

GECCO = GP + GA + ES + EP + ER + DNA + LCS + RWA + AAAA +


Genetic and Evolutionary Computation Conference 2004

June 26 - 30, 2004 (Saturday-Wednesday) Seattle, Washington USA

# Evolutionary Product Design: Rocket Science or Art?

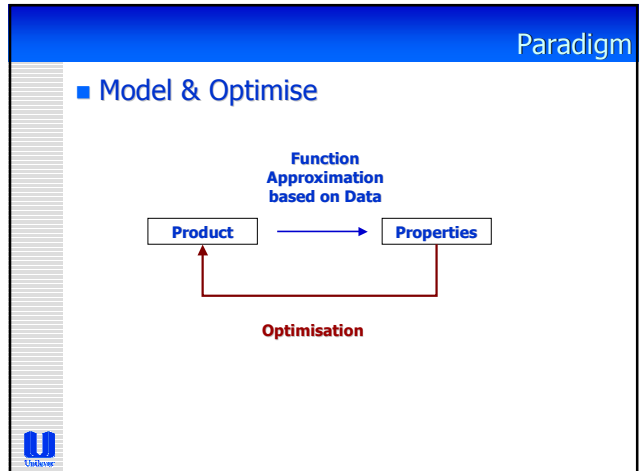
EVOLUTIONARY COMPUTATION IN INDUSTRY SESSION

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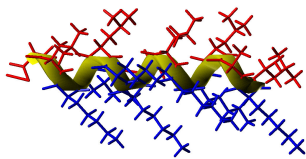
Meeting everyday needs of people everywhere





>Case 1>Problem Description  
**Design an anti-bacterial peptide**

- **Natural Representation:**
  - String of amino acids
  - **KLALALLAGSAASA**



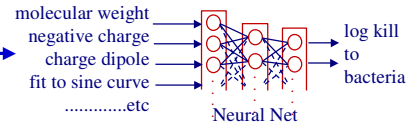
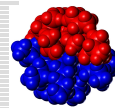
- Alpha-Helical Peptide
- Side View
  - red = hydrophobic
  - blue = hydrophilic

*Patenting computer-designed peptides, Patel, Stott, Bhakoo & Elliott, Journal of Computer-Aided Molecular Design, 12: 543-556 1998*



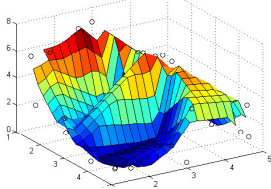
**Fitness Function**

■ **Two Step Function**

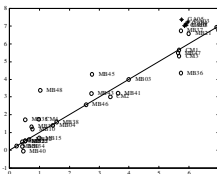
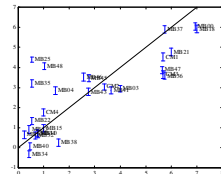


**Non-linear Mapping**

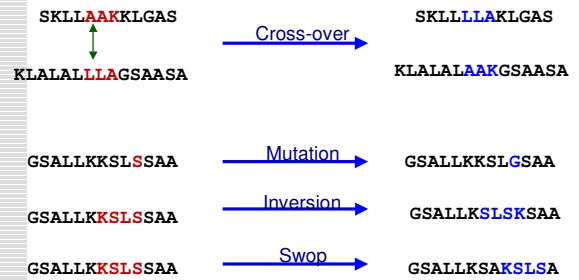
Projection of 29 peptides to Kohonen map with 3d interpolation



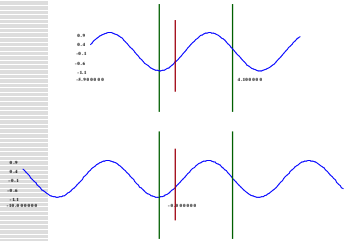
	xvalid R2 Linear	xvalid R2: Neural Net
Aureus	0.77	0.91
E Coli	0.75	0.81



**Operators**



## Special Cross-Over Operator

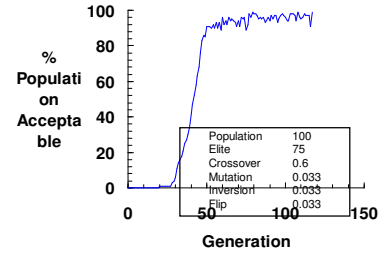


- Represent as right and left halves, separated by 'zero crossing' of hydrophobicity (red line in graph)
- Canonically align by right and left halves
- Select random (aligned) segments
- Perform cross-over



## Running the GA

- Convergence in about 50 generations



## Results: New Peptides

Peptide Sequence	predicted log kill	measured log kill
AASKAAKTLAKLLSLLKLL	7.22	> 5.96
LLKKLLRAASKALSLL	7.13	> 5.90
AAKLSKLLKTLKLL	7.35	> 5.76
KALKKLLKLASSLTAL	7.04	5.90
AASKALRTASRLARSLTLL	7.03	> 5.85

