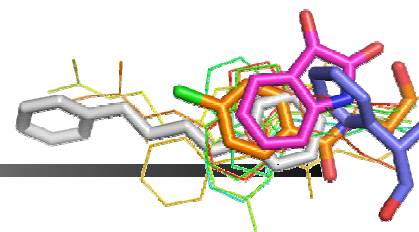


De Novo Design



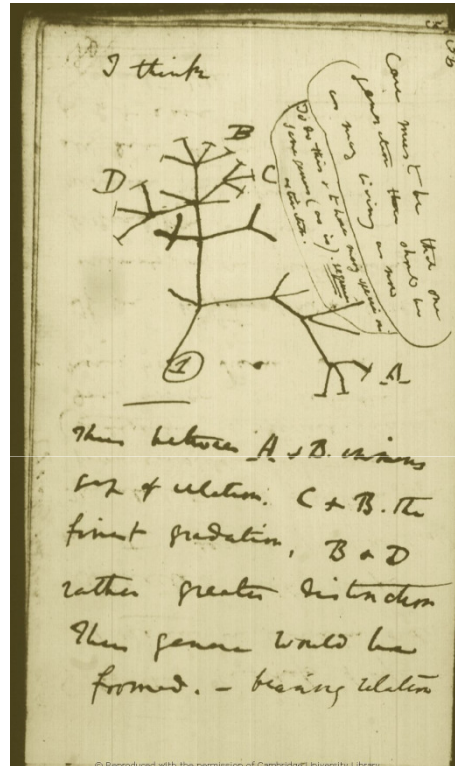
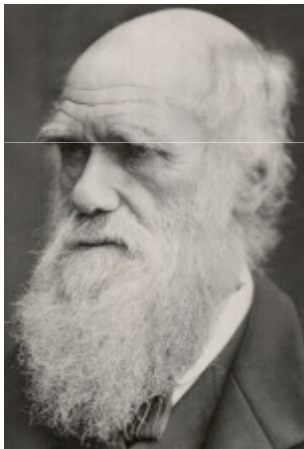
Adaptive Systems for Molecular Design

Gisbert Schneider

Beilstein Endowed Chair for Chem- & Bioinformatics

Goethe-University Frankfurt

Transmutation of Species



“[..] Thus between A & B immense gap of relation. C & B the finest gradation, B & D rather greater distinction. Thus genera would be formed. — bearing relation”

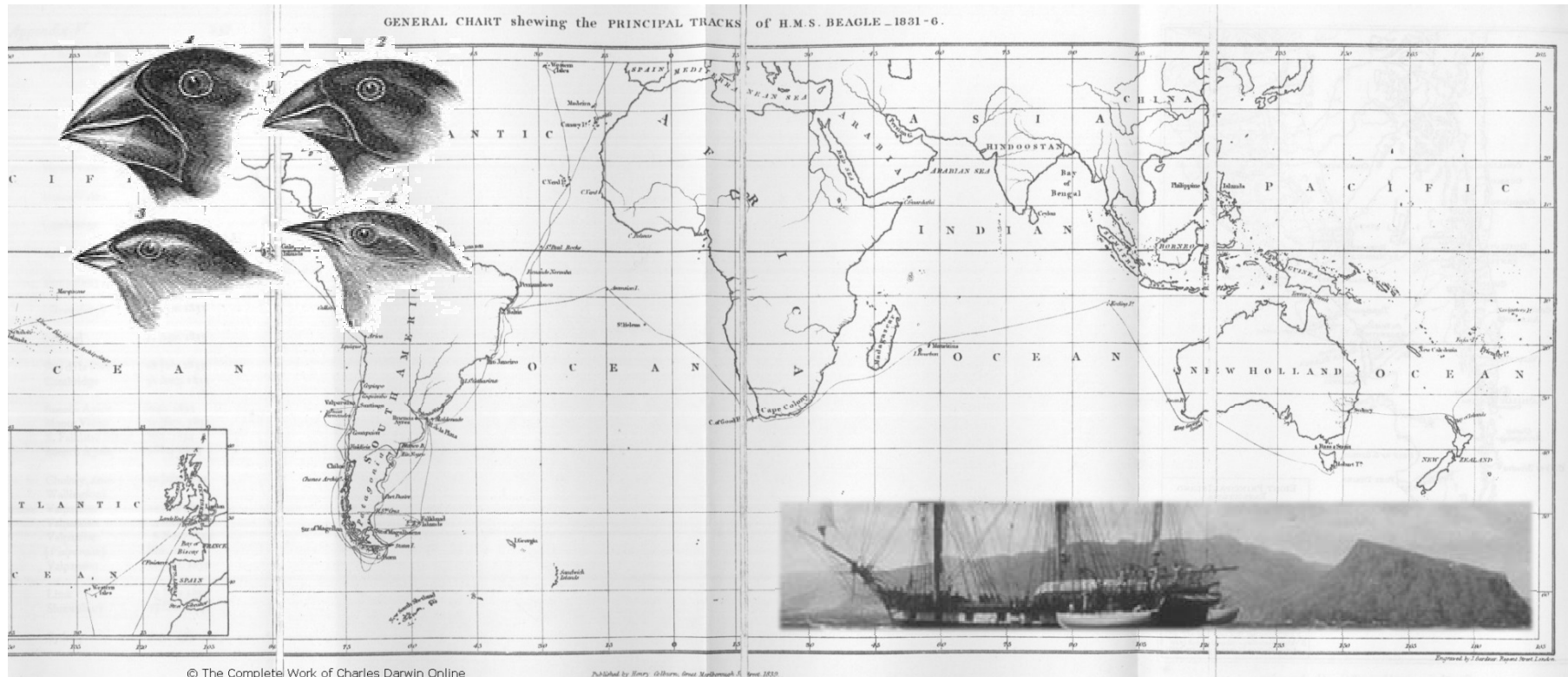
Charles R. Darwin (*1809)

Notebook B: *Transmutation of Species* (1837-1838), p.36

Source: www.darwin-online.org.uk

23 June 2008, (c) G. Schneider

Voyages to the (un)known



Darwin and Henslow, letters 1831-1860.

The growth of an idea (N. Barlow, ed. 1967).

Voyages of the H.M.S. *Beagle*

23 June 2008, (c) G. Schneider

De Novo Design Concepts

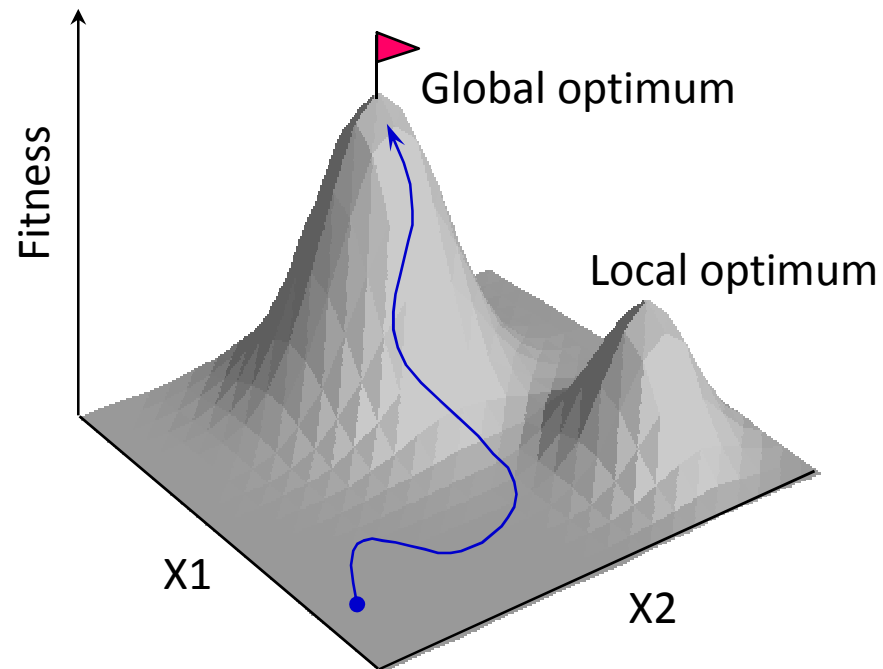
Requirements

- **Structure sampling**
- **Structure assessment**
- **Search / Optimization Method**

Implementation

- Grow
- Link
- Lattice
- Stochastic
- Primary constraints (receptor, ligand)
- Secondary constraints
- Depth-first search
- Breadth-first search
- Random search
- Evolutionary Algorithm
- Monte Carlo / Metropolis
- Exhaustive enumeration
- (Free Energy Perturbation)

Adaptive Walk in a “Fitness Landscape”

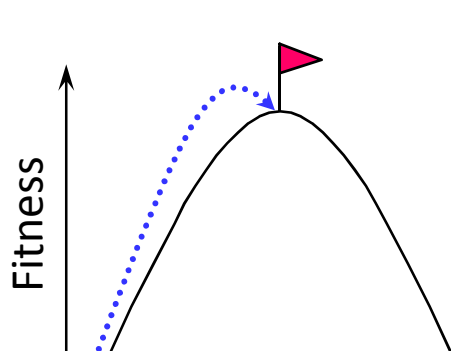


Schneider & Fechner (2005) Nat. Rev. Drug Discov. 4, 649.

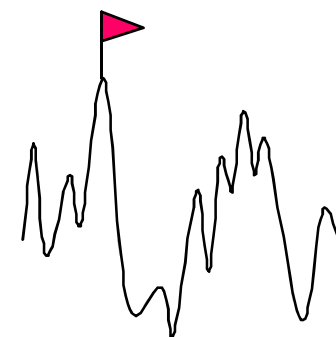
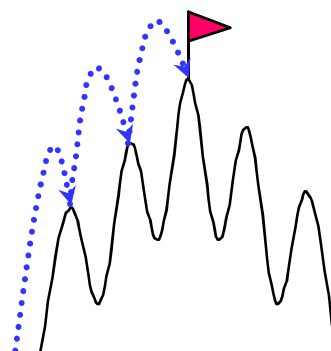
Fechner & Schneider (2007) JCI 117, 656.

