

Evolutionary search for process design

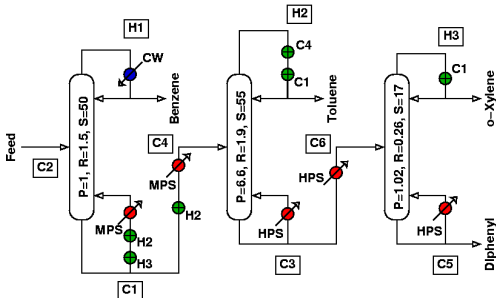
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The 13th CREST Open Workshop
12-13 May 2011

Topic

- 1 Introduction
- 2 Grammar
- 3 Visual
- 4 Conclusions

Process design



- Identify process steps and their interconnections (mass, energy, control) to achieve performance goals.
- Wish to support decision making.
- Exploration necessary for understanding.
- **Model** based.

Optimisation

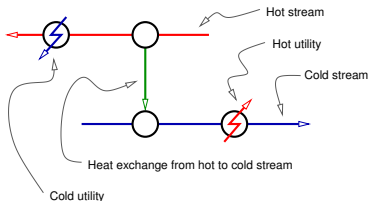
Need robust optimisation methods for process design:

- non-linear, non-convex, and discontinuous models,
- combinatorial search space,
- small, possibly non-convex, feasible regions, and
- ill- or un-defined objective function and constraint equations outside feasible regions.

Evolutionary methods can be a key element in the repertoire of optimisation tools an engineer may use.

Heat exchanger networks (HEN)

Energy consumption is often the largest cost of a process.



One means of reducing energy use is through **process integration**:

Identify matches for transfer of excess heat in one part of the process to another part.

Problem is highly combinatorial, discontinuous and non-convex.

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Evolving a HEN structure

The key to any evolutionary algorithm is the **encoding** of alternative solutions (**designs**):

- We consider a biological analogue:

Genotype the plan

Phenotype the instance

- For HEN, the aim is to use a genotype to represent overall structure: possible matches and stream splits.
- The phenotype is an instantiation of the genotype with specific matches.

We use a Lindenmayer System (L-System) to represent and to evolve genotypes.

L-system definition

An L-system is defined by a tuple,

$$G = \langle V, \omega, P \rangle \quad (1)$$

where

V the alphabet or set of symbols which can be replaced in a string by specific combinations symbols from the same set.

ω the initial configuration (set of strings).

P ($\subset V \times V^*$) is the set of replacement rules.

For the heat exchanger network synthesis problem, we define an L-system, G_{HEN} .

ESF (2009), *Engineering Optimization* 41(9):813-831.

G_{HEN} – 1. Symbols V

The alphabet includes:

- $+$, $-$ denote the cooling and heating requirements of each stream;
- S , E the start and end of each stream;
- x indication of exchange;
- s , m split and mix; and,
 - start and end of split stream segments.
 - $[,]$ start and end of split stream segments

The full alphabet, therefore, is

$$V \equiv \{-, +, S, E, s, m, [,]\} \quad (2)$$

G_{HEN} – 2. Starting representation,

ω

- The starting set of symbols is a **set** of strings, one for each stream in the network problem definition.
- Each cold stream is represented initially by the string S-E.
- Each hot stream by E+S.
- The hot and cold streams are written in opposite order to indicate the use of **counter-current** heat exchangers.

$$\text{For our example, } \omega = \overbrace{\text{E+S}}^{\text{H1}} \overbrace{\text{E+S}}^{\text{H2}} \underbrace{\text{S-E}}_{\text{C1}} \underbrace{\text{S-E}}_{\text{C2}}.$$

G_{HEN} – 3. Rule set, P

Rule	Target	Replacement	Description
R1	-	x-	Add an exchanger to a cold stream
R2	-	s [x-] [x-] m-	Split a cold stream
R3	+	x+	Add an exchanger to a hot stream
R4	+	m] x+ [] x+ [s+	Split a hot stream
R5	S	S	A do-nothing rule

Note: the rule for splitting a hot stream creates a structure that is the reverse of that created by a cold stream split rule.