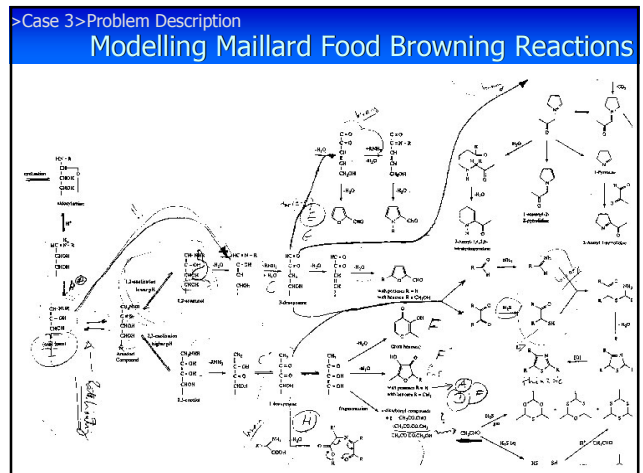
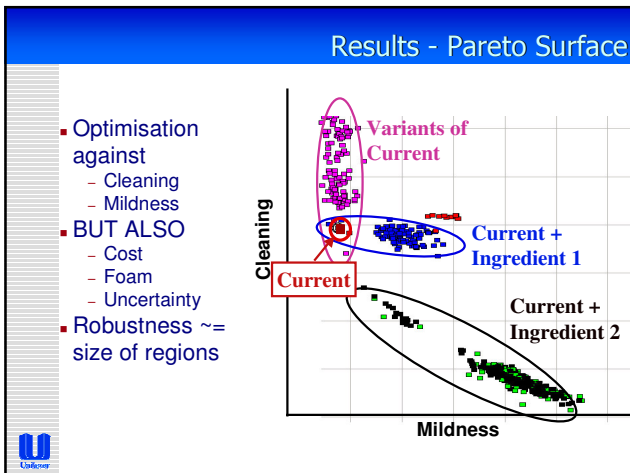
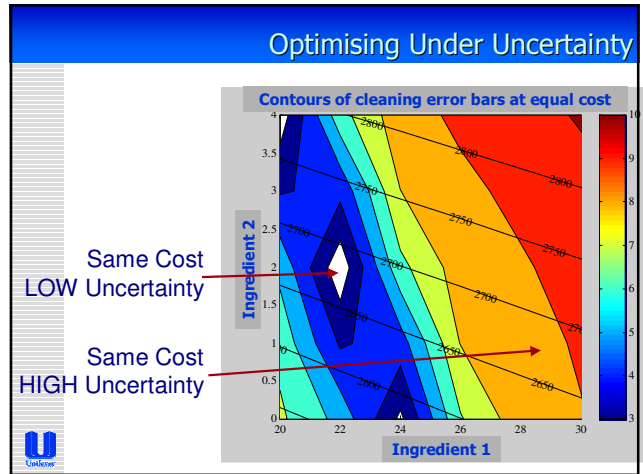
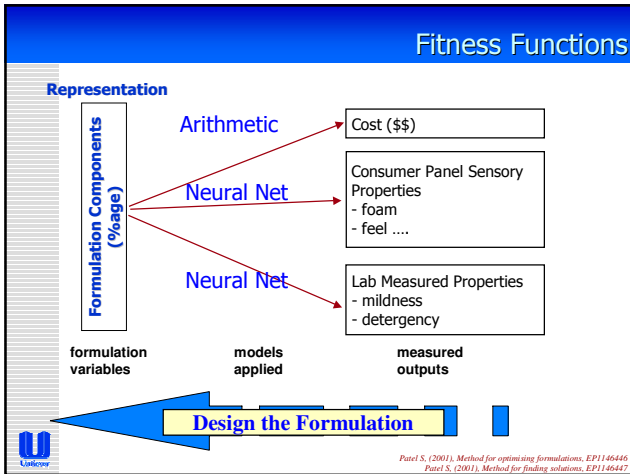


## >Case 2>Problem Description

### Multi-Criteria Formulation Optimisation

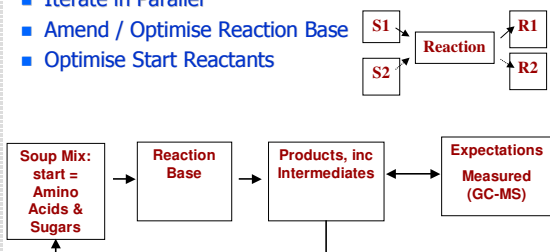
- Current Process:
  - requirements
  - formulators experience
  - past products
  - experiments
- New Process:
  - requirements
  - formulators experience
  - past products
  - experiments
  - *predictions based on past experiments*
  - *trade-off analysis based on predictions*





## Iterated Reaction Graphs

- Model Reactions by input-output
- Model reactants & products / intermediates as list of molecules
- Iterate in Parallel
- Amend / Optimise Reaction Base
- Optimise Start Reactants



Paul S. Rabone R, Russell S, Tissen J, Klaffer W, (2001) Iterated Reaction Graphs: Simulating Complex Maillard Reaction Pathways. *Journal of Chemical Information and Computer Sciences*, Vol 41, 4, 926-933