23 June 2008, (c) G. Schneider

Types of Fitness Landscapes



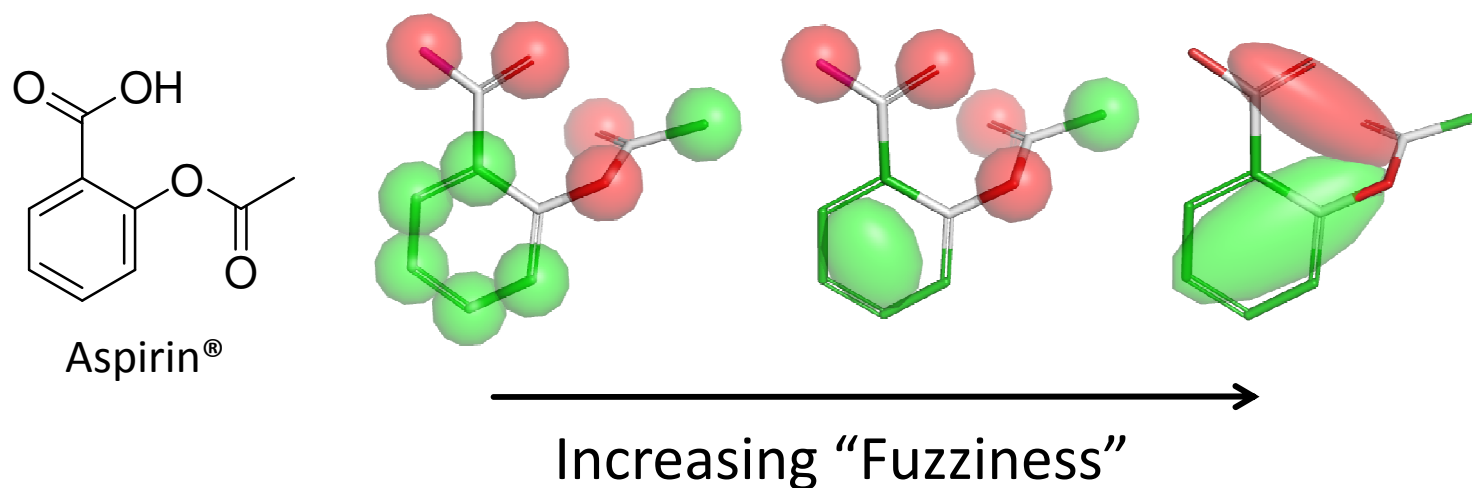
The Dream



The Nightmare

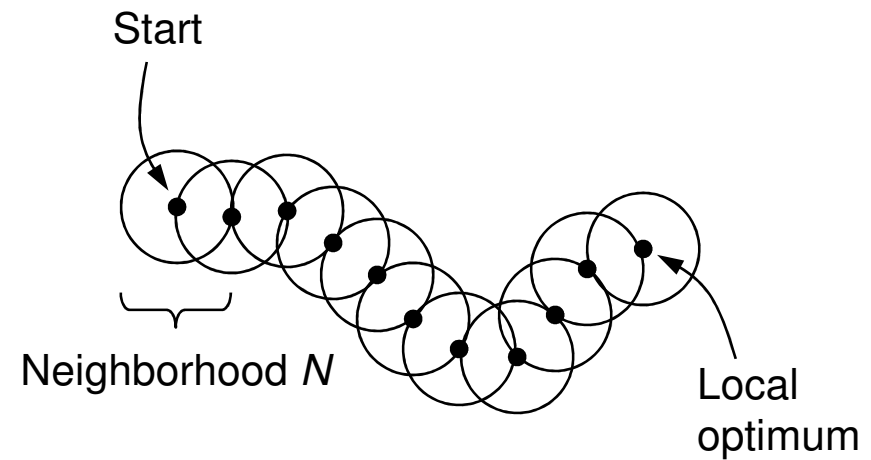
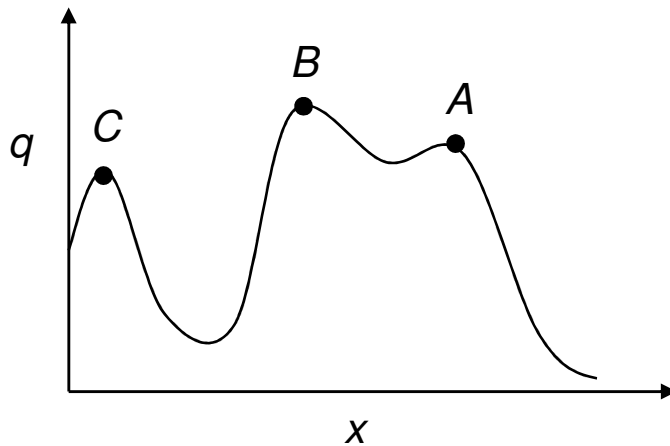
Chemical Similarity Principle!

What is a Molecule?



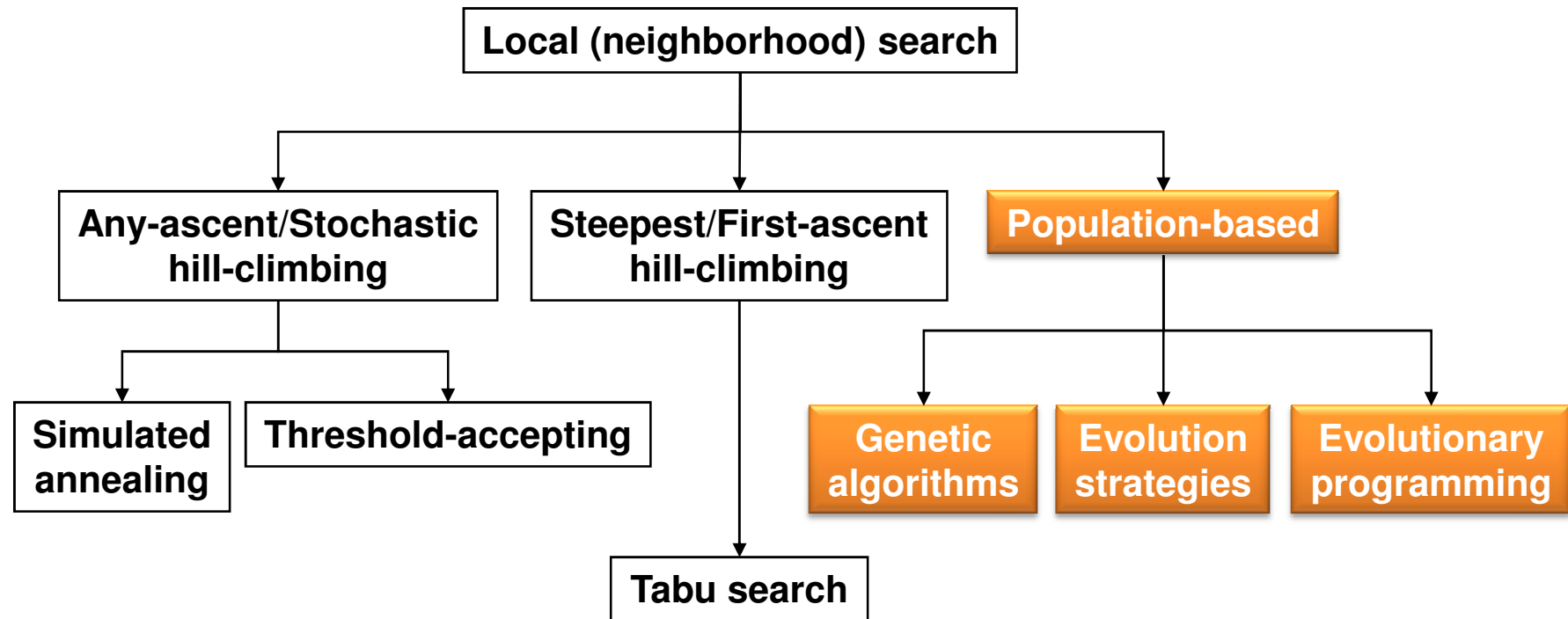
- Pharmacophoric representation on different levels of abstraction

Local Neighborhood Search



- fixed N
- variable ("adaptive") N

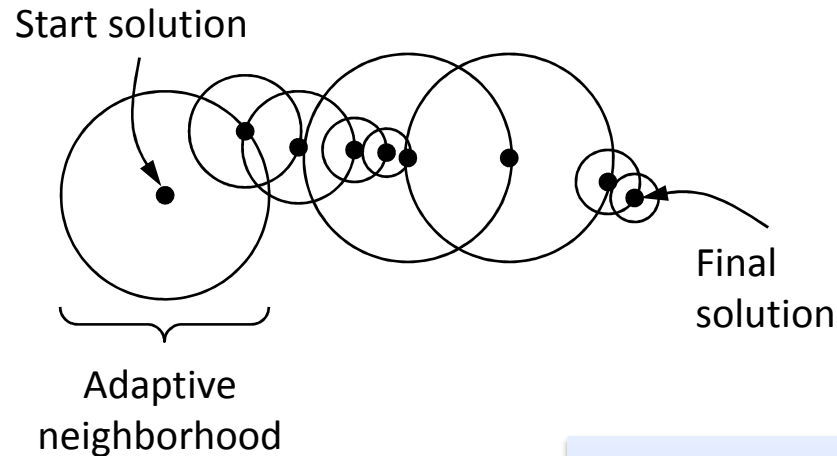
Local Neighborhood Strategies



Elements of Artificial Adaptive Systems (Koza, 1992)

- Structures that undergo **adaptation**
- Initial structures (starting solutions)
- Fitness measure that evaluates the structures
- Operations to modify the structures
- State (**memory**) of the system at each stage
- Method for designating a result
- Method for terminating the process
- Parameters that **control** the process

Why “Adaptive”?



Adaptive optimization copes with ...

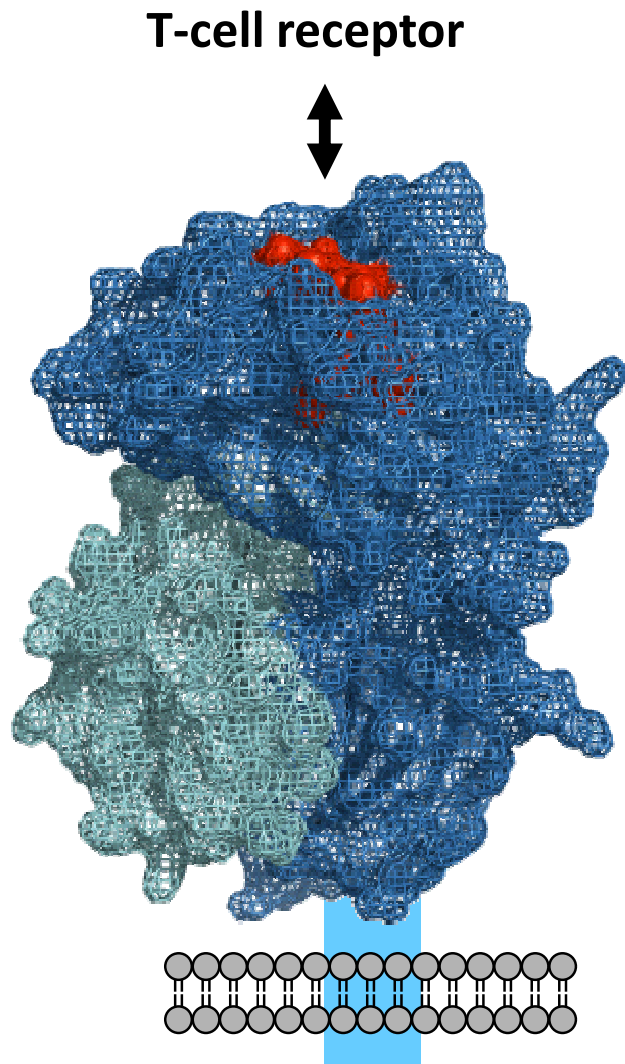
- non-additive fitness functions
- multiple fitness functions
- very large search spaces
- nonlinear system behavior

