechnology

- classical screening
 - screening culture collections
 - polluted and extreme environment
- environmental gene libraries
 - metagenomic DNA
- data-base mining
 - gene databases
 - genome sequencing projects
 - numerous uncharacterised enzymes/proteins





IF SUITABLE PROTEIN DOES NOT EXIST IN NATURE?

PROTEIN ENGINEERING

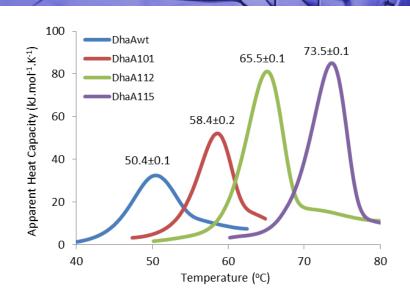
Proteins in biotechnology

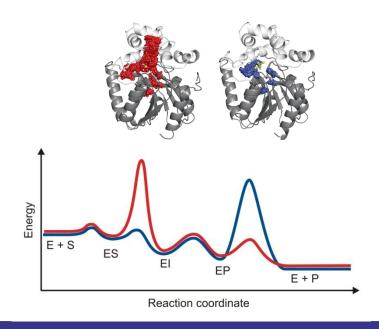
- the process of **constructing novel protein** molecules by design first principles or altering existing structure
- use of genetic manipulations to alter the coding sequence of a gene and thus modify the properties of the protein
- AIMS AND APPLICATIONS
 - technological optimisation of the protein to be suitable in particular technology purpose
 - scientific desire to understand what elements of proteins contribute to folding, stability and function

Targeted properties of proteins

- □ **structural** properties of proteins
 - stability (temperature, solvents)
 - tolerance to pH, salt
 - resistance to oxidative stress

- functional properties of proteins
 - reaction type
 - substrate specificity and selectivity
 - kinetic properties (e.g., $K_{\rm m}$, $k_{\rm cat}$, $K_{\rm i}$)
 - cofactor selectivity
 - protein-protein or protein-DNA interactions





RATIONAL DESIGN

1. Computer aided design



2. Site-directed mutagenesis



Individual mutated gene

- 3. Transformation
 - 4. Protein expression
 - 5. Protein purification
 - 6. not applied

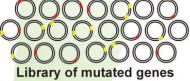


Improved protein

DIRECTED EVOLUTION

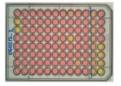
1. not applied

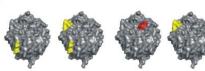
2. Random mutagenesis



Library of mutated genes (>10,000 clones)

- 3. Transformation
- 4. Protein expression
- 5. not applied
- 6. Screening and selection
 - stability
 - selectivity
 - affinity
 - activity





Selected mutant enzymes



Constructed mutant enzyme

7. Biochemical testing

Directed evolution

- directed evolution techniques emerged during mid-1990s
- inspired by natural evolution
- this form of "evolution" does not match what Darwin had envisioned
 - requires outside intelligence, not blind chance
 - does not create brand new species, macroevolution,
 but only improvements of molecules, molecular evolution
 - does not take millions of years, but happens rapidly

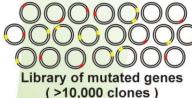
Directed evolution

- evolution in test tube comprises two steps
 - random mutagenesismutant library building
 - screening and selection
 identification of desired biocatalyst
- prerequisites for directed evolution
 - gene encoding protein of interest
 - method to create mutant library
 - suitable expression system
 - screening or selection system

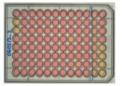


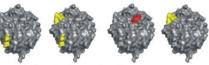
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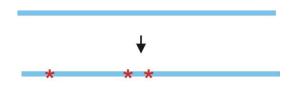
7. Biochemical testing