G_{HEN} summary

- The L-system for HEN design is **context free**.
- It is **nondeterministic**, complementing the evolutionary algorithm.
- The majority of strings (words generated from V^* starting with ω) represent a valid genotype for the HEN design problem.
- The genotype describes a configuration with
 - locations of splits and
 - locations for integrated exchangers.
- Genotype is **instantiated** into a phenotype for an actual design.

Phenotype instantiation

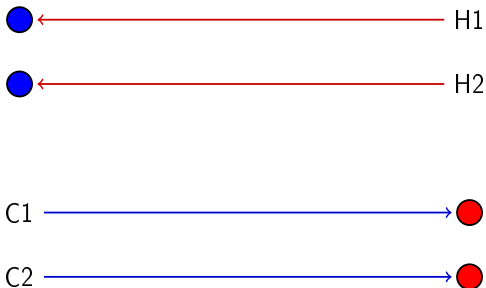
- A genotype describes the overall structure of a HEN.
- To **instantiate** a phenotype from a genotype:
 - 1 link the integrated exchangers and
 - 2 define the appropriate optimisation problem to size exchangers and determine split factors.
- Linking exchangers is **non-deterministic** so a single genotype may lead to different phenotypes.
- The **do-nothing** rule, R5, allows for multiple instances of the same genotype in a population.

Example (re-visited)

Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }

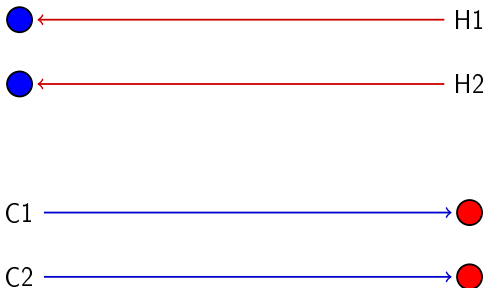
Example (re-visited)

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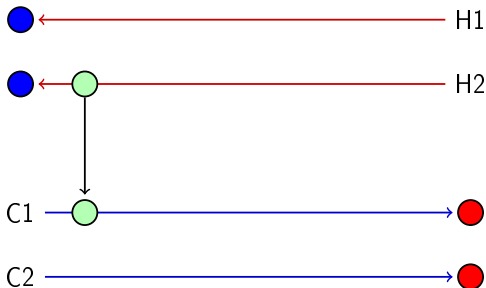
Example (re-visited)

Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }
+R1: { H1:E+S, H2:E+S, C1:Sx-E, C2:S-E; }



Example (re-visited)

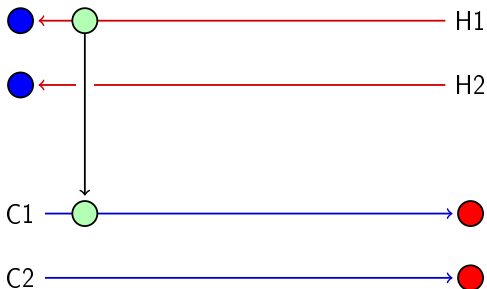
Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }
 +R1: { H1:E+S, H2:E+S, C1:Sx-E, C2:S-E; }



H1:E+S, H2:Ex(C1)+S, C1:Sx(H2)-E, C2:S-E;

Example (re-visited)

Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }
 +R1: { H1:E+S, H2:E+S, C1:Sx-E, C2:S-E; }



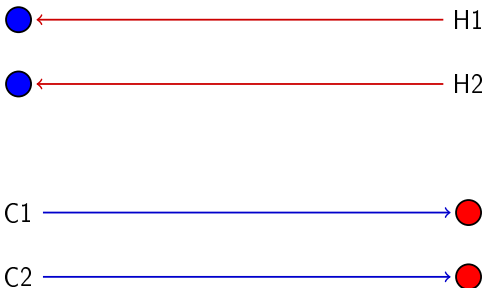
H1:Ex(C1)+S, H2:E+S, C1:Sx(H2)-E, C2:S-E;

Example (re-visited)

Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }

+R1: { H1:E+S, H2:E+S, C1:S_x-E, C2:S-E; }

+R2: { H1:E+S, H2:E+S, C1:S_x-E, C2:S_s[x-][x-]_m-E; }

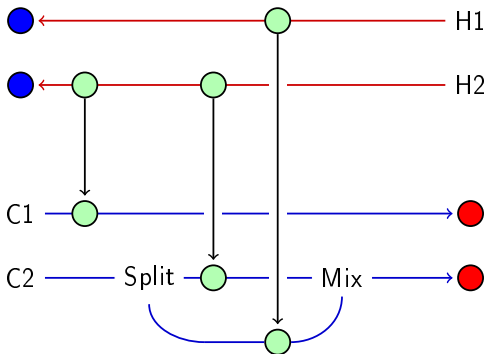


Example (re-visited)

Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }

+R1: { H1:E+S, H2:E+S, C1:Sx-E, C2:S-E; }

+R2: { H1:E+S, H2:E+S, C1:Sx-E, C2:Ss[x-][x-]m-E; }



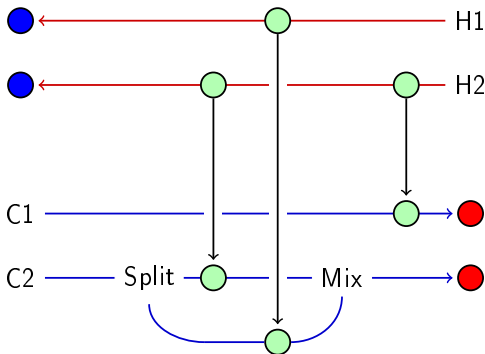
H1:Ex(C2)+S, H2:Ex(C1)x(C2)+S, C1:Sx(H2)-E, C2:Ss[x(H2)-][x(H1)-]m-E;

Example (re-visited)

Initial: { H1:E+S, H2:E+S, C1:S-E, C2:S-E; }

+R1: { H1:E+S, H2:E+S, C1:Sx-E, C2:S-E; }

+R2: { H1:E+S, H2:E+S, C1:Sx-E, C2:Ss[x-][x-]m-E; }

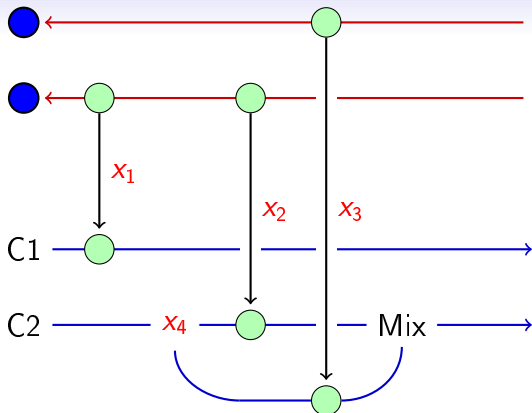


H1:Ex(C2)+S, H2:Ex(C2)x(C1)+S, C1:Sx(H2)-E, C2:Ss[x(H2)-][x(H1)-]m-E;

Embedded optimisation problem

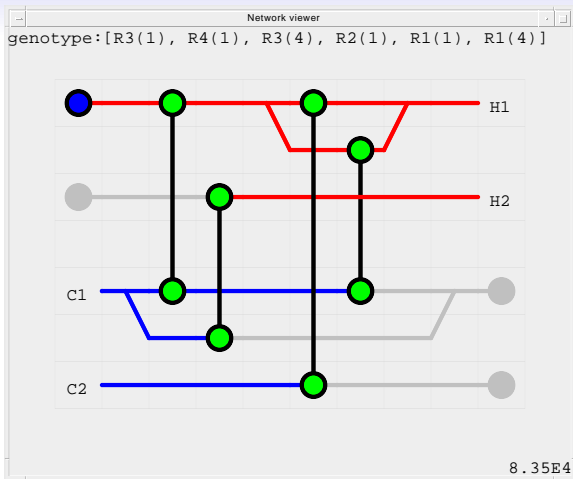
Phenotype \Rightarrow network
 structure \Rightarrow a nonlinear
 programme with decision
 variables:

- 1 $x_i \in [0, 1]$, the fraction to exchange:
 $Q_i = x_i Q_{i,\max}$
- 2 $x_j \in [0, 1]$, the split fraction.