Learning from Nature

- **Genetic algorithms**
- **Evolution strategies**
- **Particle Swarm Optimization (PSO)**
- **Ant Colony Optimization (ACO)**

Ant Colony Optimization

A Simple Combinatorial Case: Peptide Design

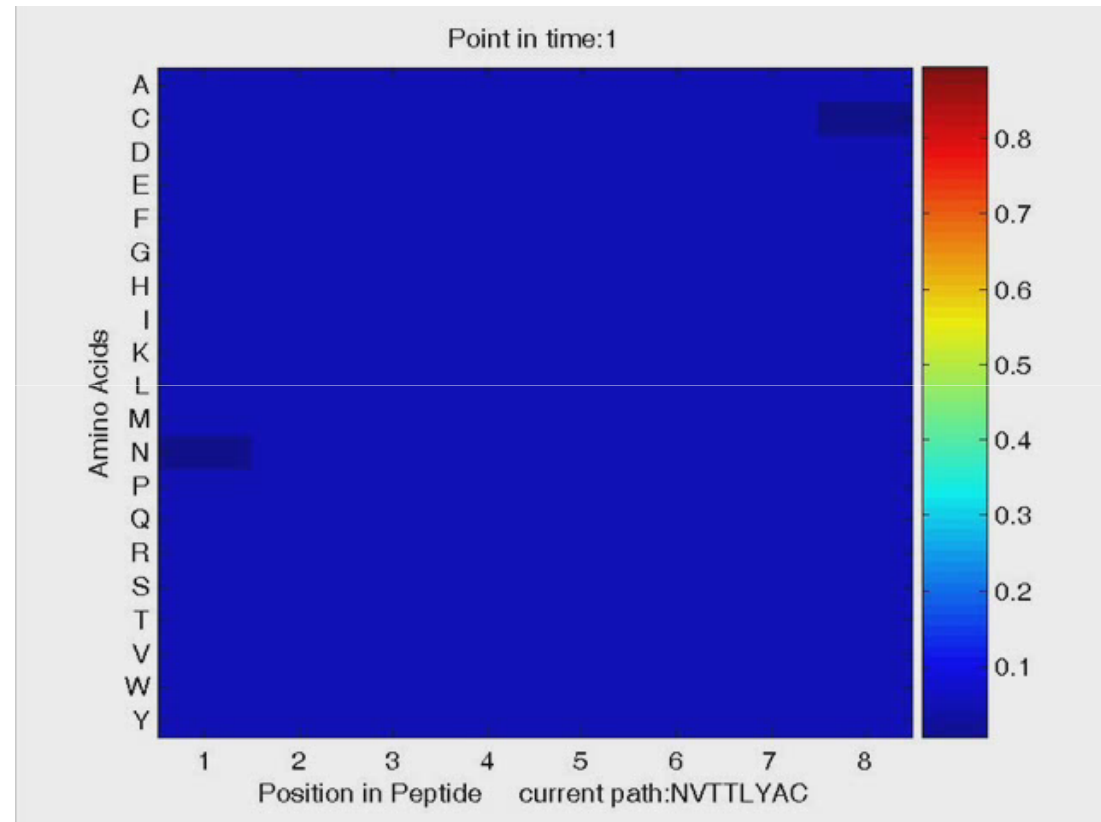
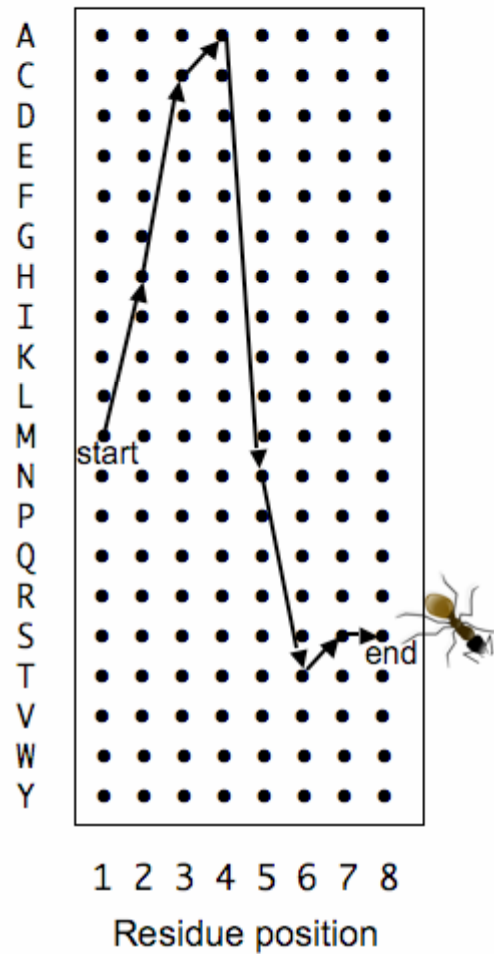


- Design of novel antigens presented by MHC I (H-2Kb)
- Length: 8 residues (20^8 possibilities)
Neural network ensemble + jury
Artificial ant system

Assay confirms the predictions:
89% correct for binding peptides
95% correct for non-binding peptides

Hiss *et al.* (2007) PEDS 20, 99.
Schneider *et al.* (1998) PNAS 95, 12179.

Ant System for Combinatorial Design



Which Search Strategy Should be Applied?

Global & local search control guided by

- Random parameter variation
- Stepwise systematic parameter variation
- Gradients in the fitness landscape ...

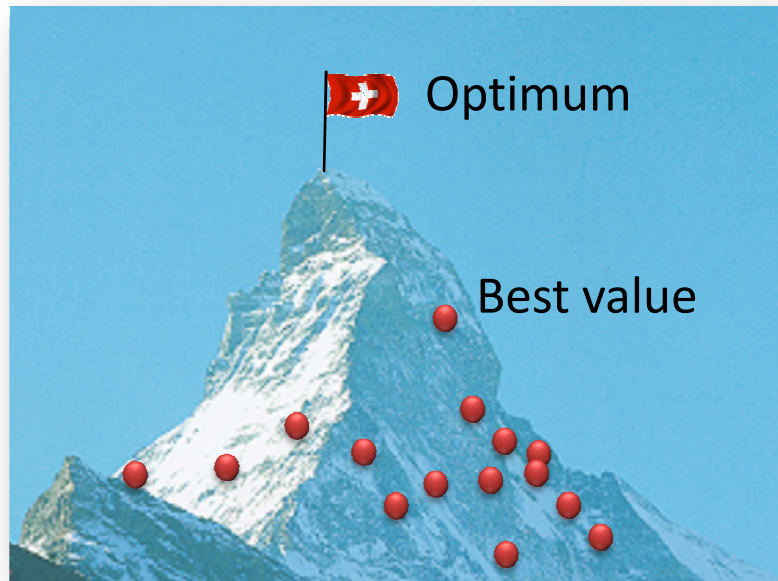
There is no one best method.

Evolutionary algorithms are

- easily implemented
- robust
- able to cope with high-dimensional optimization tasks

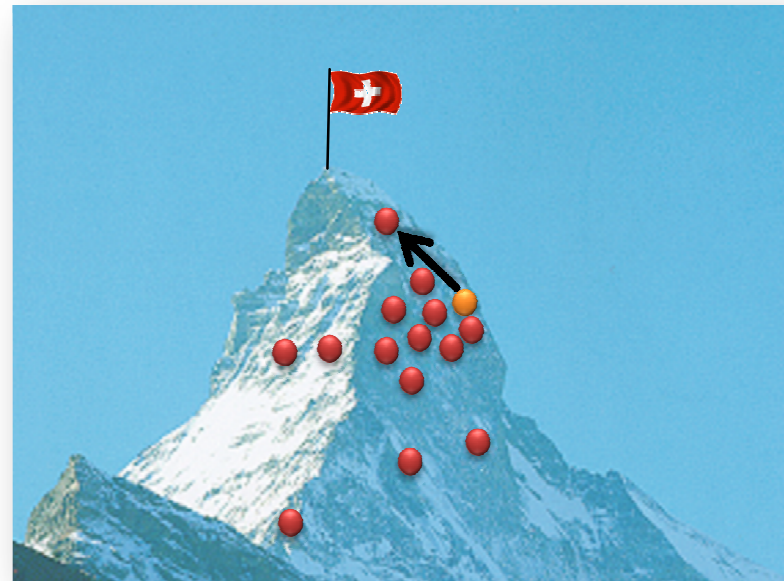
Principle of Evolutionary Strategies

Step 1: Random search



Generate a *diverse* set of parameter values and hope for a hit

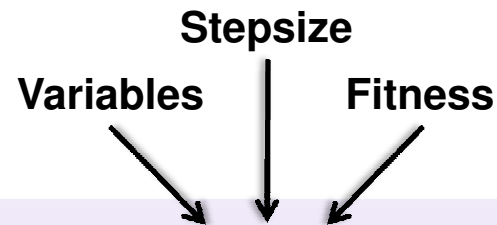
Step 2 and following:
“Local hill climbing”



Generate *localized distributions* of parameter values and improve steadily

Adaptive neighborhood

Algorithm of the $(1,\lambda)$ Evolution Strategy (Rechenberg, 1973)



Initialize parent (x^P, s^P, F^P)

For each generation:

Generate λ variants (x^V, s^V, F^V) around the parent:

$$s^V = \text{abs}(s^P + G)$$

$$x^V = x^P + s^V \cdot G$$

Calculate fitness F^V

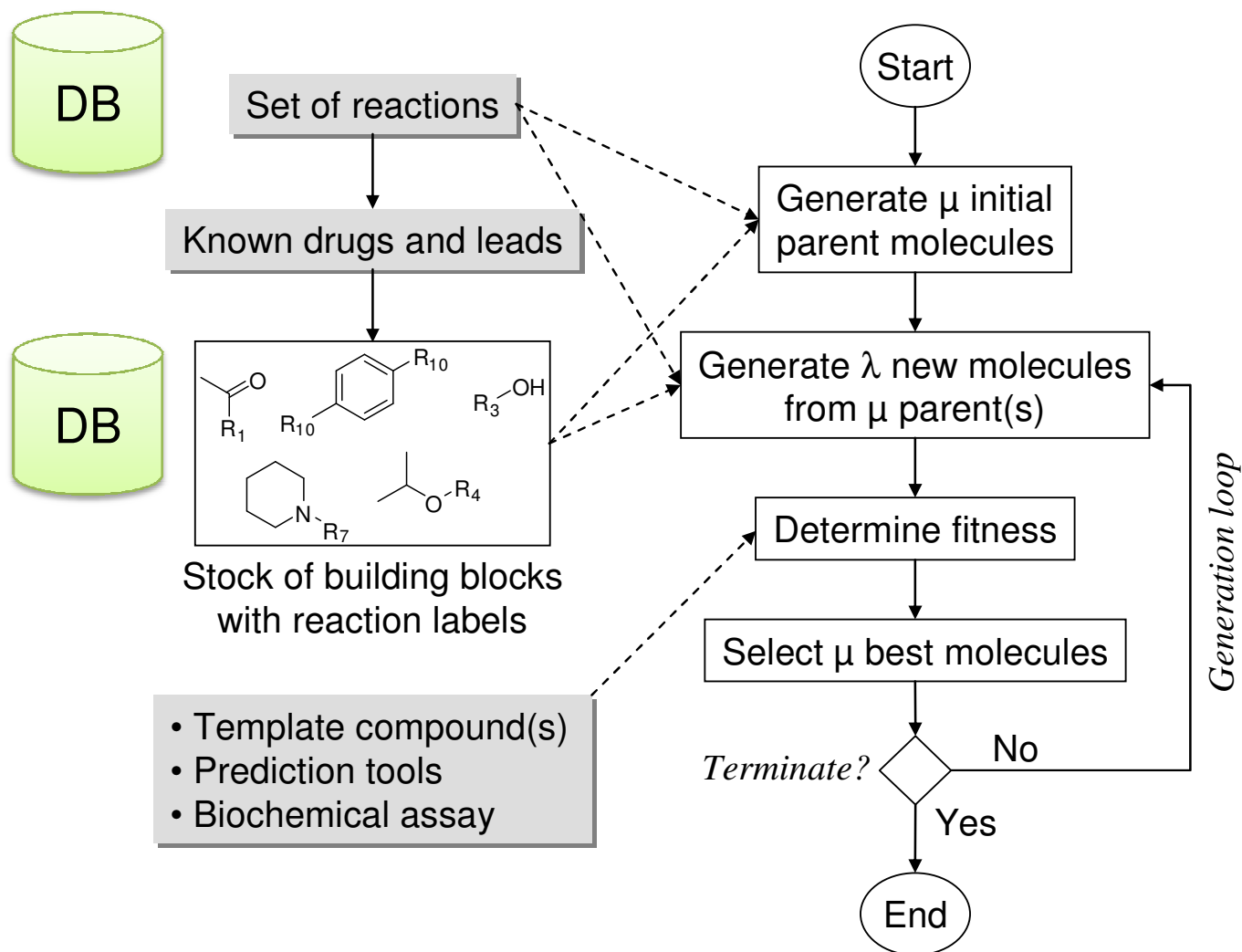
Select best variant according to F^V

Set $(x^P, s^P, F^P) = (x^V, s^V, F^V)^{\text{best}}$

$$G(i, j) = \sqrt{-2\ln(i)} \cdot \sin(2\pi j) \quad \text{in } [-\infty, \infty]$$

i, j : pseudo-random numbers in $]0, 1]$

De novo Fragment Assembly



Schneider et al. (2000) Angew. Chem. Int. Ed. 39:4130

23 June 2008, (c) G. Schneider

Operators of Evolutionary Optimization

Variation



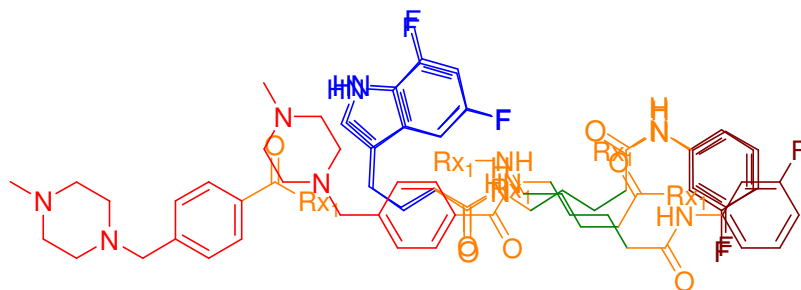
- **Mutation**
- **Crossover**
- **Selection**

Adds new information to a population

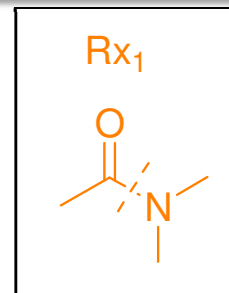
Exploits information within a population

Breeding New Molecules

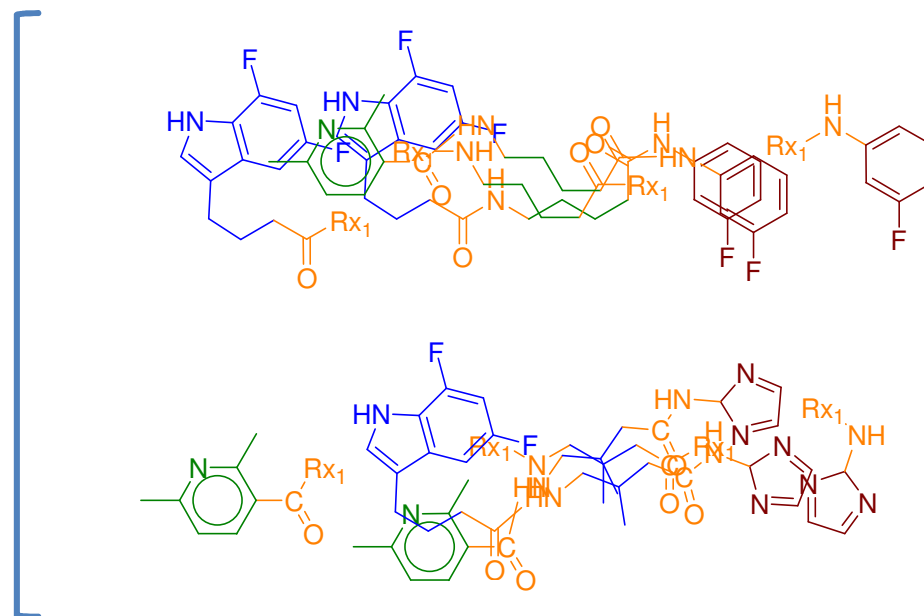
Mutation



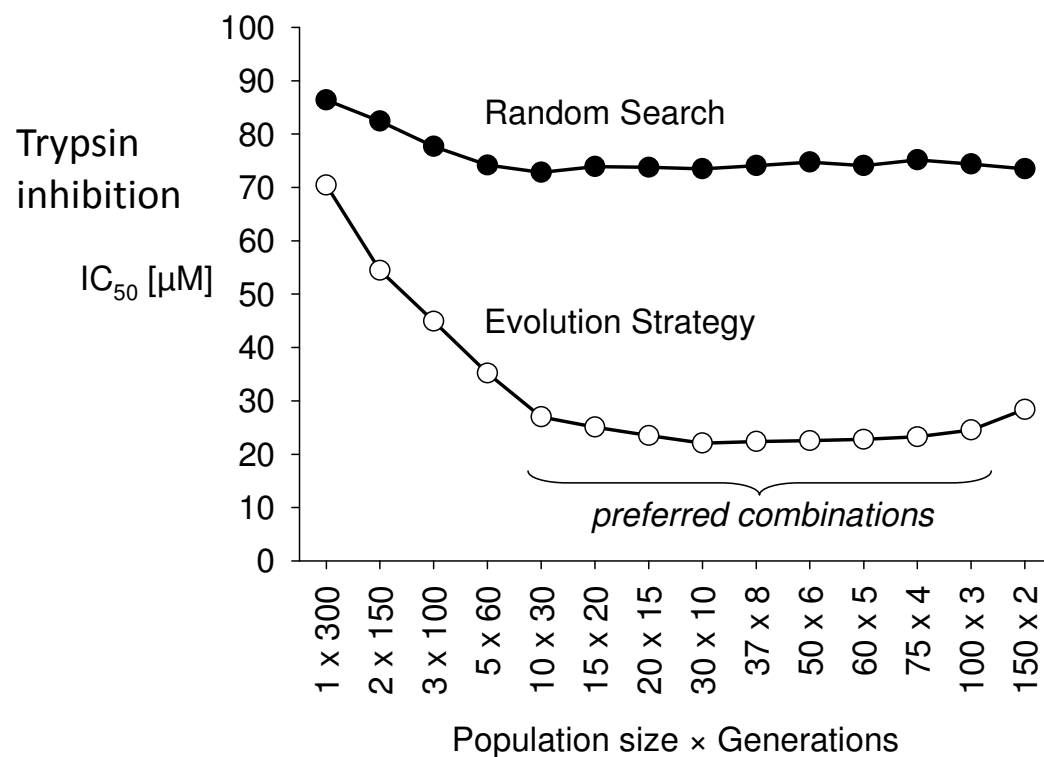
Virtual reaction



Crossover



How Many Iterations? How Many Compounds?



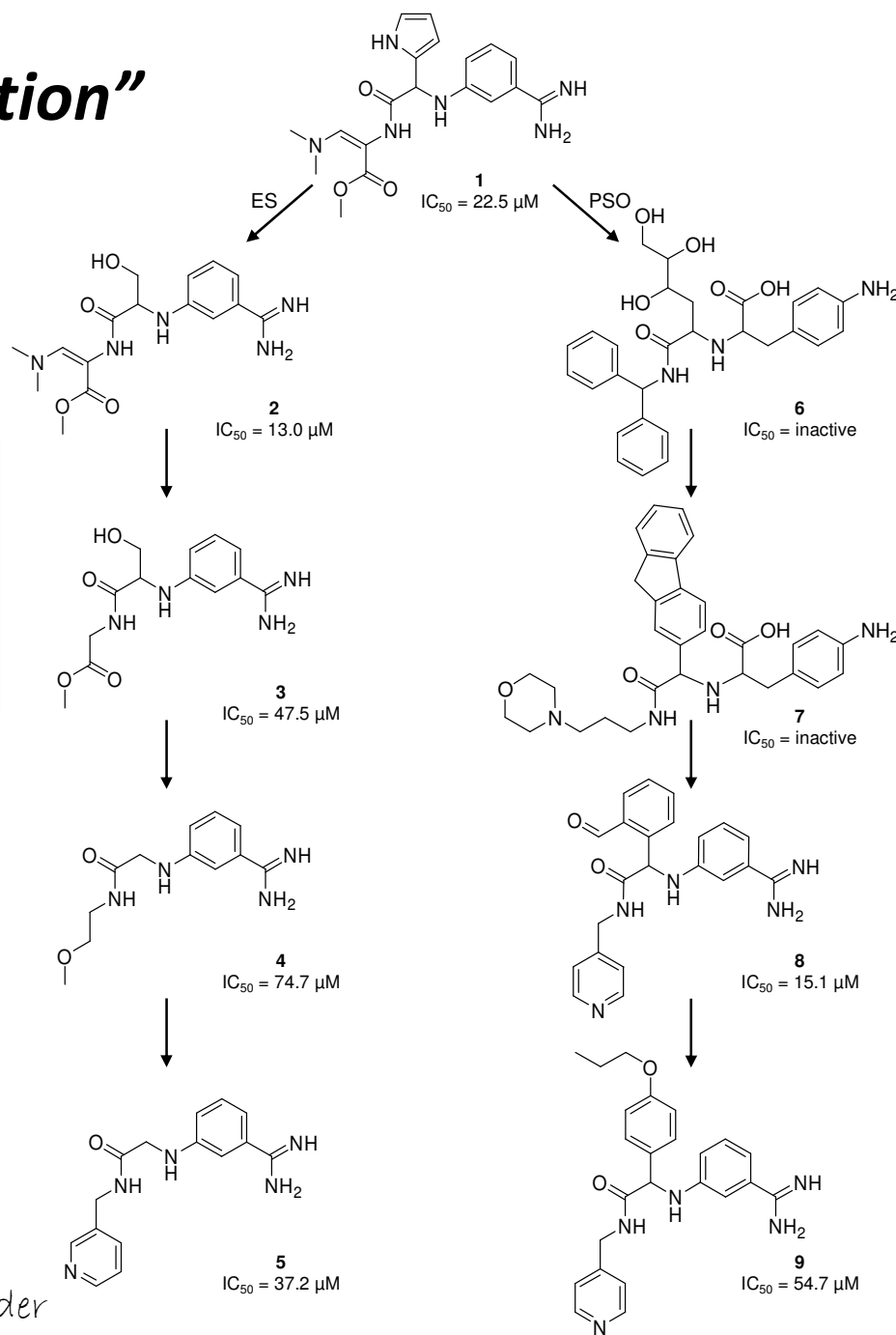
Limited resources:
300 compounds
There is an optimum!