## Representation

- List of Molecules in SMILES notation

```
C(O)C(O)C(O)C(O)C(O)C=O
C(O)C(O)C(O)C(O)C(O)C=O
C(O)C(O)C(O)C(O)C(O)C=O
C(O)C(O)C(O)C(O)C(O)C=O
...
NC(C(C)O)C(=O)O
NC(C(C)O)C(=O)O
.....
```

**Sugars**  
**Amino Acids**

## 'Operators' = Chemical Reaction

Chemical Reaction =  
breaking bonds  
making bonds

3-amino-2-butanone  $\rightarrow$  tetramethyl pyrazine precursor

## Output

- Reaction Triplets := <Substrate Reaction Product>

```
C1=O(C)C(-CC(O)O)O R1_1_6_endiol C1=C(C)C(-O)O(O)O 1
C=O(C)C(C)C(O)O(O)O R1_1_6_endiol C1=C(C)C(C)C(O)O(O)O 1
C1=CC(O)C(O)O R1_1_2_enamino C1=C(C)O(NC)C(O)O 1
C1=CN=C(C)O(O) R1_4_2_strecker C1=CN(O)C(O)C(O)O 1
```

- Chain to Form a Graph, eg one path:

```
<Start= C(O)C(O)C(O)C(O)C(O)C=O R1_3_6_sub_enol C=O(C)C(O)C(O)C(O)C=O R1_1_8_2bunsaturated C=O(C)C(O)C(O)C=O
R1_1_8_2bunsaturated C=O(C)C(O)C(O)C(O)C=O R1_2_2_retriosid C=O(C)O(O)C(O)C=O R1_1_1_Schiff base
CO(O)N=CC(C)C(O)C(O)C(O)O R1_4_1_Strecker CO(O)C=NC=C(C)C(O)O R1_4_2_strecker CO(O)C=O R1_1_6_endiol ...
```

- List of products

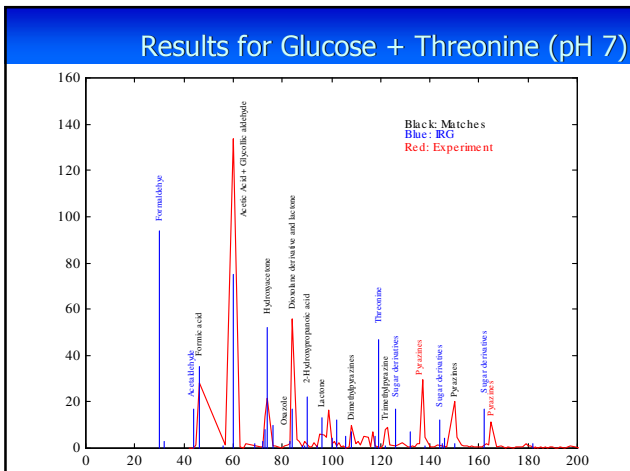
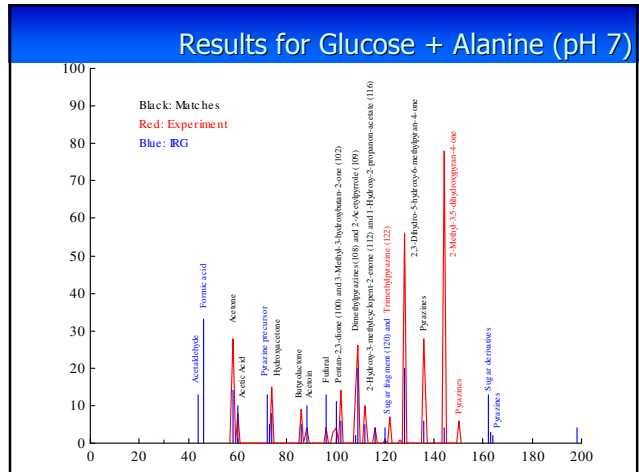
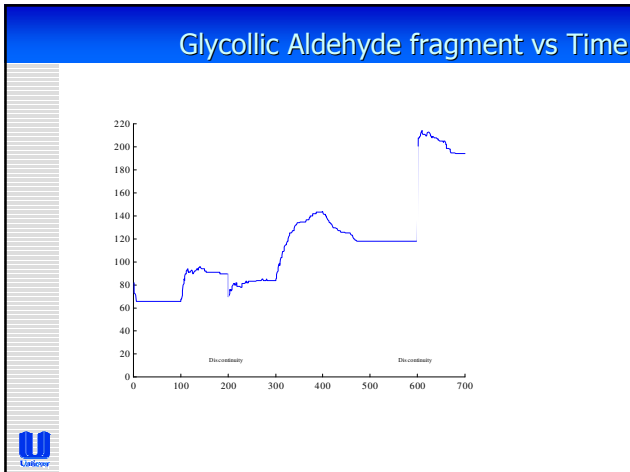
```
1 x C1=O(C)O(NC)C(O)O
1 x C1=CC(O)C(O)C(O)C=O
1 x C1=O
```

- Virtual Mass Spec

- (in time)

- Distance Measure

- to expected products



### Conclusion

- Model and Optimise paradigm is powerful
- Interplay between representation, operators and fitness function is critical
- Multi-criteria optimisation offers choices

**U**  
Unilever