



A Multi-objective Genetic Algorithm

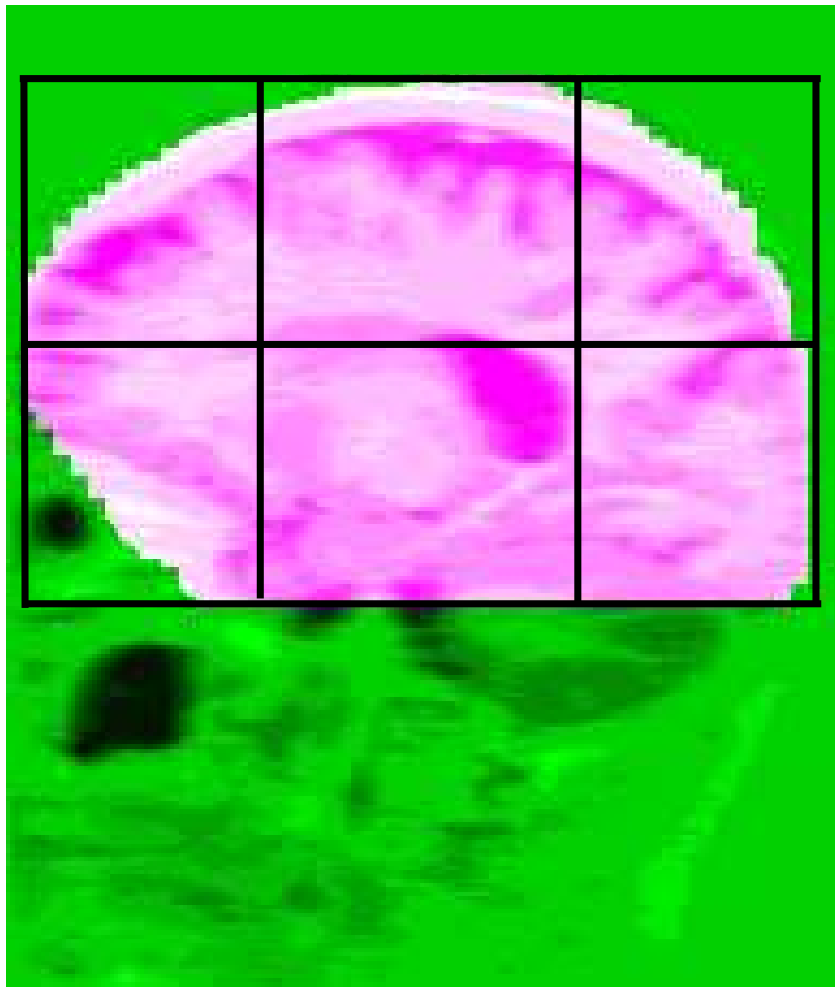
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Imaging Science and Biomedical Engineering

Overview

- Diagnosis of dementing diseases using the distribution of cerebral atrophy
- Optimisation of the technique using a genetic algorithm
 - preselection and static extraction
 - novel mating restriction
 - self adaptive mutation parameters
 - a multi-objective cost function test suite
- Current state of the project

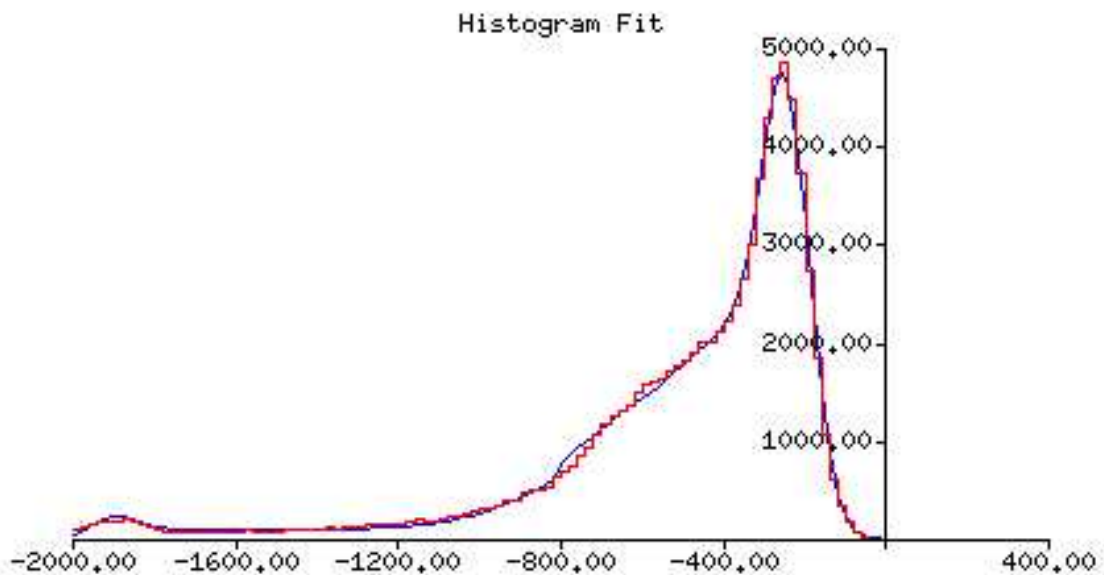
Atrophy Analysis: Introduction

- Original technique developed by Thacker et. al. (2002) Radiology 224 p278-285
 - hypothesis: dementing diseases result in specific patterns of cerebral atrophy
 - use minimal parameter atrophy measurement
 - co-register to a standard coordinate system
 - divide head into twelve equi-sized boxes