Schüller & Schneider (2008) JCIIM, in press.

23 June 2008, (c) G. Schneider

"Molecular Evolution"

Trajectories through inactive regions of chemical space are possible

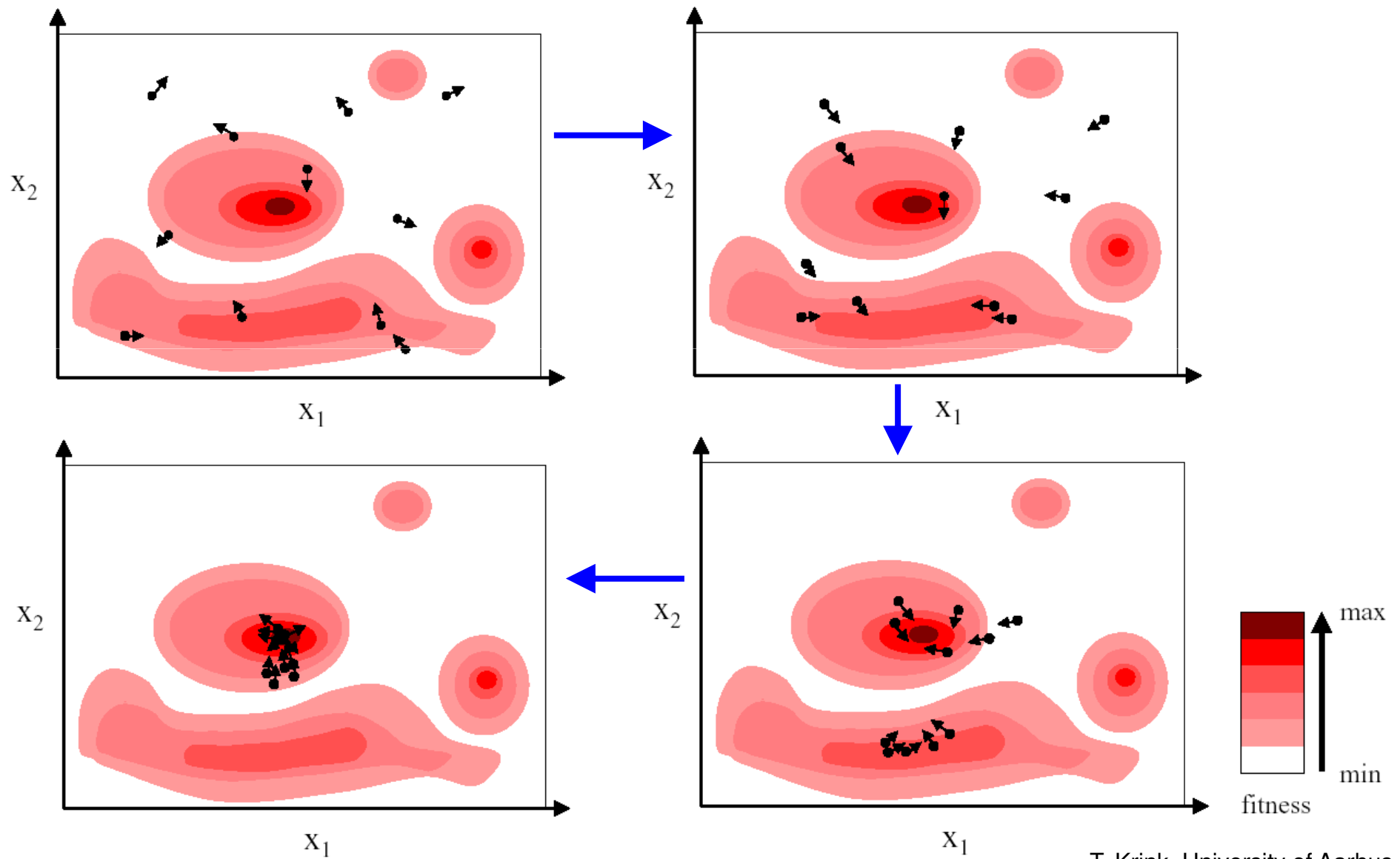


23 June 2008, (c) G. Schneider

Particle Swarm Optimization (PSO)

- Biological motivated optimization paradigm
- Individuals move through a fitness landscape
- Particles can change their **movement direction** and **velocities**
- **“Social” interactions** between individuals of a population
- **Each** particle knows about its **own** best position
- **All** particles know about the **overall** best position

Particle Swarm Optimization (PSO)



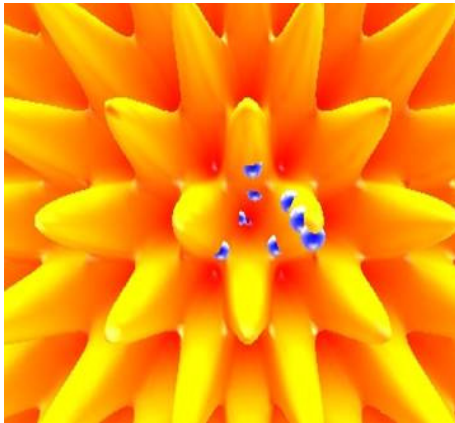
PSO: Update Rule 1 (Standard Type)

$$v_i(t+1) = v_i(t) + \underbrace{2 \cdot r_1 \cdot (p_i - x_i(t))}_{\text{Individual memory}} + \underbrace{2 \cdot r_2 \cdot (p_b - x_i(t))}_{\text{Social memory}}$$

- v:** Velocity vector
- x:** Coordinates of a particle
- i:** Dimension
- t:** Time (epochs)
- r_1, r_2 :** Random numbers in]0,1]
- p_i, p_b :** Best positions (individual and social)

PSO: Individual and Social Memory

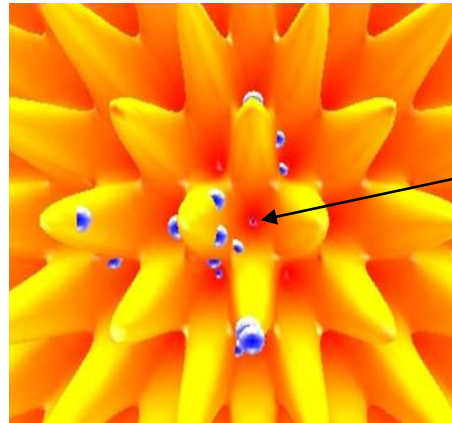
PSO convergence after 100 epochs (Griewangk's function)