Selected mutant enzymes

Methods to create mutant libraries

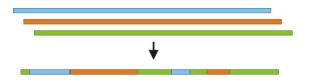
- □ technology to generate large diversity
 - NON-RECOMBINING

one parent gene -> variants with point mutations



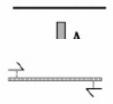
RECOMBINING

several parental homologous genes -> chimeras



Non-recombining mutagenesis

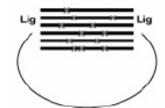
- UV irradiation or chemical mutagens (traditional)
- mutator strains lacks DNA repair mechanism mutations during replication (e.g., Epicurian coli XL1-Red)
- error-prone polymerase chain reaction (ep-PCR)
 - gene amplified in imperfect copying process
 (e.g., unbalanced deoxyribonucleotides concentrations,
 high Mg²⁺ concentration, Mn²⁺, low annealing temperatures)
 - 1 to 20 mutation per 1000 base pairs
- saturation mutagenesis
 - randomization of single or multiple codons
- other methods
 - gene site saturation mutagenesis
 - cassette mutagenesis (region mutagenesis)





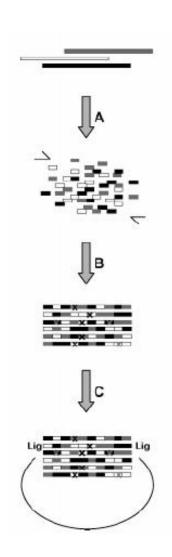






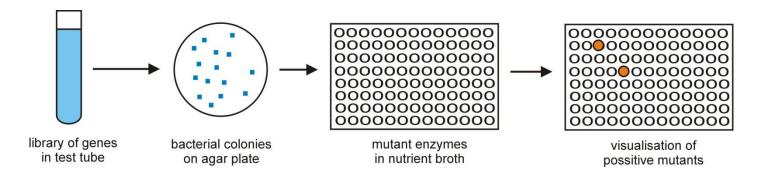
Recombining mutagenesis

- also refered to as "sexual mutagenesis"
- DNA shuffling
 - fragmentation step
 - random reassembly of segments
- □ StEP staggered extension process
 - simpler then shuffling
 - random reannealing combined with limited primer extension
- other methods
 shuffling of genes with lower homology down to 70%
 (e.g., RACHITT, ITCHY, SCRATCHY)



Screening and selection

- most critical step of direct evolution
- ☐ isolation of positive mutants hiding in library
 - HIGH THROUGHPUT SCREENING
 individual assays of variants one by one
 - DIRECT SELECTION
 display techniques (link between genotype and phenotype)

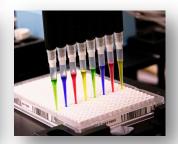


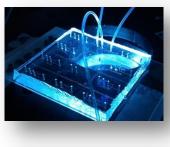
(Utra) High throughput screening

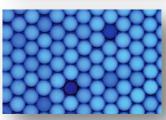
- common methods not applicable
- □ agar plate (pre)screening
- microtiter plates screening
 - 96-, 384- or 1536-well formate
 - robot assistance (colony picker, liquid handler)
 - 10⁴ libraries
 - volume 10 100 uL
- microfluidic systems
 - water in oil emulsions (up to 10 kHz)
 - FACS sorting (10⁸ events/hour)
 - 10⁹ libraries
 - volume 1 10 pL

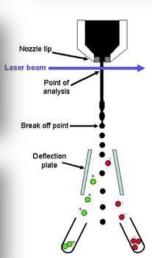








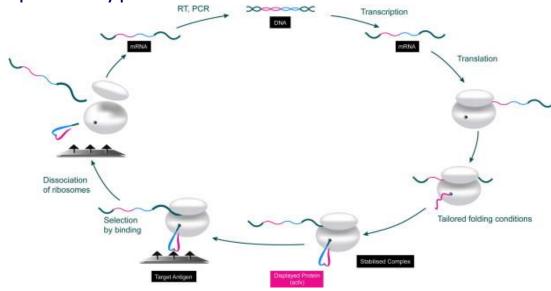




Direct selection

- □ not generally applicable (mutant libraries >10 variants)
- ☐ link between genotype and phenotype
- display technologies
 - ribosome display
 - phage display
- life-or-death assay
 - auxotrophic strain
 - toxicity based selection