Atrophy Analysis: CSF Segmentation

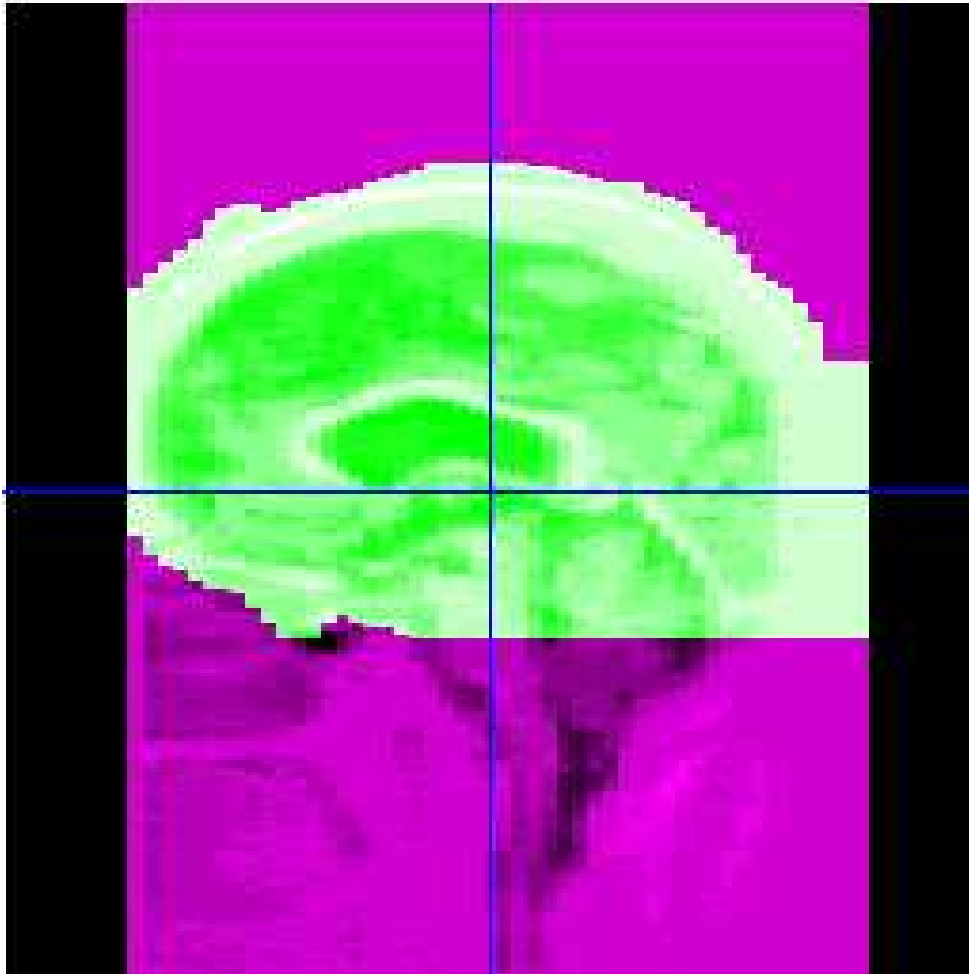
- Segment the CSF
 - use T2 IR scans
 - plot histogram of interior of head
 - fit with Gaussians + triangular distributions convolved with Gaussians