$$n_1 = 1$$

$$n_2 = 2$$

**Higher weight on
the social memory**

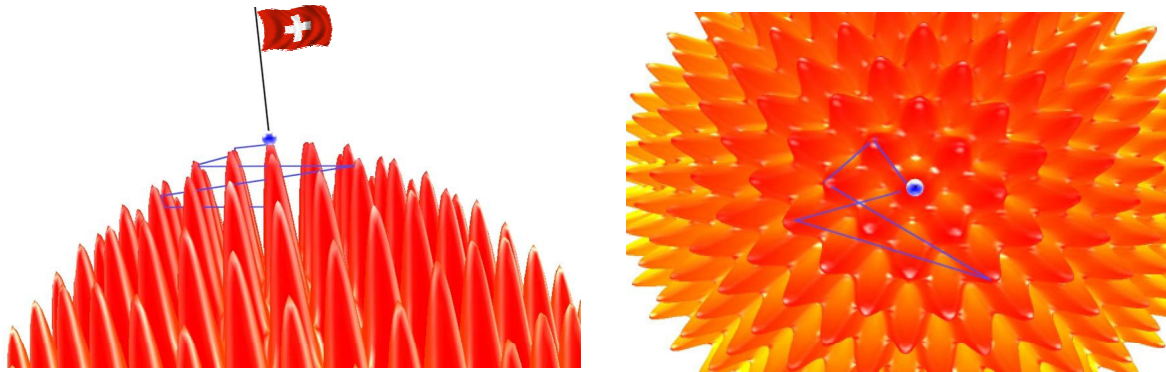


Global optimum

$$n_1 = 2$$

$$n_2 = 1$$

Visualization of Particle Trajectories

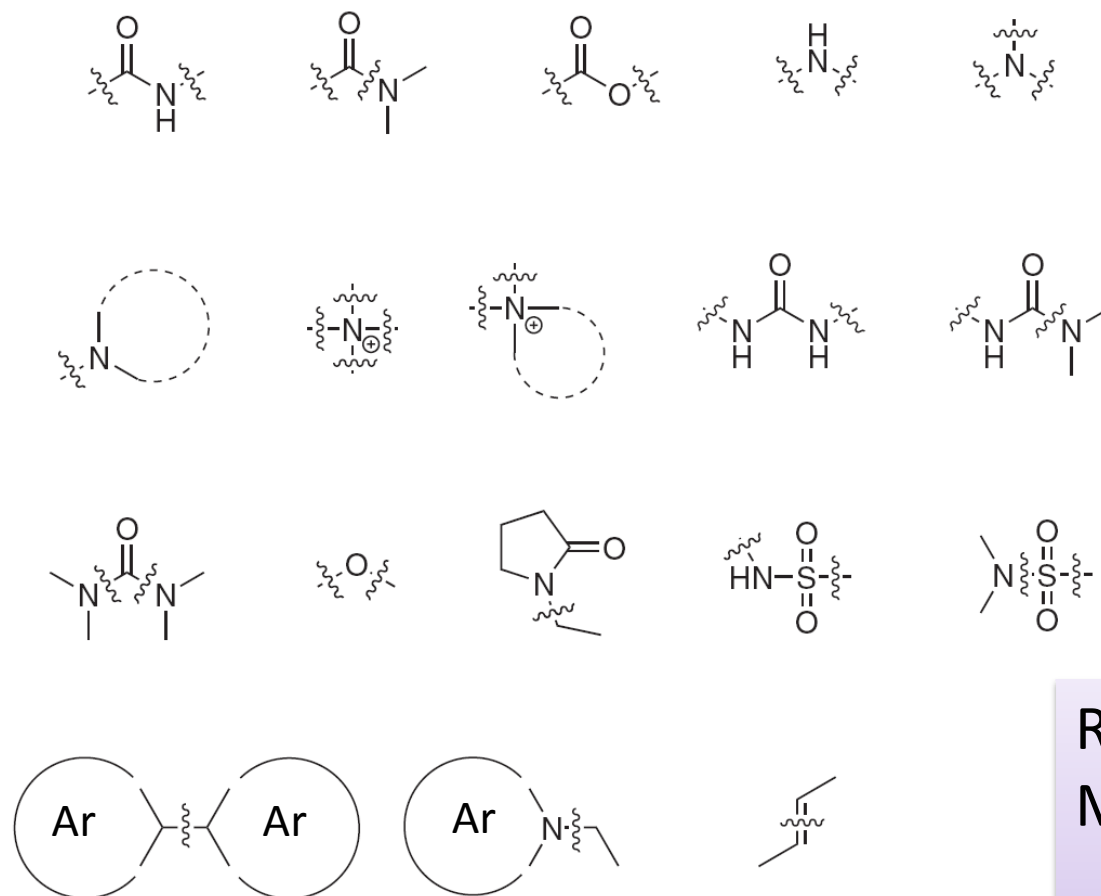


- The particle visited several local optima before finding the global optimum

COLIBREE®

Combinatorial Library Breeding

Disassembly Rules

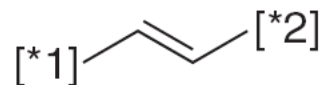
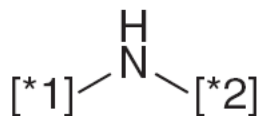
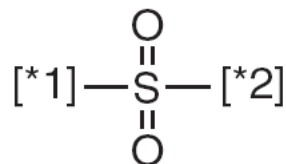
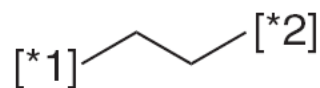
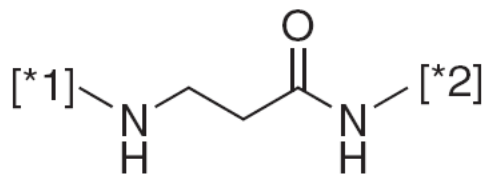
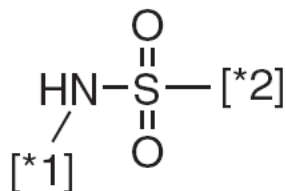
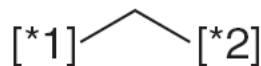
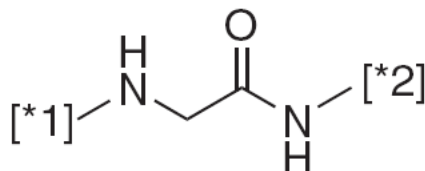
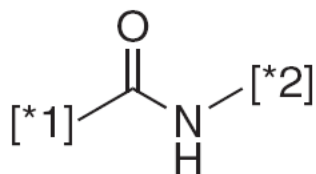
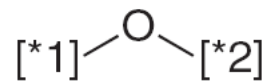
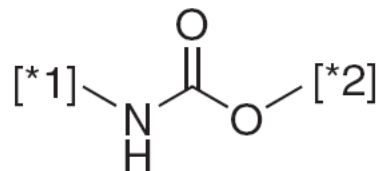
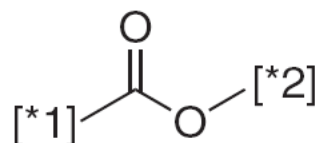


Retain building blocks with
MW < 200 Da

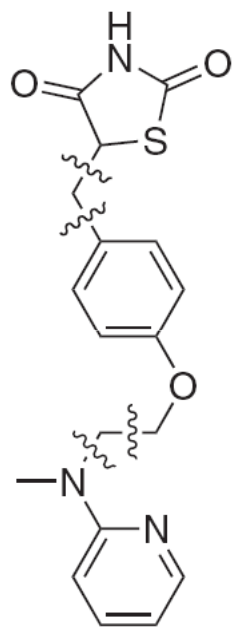
7,184 Building Blocks
avg MW = 141 ± 35 Da

Hartenfeller et al. (2008) Chem. Biol. Drug Des., in press
23 June 2008, (c) G. Schneider

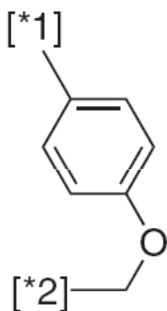
Linker Library



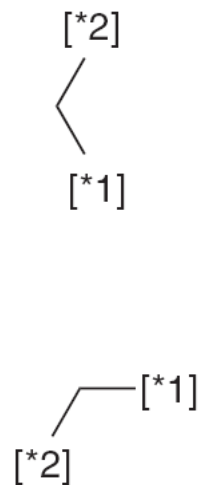
Molecule Dissection



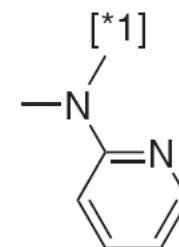
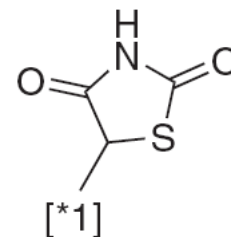
Rosiglitazone