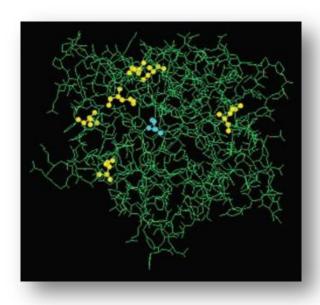


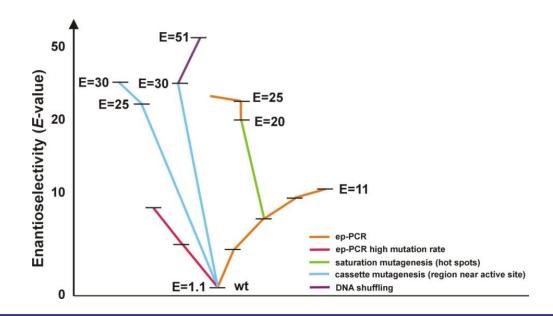




Example of Directed evolution

- directed evolution of enantioselectivity
 - lipase from P. aeruginosa (E-value improved from 1.1 into 51)
 - spectrophotometric screening of (R)- and (S)-nitrophenyl esters
 - 40 000 variants screened
 - the best mutant contains six amino acid substitutions





RATIONAL DESIGN

1. Computer aided design

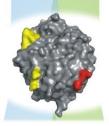


2. Site-directed mutagenesis



Individual mutated gene

- 3. Transformation
 - 4. Protein expression
 - 5. Protein purification
 - 6. not applied



Improved protein

DIRECTED EVOLUTION

1. not applied





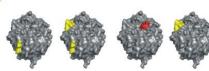
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6. Screening and selection

- stability
- selectivity
- affinity
- activity





Selected mutant enzymes



Constructed mutant enzyme

7. Biochemical testing

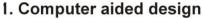
Rational design

- emerged around 1980s as the original protein engineering approach
- knowledge based combining theory and experiment
- protein engineering cycle:

"structure-theory-design-mutation-purification-analysis"

- difficulty in prediction of mutation effects on protein property
- de novo design

Principal of rational design



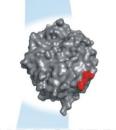


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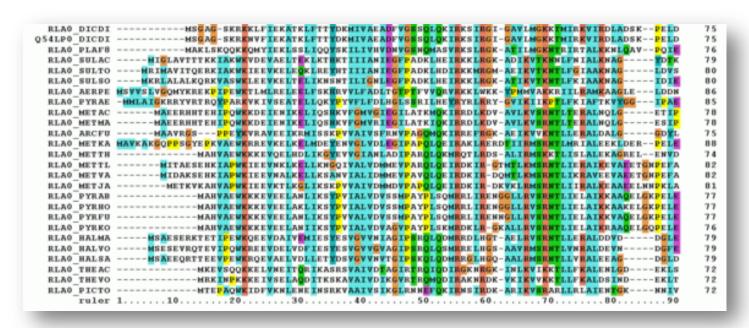
7. Biochemical testing

- rational design comprises:
 - design understanding of protein functionality
 - experiment construction and testing of mutants
- prerequisites for rational design:
 - gene encoding protein of interest
 - 3D structure (e.g., X-ray, NMR)
 - structure-function relationship
 - computational methods and capacity
 - (multi)side directed mutagenesis techniques
 - efficient expression system
 - biochemical tests

Design



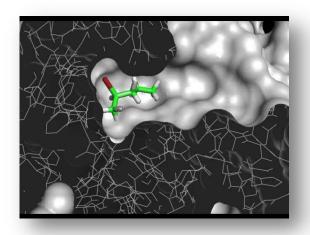
- homologous wild-type sequences are collected and compared
- identifying amino acid residues responsible for differences
- reconstruction transfer differences from one enzyme to another
- new design combination of possitive mutation from all parental proteins in one construct, new protein better than all parental