- place threshold at CSF probability = 0.5
- produce binary CSF maps

Atrophy Analysis: Masking

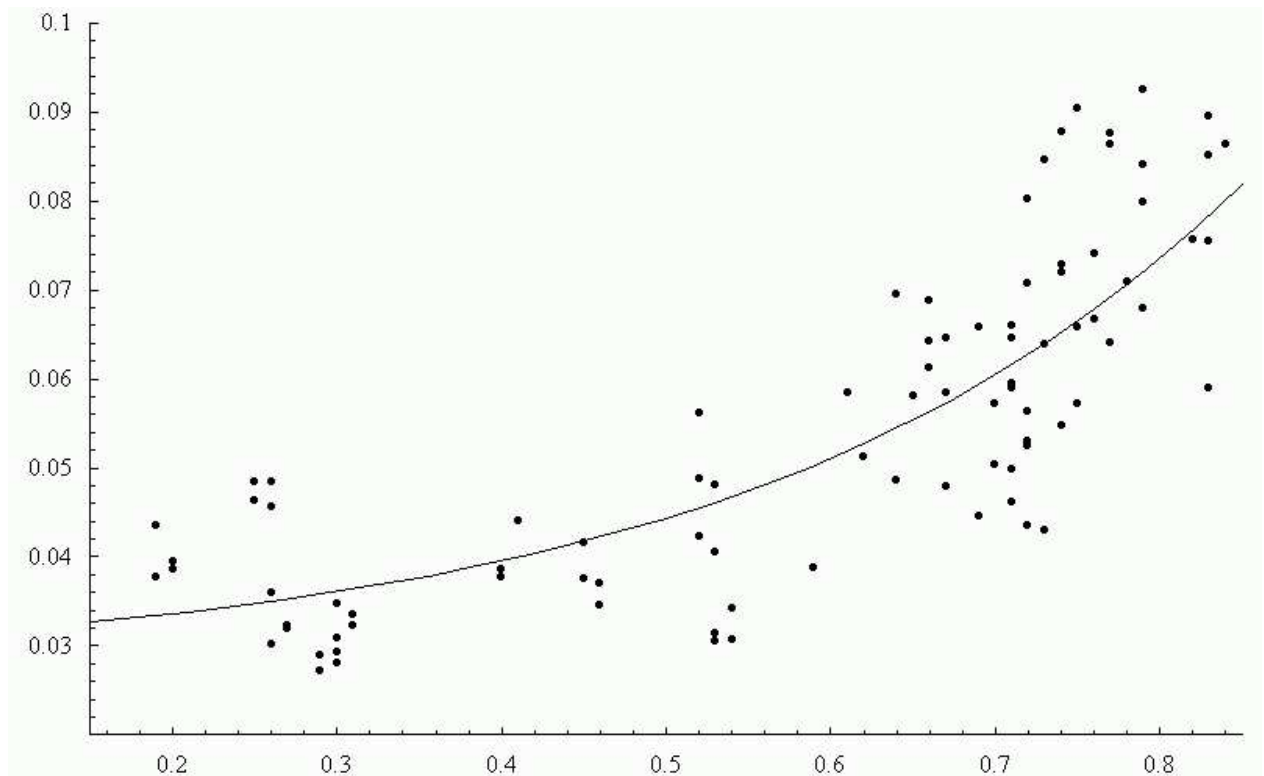
- Binary images include all fluid spaces
 - CSF, eyes, sinuses
- Delete eyes, sinuses etc. using binary masks



- Masks also define lower extent of the space: all other limits defined by outside of head

Atrophy Analysis: Age Correction

- Count CSF in the twelve boxes
- Normalise for head size (divide by total volume)
- Normalise for age-related atrophy



- Age normalisation
 - 94 normals: age range 19 to 85
 - fit curves

$$volume = a + b \exp^{c \text{ age}}$$

- normalise to mean age point

Atrophy Analysis: Variable Reduction

- Calculate reduced variables
 - F = sum front four volumes
 - M = sum middle four volumes
 - B = sum back four volumes
 - P = sum left six volumes
 - S = sum right six volumes
 - U = sum top six volumes
 - L = sum bottom six volumes
- Calculate relative variables

$$W_1 = \frac{\sqrt{M} - \sqrt{F}}{\sqrt{2}}$$

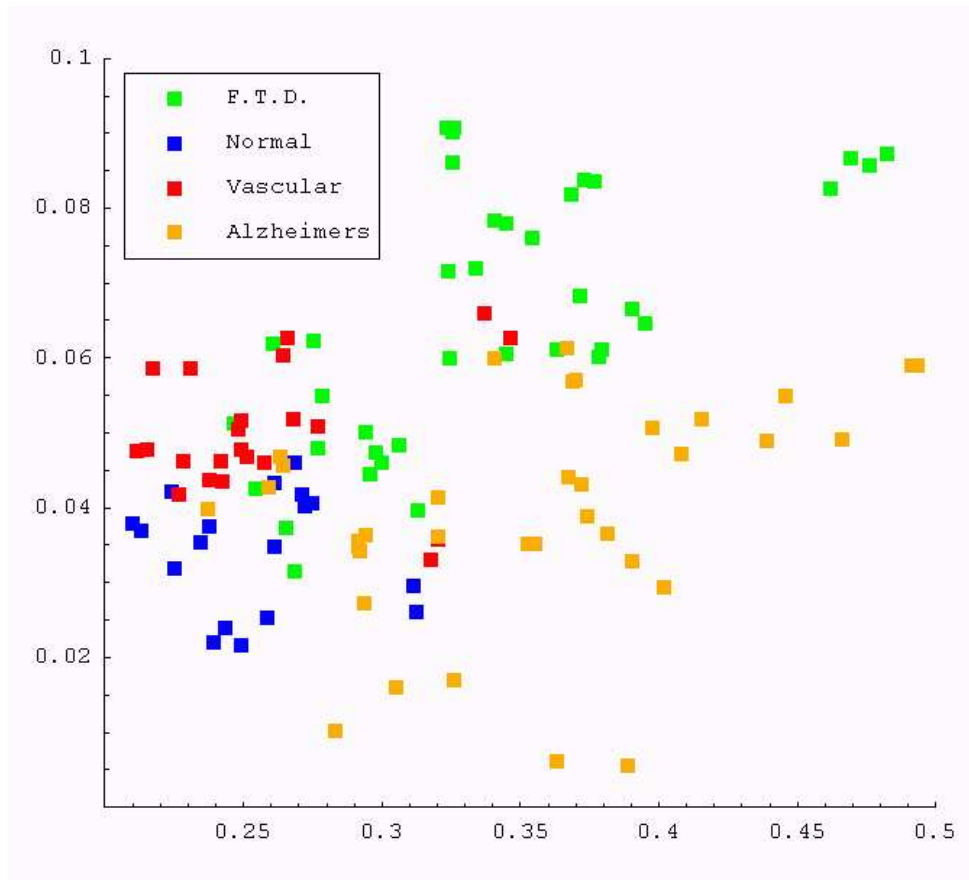
$$W_2 = \frac{\sqrt{M} - \sqrt{B}}{\sqrt{2}}$$