Scaffold

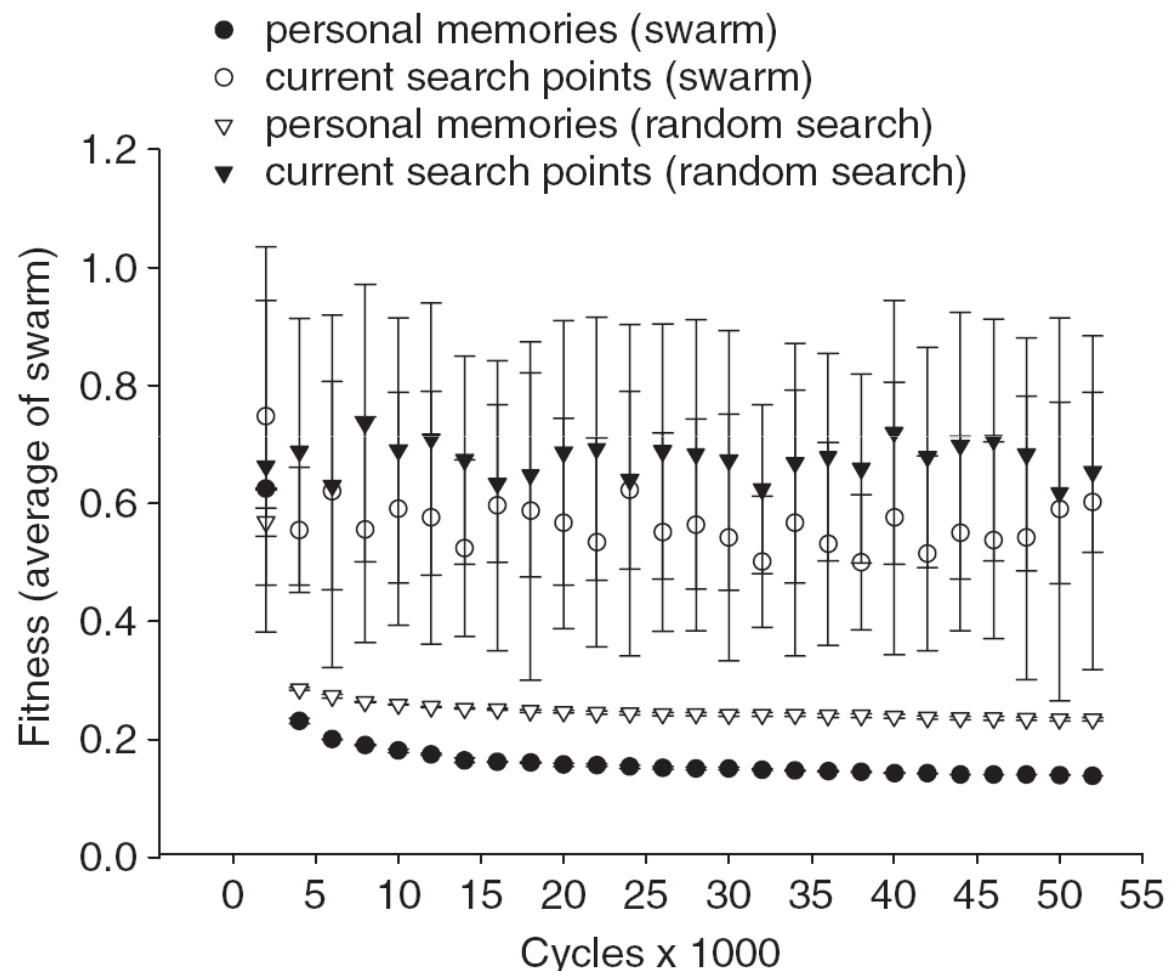


Linker



Building blocks

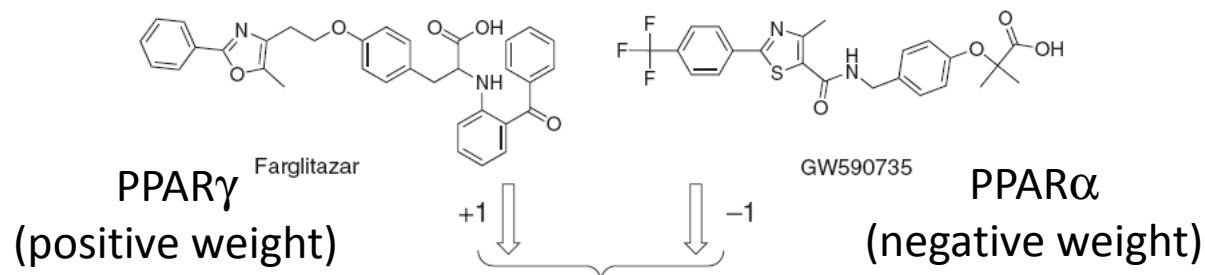
PSO Progress Over Time



Library size:
40 molecules

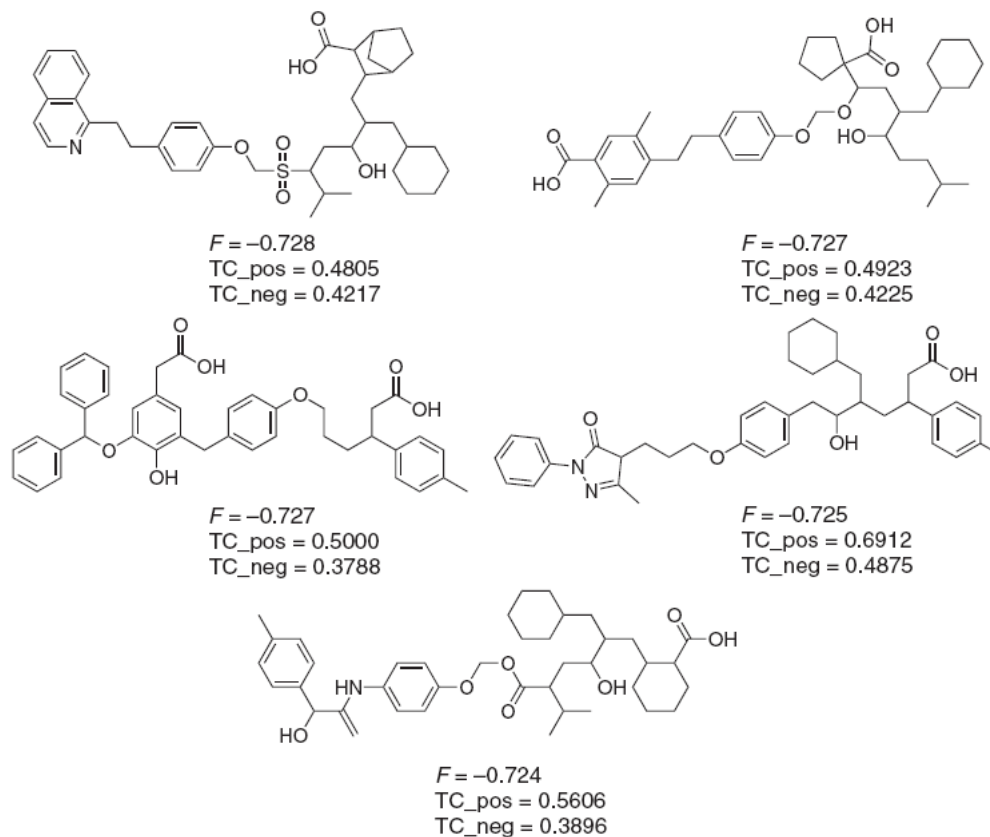
Fitness:
Distance to Reference
(minimization!)

Positive and Negative Design



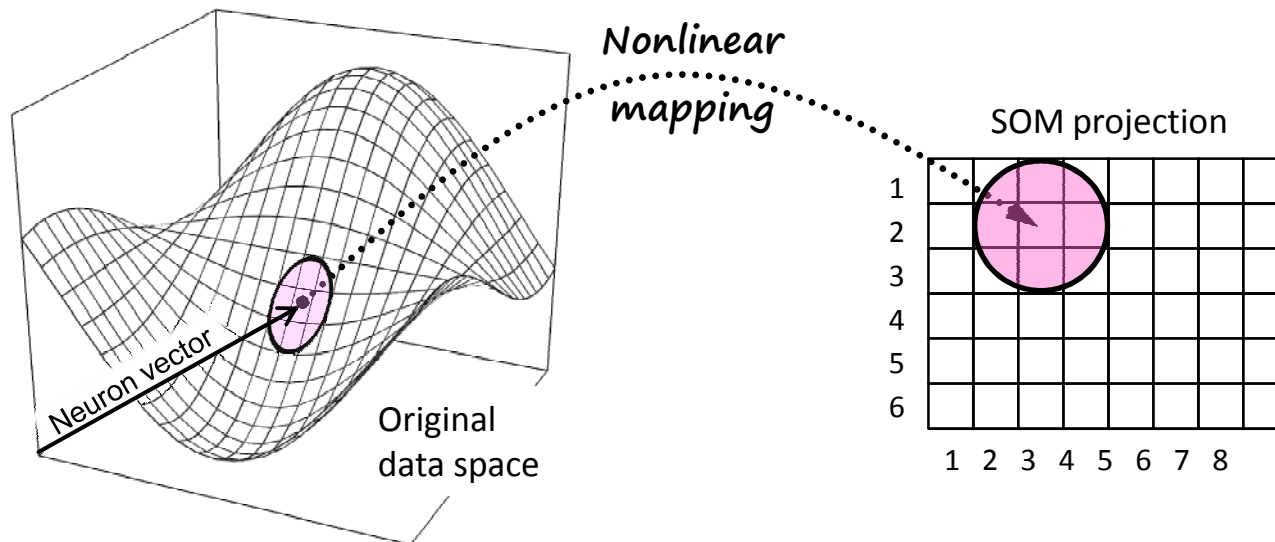
$$F(\mathbf{X}) = \sum_{j=1}^r w_j \bullet D_{\text{Euklid}}(\mathbf{X}, \mathbf{Y}_j)$$

$$D_{\text{Euklid}}(\mathbf{X}, \mathbf{Y}) = \sqrt{\sum_{i=1}^n (x_i - y_i)^2}$$



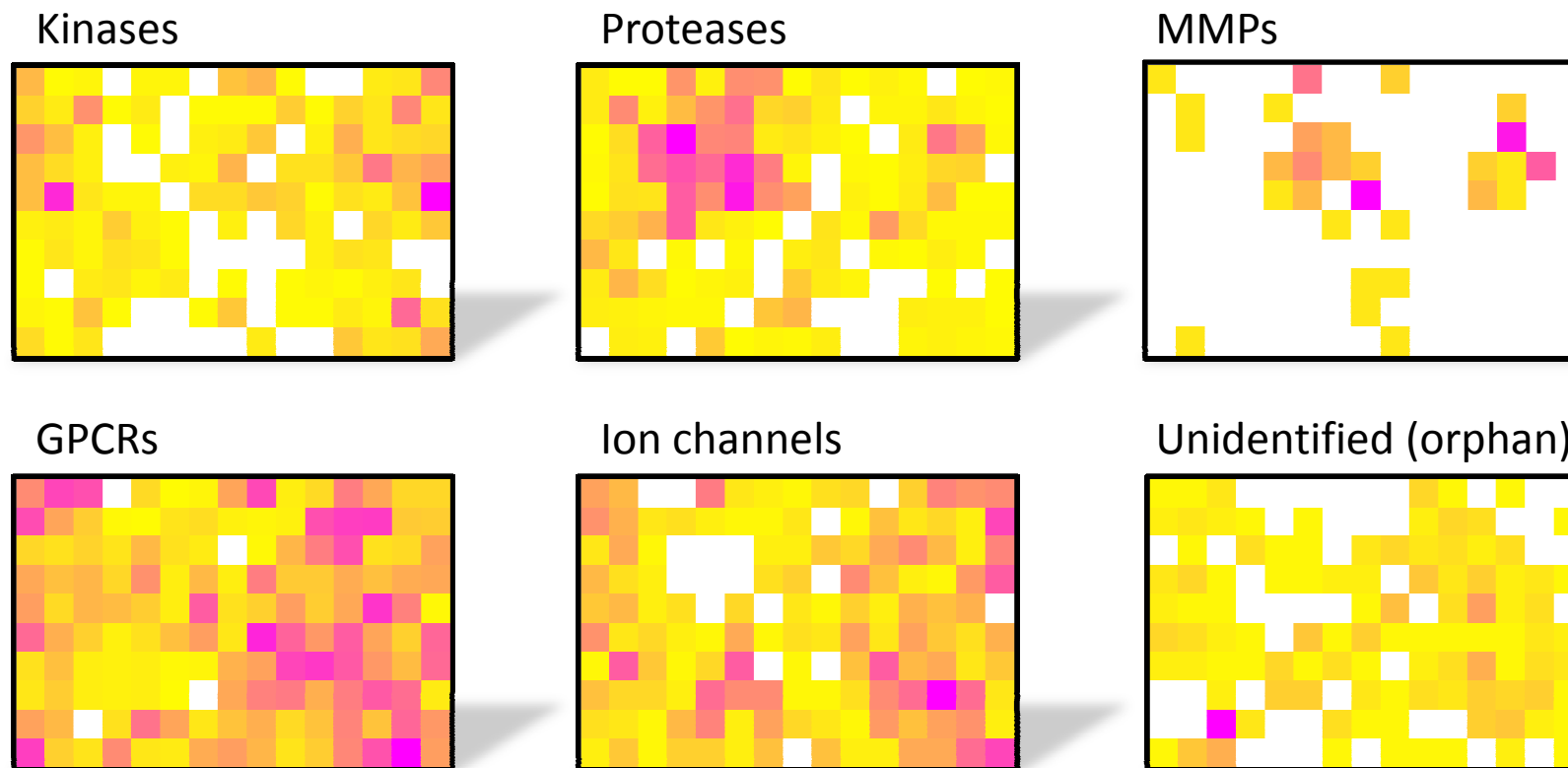
Self-Organizing Map (SOM)

Kohonen , 1982



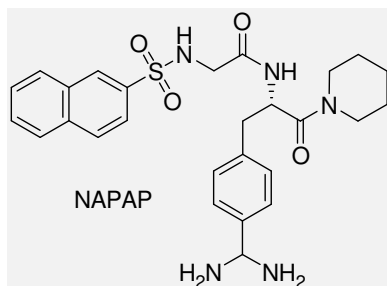
- Nonlinear projection of high-dimensional space
- Topology-preserving projection
- Data clustering
- Prediction with probability estimation

Pharmacophore Road Map of Chemical Space

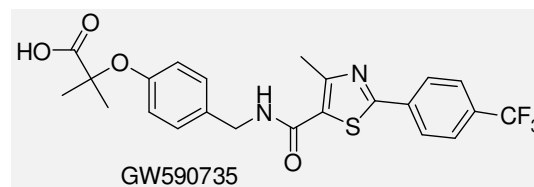
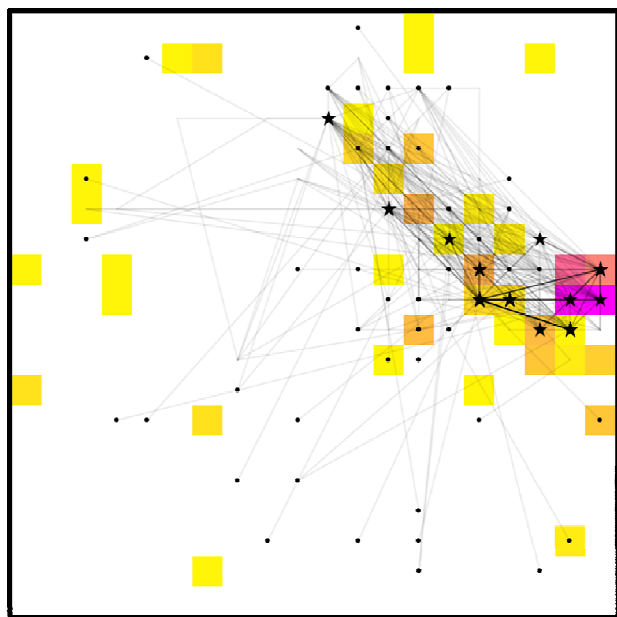