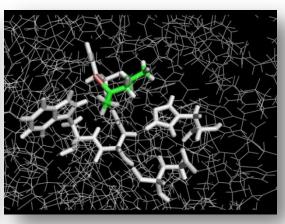


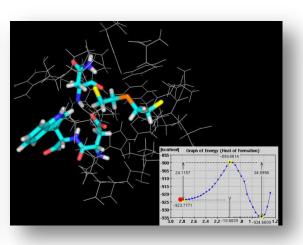
Design

STRUCTURE-BASED APPROACH

- prediction of enzyme function from structure alone is challenging
- protein structure (X-ray crystallography, NMR, homology models)
- molecular modelling
 - molecular docking
 - molecular dynamics
 - quantum mechanics/molecular mechanics (QM/MM)

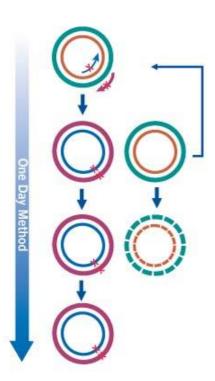






Construction

- ☐ site-directed mutagenesis
 - introducing point mutations
- multi site-directed mutagenesis
- gene synthesis
 - commercial service
 - codone optimisation

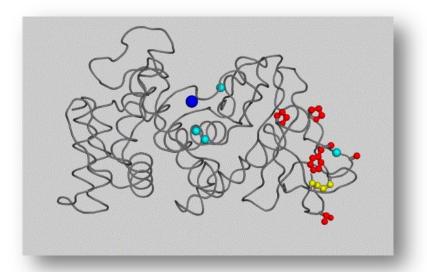






- rational design of protein stability
 - stability to high temperature, extreme pH, proteases etc.
 - stabilizing mutations increase strength of weak interactions
 - o salt bridges and H-bonds
 Eijsink et al., Biochem. J. 285: 625-628, 1992
 - S-S bonds
 Matsumura et al., Nature 342: 291-293, 1989
 - o addition of prolines
 Watanabe et al., Eur. J. Biochem. 226: 277-283, 1994
 - less glycines
 Margarit et al., Protein Eng. 5: 543-550, 1992
 - o oligomerisation
 Dalhus et al., J. Mol. Biol. 318: 707-721, 2002

- engineering protein to resist boiling
 - reduced rotational freedom
 Thr56Ala, Gly58Ala, Ser65Pro and Ala96Pro
 - introduction of disulfide bridge Gly8Cys + Asn60Cys
 - improved internal hydrogen bond Ala4Thr
 - filling cavity Tyr63Phe



Half-lifes (min.)	80°C	100°C
wild type	17.5	>0.5
8-fold mutant	stable	170

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7. Biochemical testing

DIRECTED EVOLUTION

1. not applied

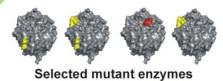


- 2. Random mutagenesis

 OOOOOO

 Library of mutated genes
 (>10,000 clones)
- 3. Transformation
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 - stability
 - selectivity
 - affinity
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7. Biochemical testing

RATIONAL DESIGN

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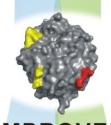


2. Site-directed mutagenesis



Individual mutated gene

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IMPROVED ENZYME

DIRECTED EVOLUTION

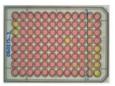
SEMIRATIONAL DESIGN

2. Random mutagenesis



Library of mutated genes (>10,000 clones)

- 3. Transformation
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 - stability
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Constructed mutant enzyme