$$W_3 = \frac{\sqrt{F} + \sqrt{M} + \sqrt{B}}{\sqrt{3}}$$

$$W_4 = \frac{\sqrt{P} - \sqrt{S}}{\sqrt{2}}$$

$$W_5 = \frac{\sqrt{U} - \sqrt{L}}{\sqrt{2}}$$

Atrophy Analysis: Results



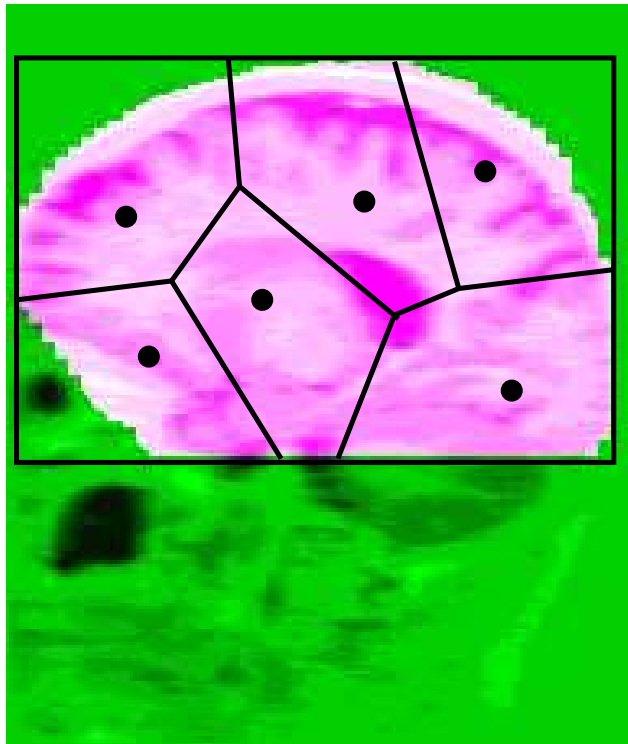
W_3 vs. W_2

- Cross-validated Parzen classifier results

Diagnosis	Normal	FTD	Vascular	Alz.
Normal	7	2	8	1
FTD	5	21	3	7
Vascular	3	2	13	4
Alz.	1	3	6	28

Motivation for GA Optimisation

- CSF boxes not related to brain structure
 - sub-optimal e.g. FTD often produces degeneration of frontal lobes
- Optimise boundary positions
 - replace boundaries with centre points
 - tessellate space with Voronoi cells



- point pairs form natural building blocks
- cost function noisy, form unknown
- use GA