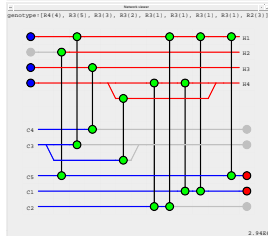
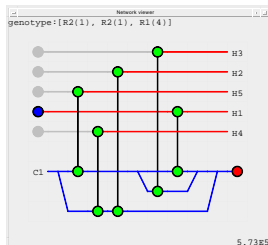
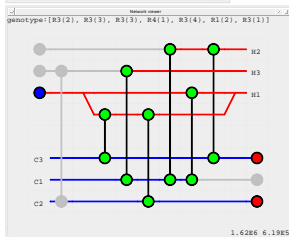
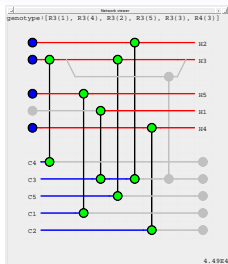


The NLP represents a superstructure and is solved using a hybrid stochastic & direct search procedure.

Example (solution)



Other case studies



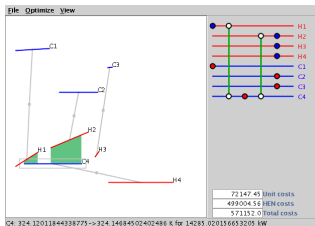
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Visual representation

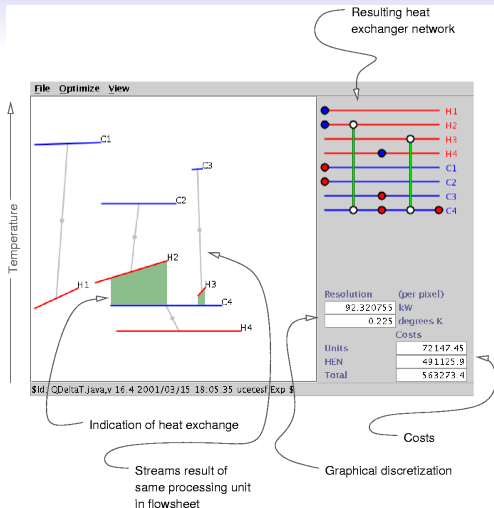
For a given process configuration, we can display the hot and cold streams visually and support interaction, where

- x-axis for position independent duties,
- y-axis for temperature, and
- hot stream overlapping cold stream indicates heat integration.



Allow user to manipulate process by moving streams (the **tail wagging dog** approach): streams can be moved horizontally for different integrations and moved vertically or stretched horizontally to change underlying unit designs.

Demonstration of interaction



- Interface encourages exploration and allows for insight.
- However, space is complex so best solutions are not always immediately obvious.

Discrete design representation

Representation

A graphical view of process heat requirements defines left and right end-points for each hot and cold stream in the process:

$$\{(x_{a,i}, y_{a,i})\}$$

$$\{(x_{b,i}, y_{b,i})\}$$

$i = 1, \dots, n_s$ and
 $x, y \in \mathbb{Z}^+$, suitable for
 manipulation by
 evolutionary algorithms.

- 1 Define list of intervals

$$I \leftarrow \bigcup_1^{n_s} \{\{x_{a,i}\} \cup \{x_{b,i}\}\}$$

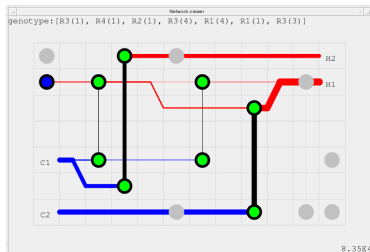
- 2 For each interval $[l_j, l_{j+1}]$:
 - 1 Generate list of active streams, A .
 - 2 Sort A from top to bottom using y_b values.
 - 3 Generate match for each hot stream immediately above cold stream in A .
 - 4 Generate utility match for all other streams.
- 3 Coalesce adjacent similar matches.
- 4 Design exchanger for each match.
- 5 Cost all exchangers and utility use.

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Summary

- Optimization problems in design are typically complex.
- Evolutionary methods can provide a useful tool for design.
- The key is the representation of possible solutions.



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<http://www.homepages.ucl.ac.uk/~ucecesf/>