7. Biochemical testing



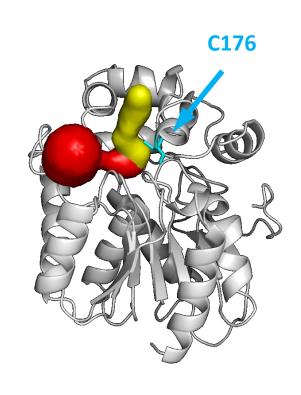


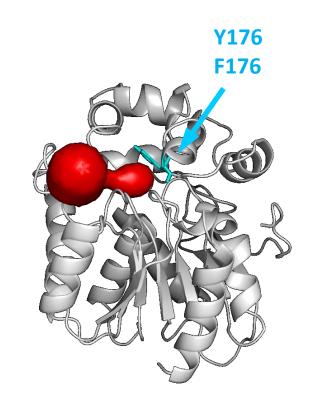




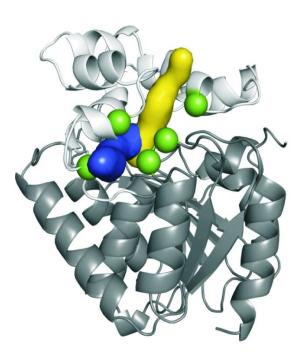
Selected mutant enzymes

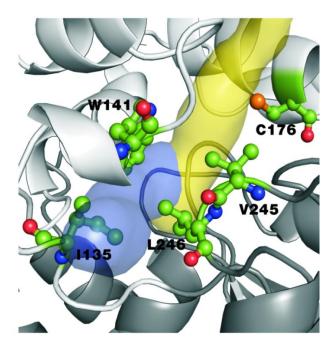
- conversion of 1,2,3-trichloropropane
 by DhaA from *Rhodococcus erythropolis* Y2
- □ **DIRECTED EVOLUTION** importance of access pathways

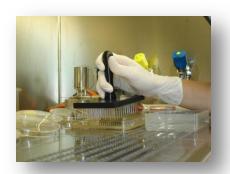


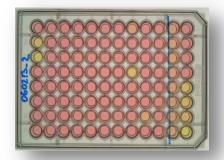


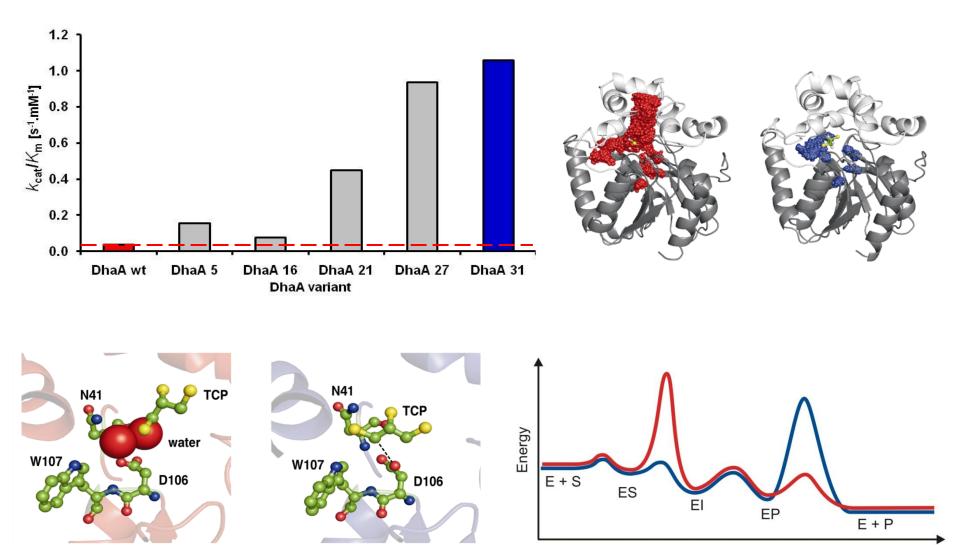
- conversion of 1,2,3-trichloropropane
 by DhaA from Rhodococcus erythropolis Y2
- □ **DIRECTED EVOLUTION** importance of access pathways
- SEMI-RATIONAL DESIGN hot spots in access tunels
- ☐ library of **5,300 clones** screened











Reaction coordinate