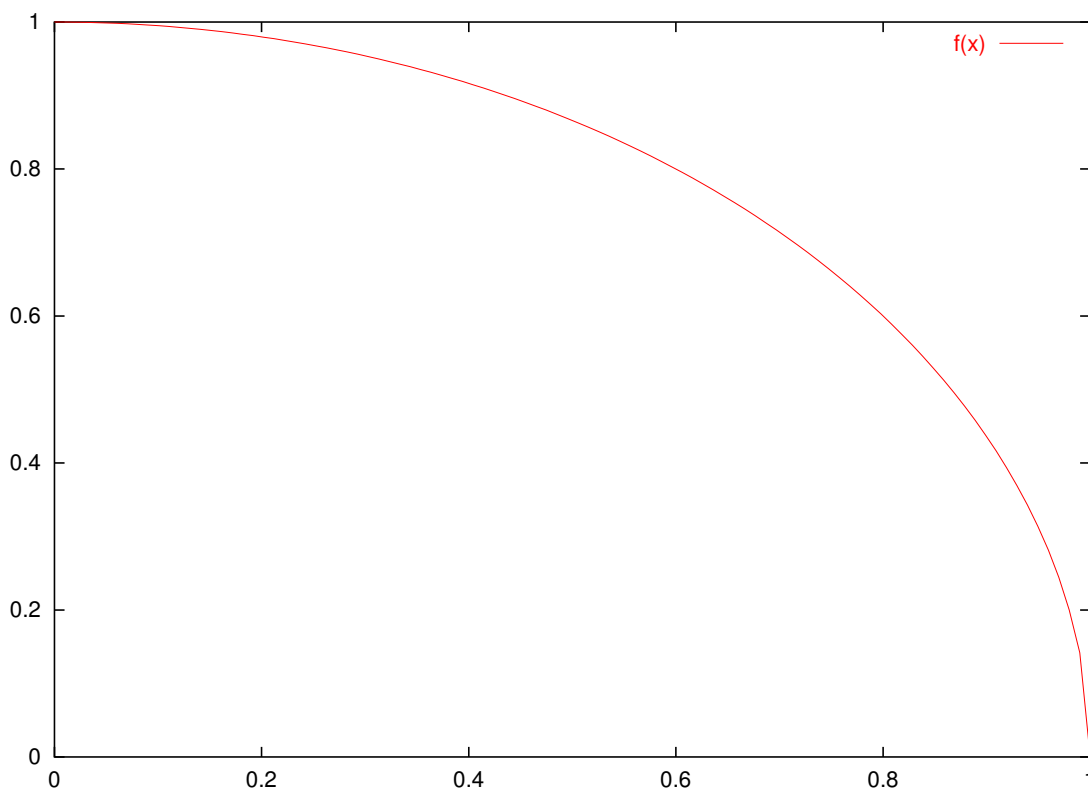
GA Goals

- Multi-objective cost function
 - use diagonal elements of confusion matrix
 - diversity preservation essential
- Diversity preservation
 - global convergence will occur if any pair of solutions can interact
 - nature: strict mating restrictions (inter-species)
+ dynamic cost function (intra-species)
 - convergence to occur locally, but not globally
- Minimise free parameters
 - population size, mutation probabilities
 - diversity preservation strategy parameters (e.g. crowding factor)

Multi-objective Fitness Evaluation

- Non-Pareto approaches
 - e.g. Schaffer (1985) VEGA
 - linear weighting of objectives
 - does not cover concave regions of cost surface
- E.g. minimise x and y for

$$y \geq \sqrt{1 - x^2}$$



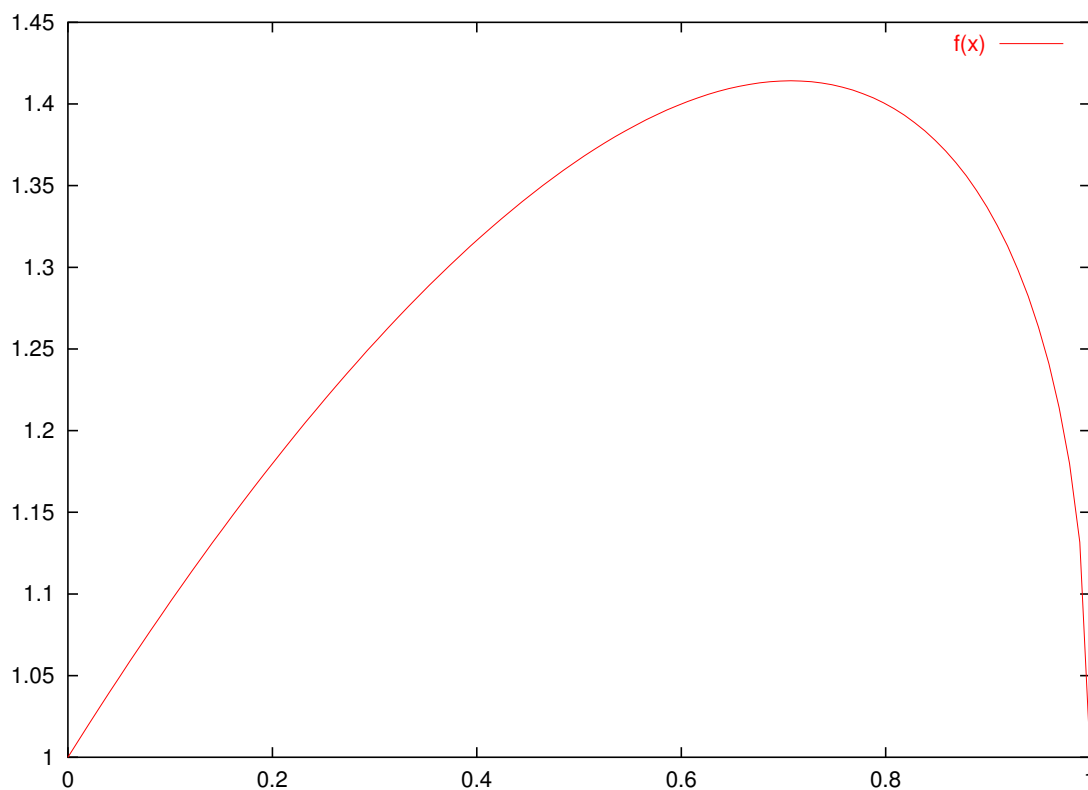
- cost is linear weighting of x and y

$$f = \alpha x + \beta y$$

Multi-objective Fitness Evaluation 2

- For solutions on the Pareto front

$$f = \alpha x + \beta \sqrt{1 - x^2}$$



f vs. x for $\alpha = 1$ and $\beta = 1$

- Only the ends of concave regions of the cost surface will be found

Multi-objective Fitness Evaluation 3

- Pareto-based approaches
 - e.g. Goldberg (1989) Pareto ranking
 - find non-dominated solutions \rightarrow rank 1
 - remove rank 1 solutions
 - iterate \rightarrow rank 2, 3...
 - probability of reproduction based on rank
- Fitness evaluated purely on dominance basis