- Topological pharmacophores (CATS descriptor)
- COBRA collection (10,500 drugs and lead compounds)

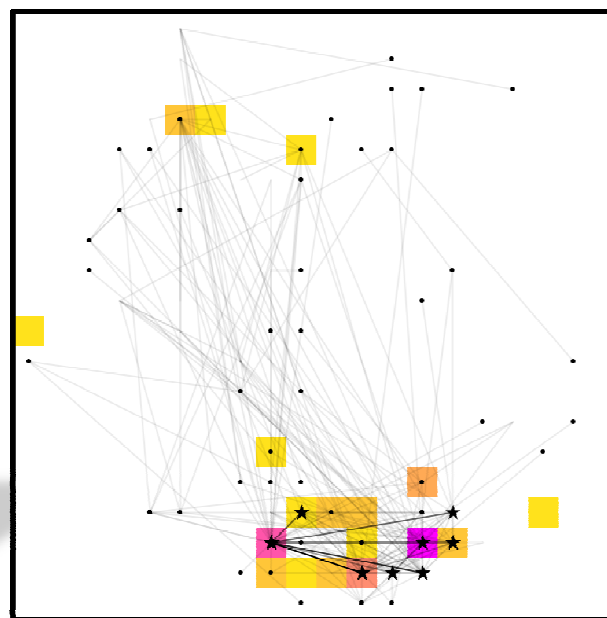
Land Ho! Exploration of Chemical Space



Thrombin inhibitors



PPAR α modulators

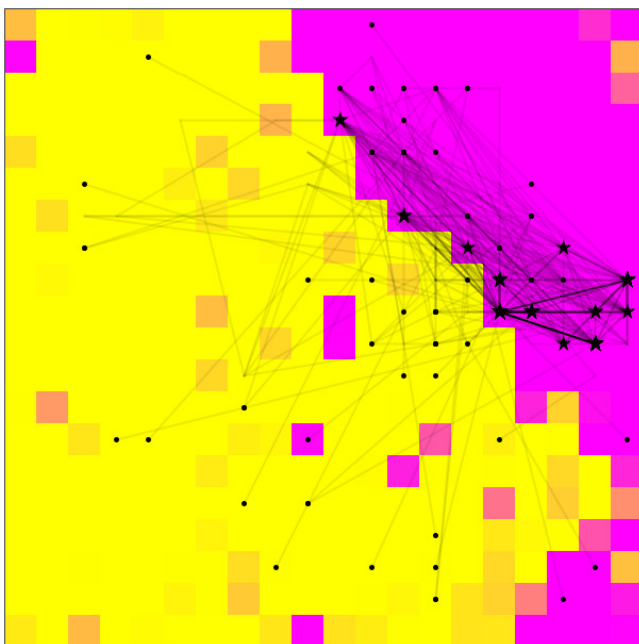


Let's Be Positive!

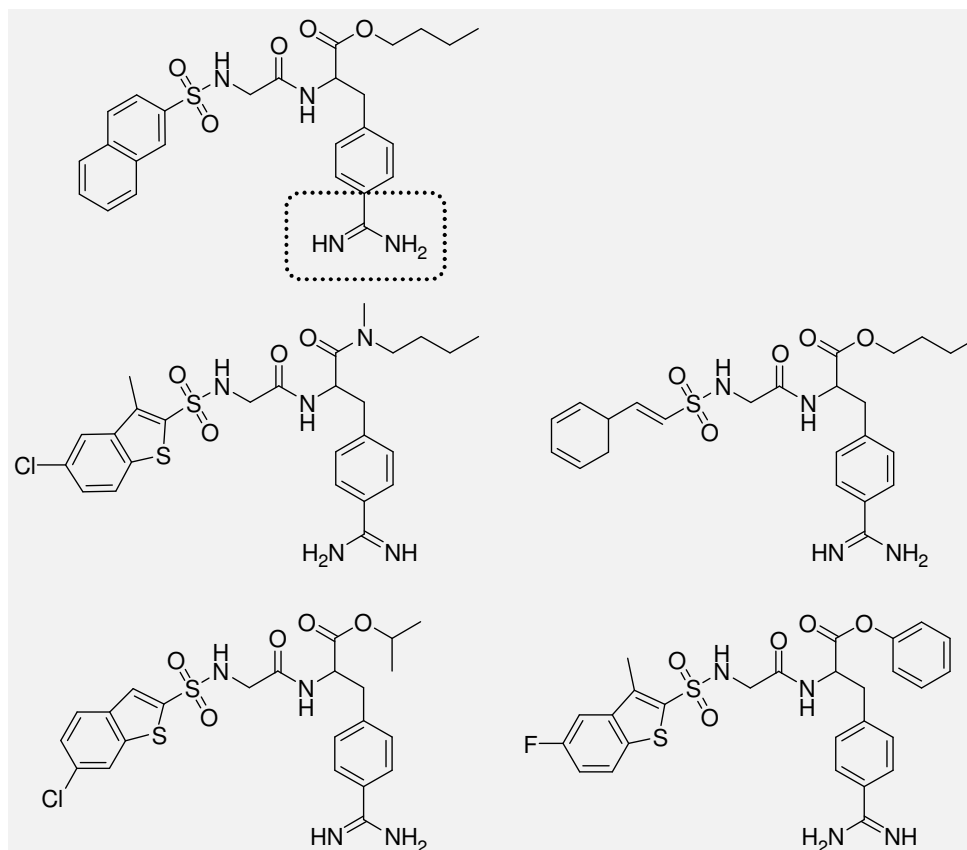
De Novo Design of Thrombin Inhibitors

CATS descriptor no. 10:

PP0 = $\oplus$

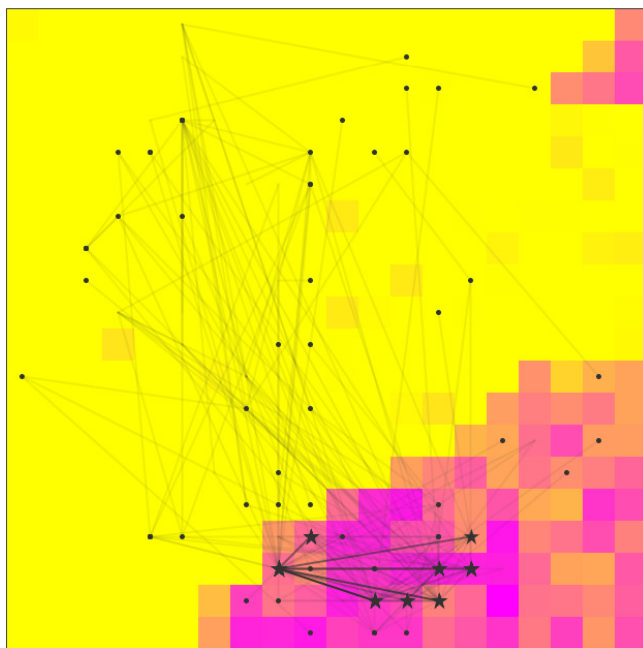
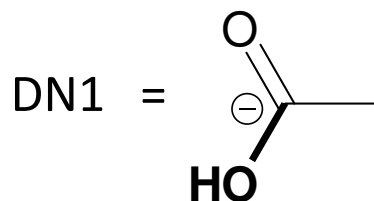


Top-ranking designed compounds

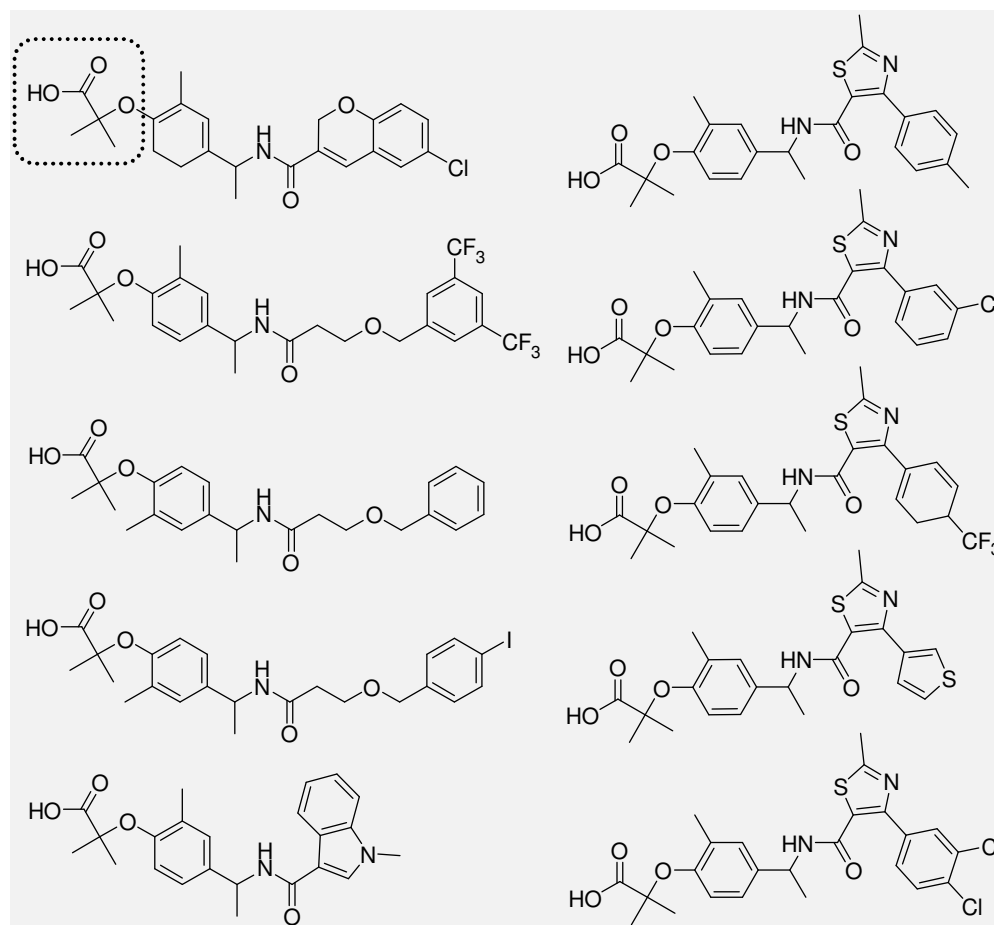


Climbing “Acid hill”: De Novo Design of PPAR α Agonists

CATS descriptor no. 19:



Top-ranking designed compounds



Conclusions

- Fragment-based design often results in bioactive compounds
- “Novelty” of the designs depends on local convergence
- Adaptive optimization enables “scaffold-hopping”
- Visual inspection / expert knowledge is crucial

Acknowledgement



Prof. Manfred Schubert-Zsilavecz
Prof. Holger Stark
Prof. Dieter Steinhilber



Dr. Karl-Heinz Baringhaus



Dr. Tanja Weil

modlab[®] Team
www.modlab.de