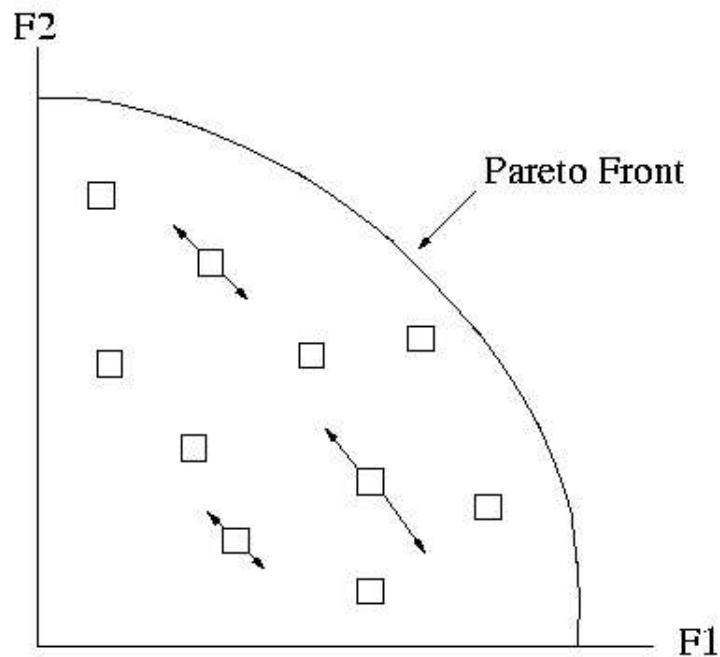
The Larcombe GA: Reproduction

- Larcombe (1996) Ph.D. thesis: GA optimisation of chip layer partitioning problem
- Avoid fitness proportional selection
 - select single pair of solutions per iteration
 - fitter offspring replace parents in a first-come, first-served competition
 - c.f. Cavicchio's preselection
- What about the Fundamental Theorem?

$$m(H, t + 1) \geq m(H, t) \frac{f(H)}{\bar{f}} \dots$$

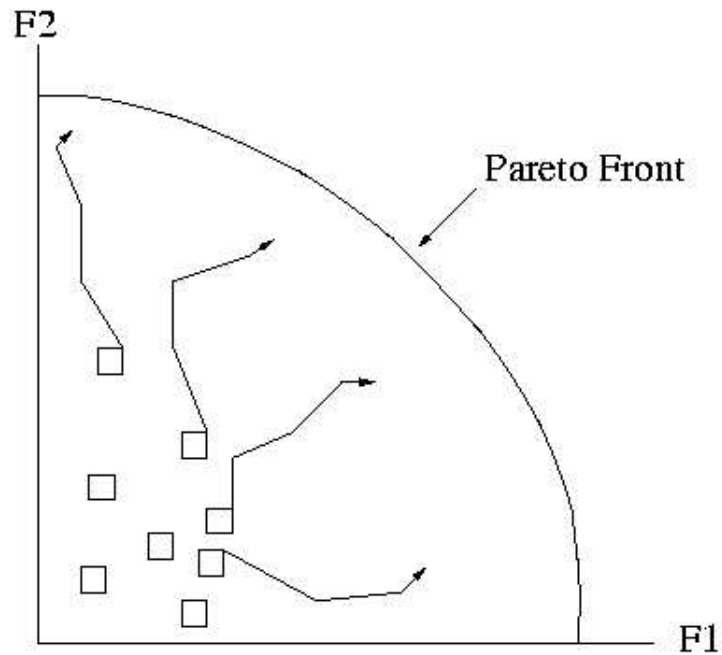
- schemata can double per iteration
 - ⇒ quadratic growth
- c.f. fitness scaling with scaling factor = 2

Fitness and Direction Vectors



- Parent-offspring pairs can continually replace each other
 - oscillatory behaviour
- Add direction flags to genome
 - one flag for each objective: initialised at 0
 - child replaces parent if it wins on any one objective
 - children of that child must win on the same objective
 - more directions can be set to 1 at any time

Static Extraction



- Solutions can become stuck at local minima
 - super-fit individuals: *statics*
- Remove these after n failed reproduction attempts
 - place in a secondary population:
static extraction
 - replace with *random immigrants*
 - population size effectively infinite
- Diversity preservation + free parameter reduction

Coding and Mutation

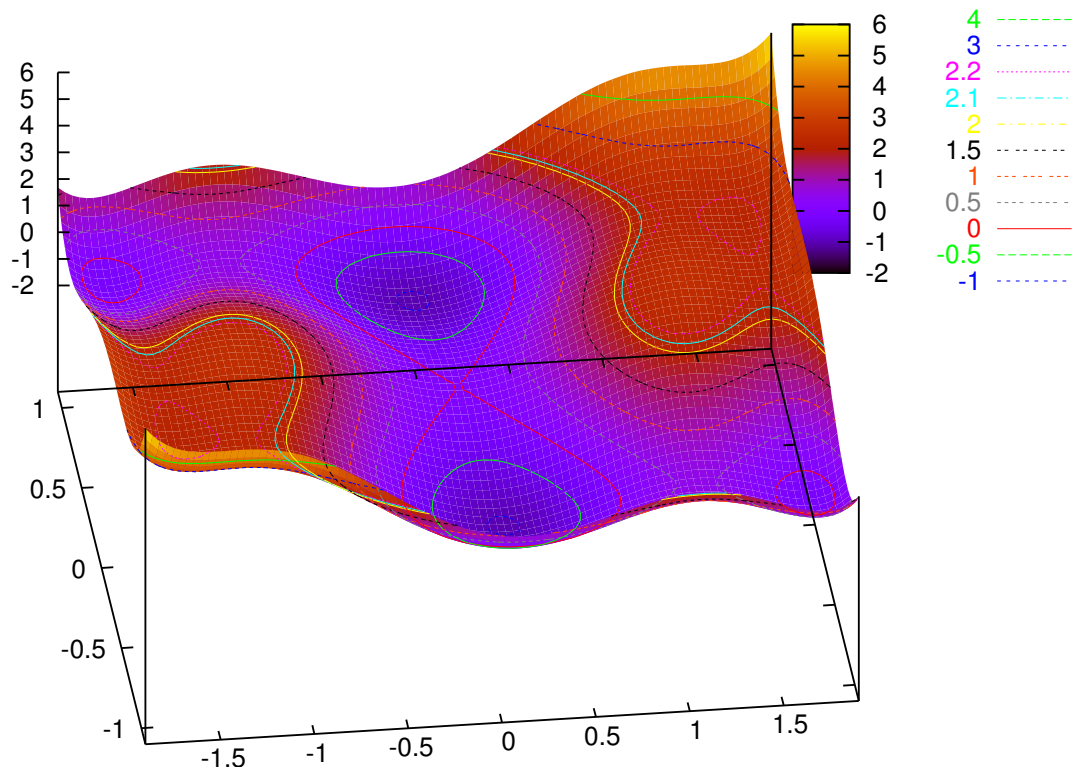
- Larcombe: binary coding
- Atrophy analysis: list of 6 real-valued points
 - point pairs form natural building blocks
 - distance parameter required for mutation
- Self-adaptive mutation
 - add mutation parameters to genome as a header
 - points: probability and fractional distance
 - header: probability and probability change
 - allow mutation probabilities to mutate
- Previous work (e.g. Cavicchio, 1970)
 - centralised control

Crossover

- For atrophy analysis
 - treat each point as a single gene
 - implement single point crossover
- For testing on simple functions
 - code a single point on a test function
 - new crossover routine required
- Previous work
 - Ono, Kita and Koboyashi, GECCO 1999
 - many techniques e.g. place children randomly on line between parents
 - adopt simple method: place children randomly in hypercube defined by parents

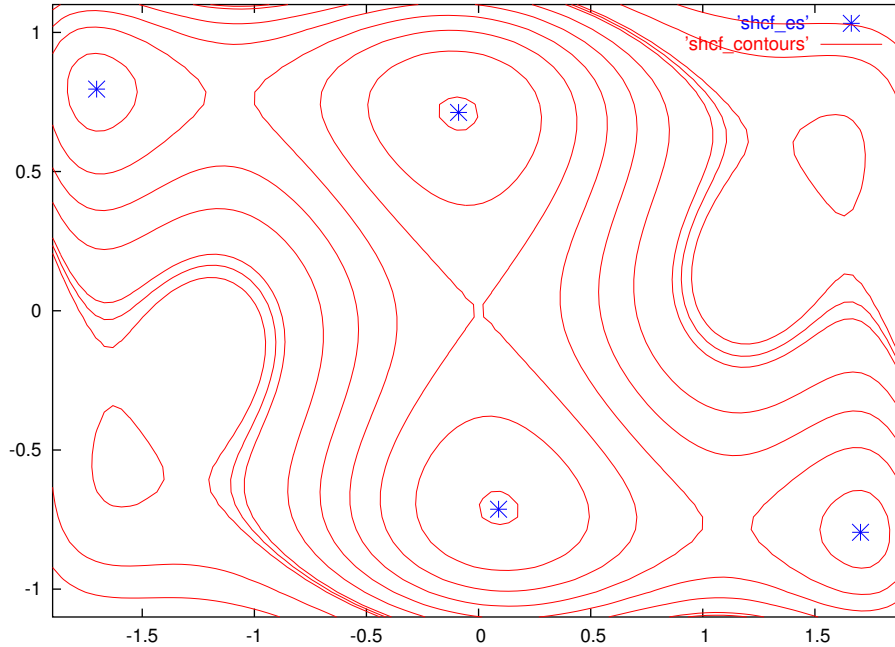
Initial Testing

- Implementation:
 - Larcombe GA: static extraction, random immigration
 - additions: real-valued representation, self-adaptive mutation
 - one free parameter left: extraction age
- Initial testing: six-hump camelback function

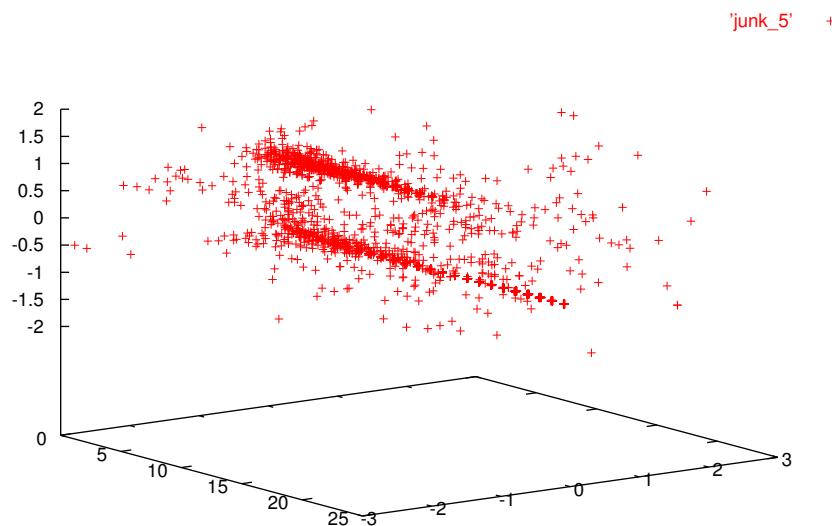


Initial Testing 2

- Extracted set at generation 25:



- General population:



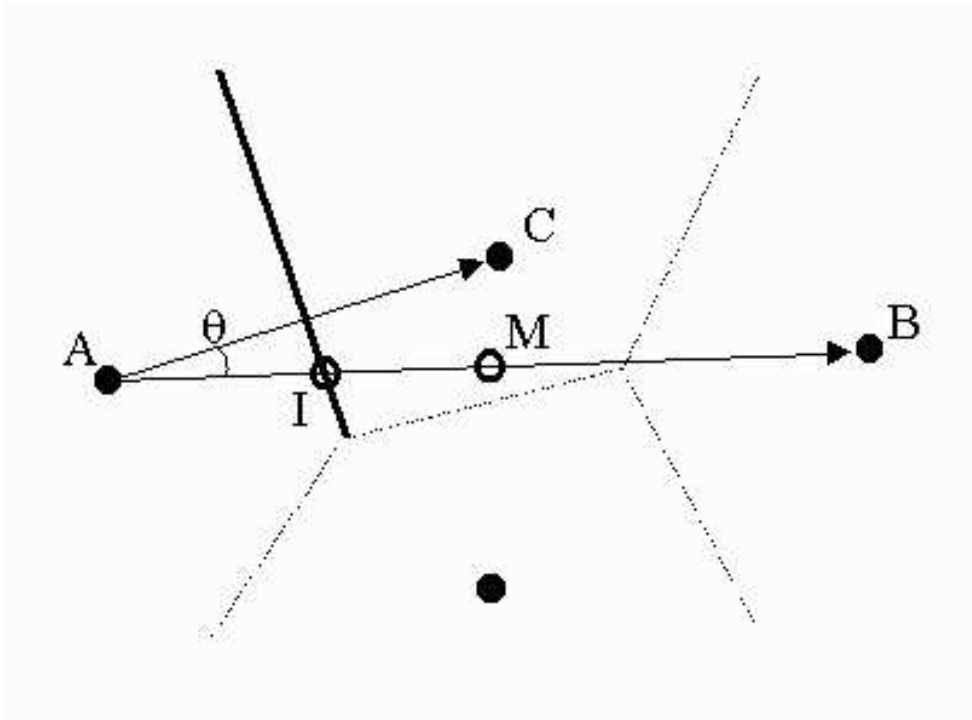
– diversity loss due to random selection

Diversity Preservation

- c.f. fitness scaling
 - start of run: super-fit individuals
 - end of run: lack of fitness differentiation
 - multi-modal function: loss of global optima
- Larcombe GA
 - static extraction limits super-fit individuals
 - global optima still lost at end of run
- Additional diversity preservation required
 - add mating restriction
 - aim: population convergence should be local but not global
 - convergence in extracted set instead of population

Voronoi Mating Restriction

- Avoid additional parameters: use algorithm's definition of local vs. global
 - restrict mating to Voronoi neighbours



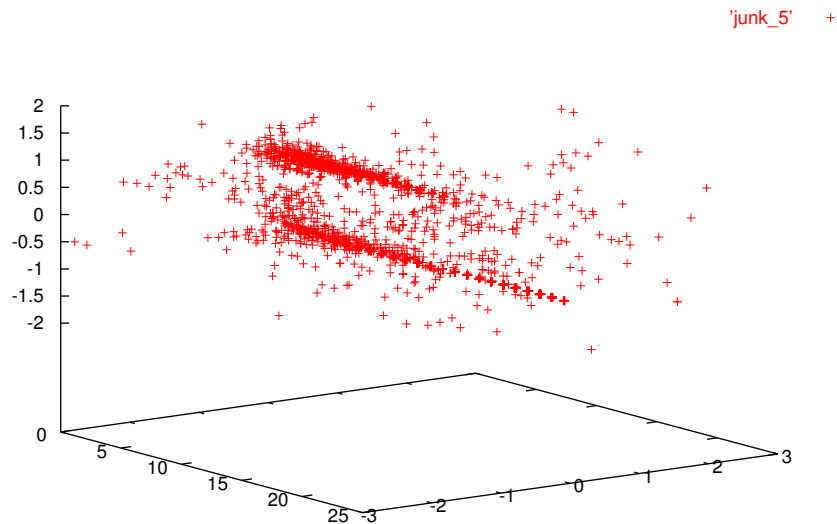
- rejection criterion

$$|\mathbf{AM}| > |\mathbf{AI}| \Rightarrow \frac{|\mathbf{AC}|^2}{\mathbf{AB} \cdot \mathbf{AC}}$$

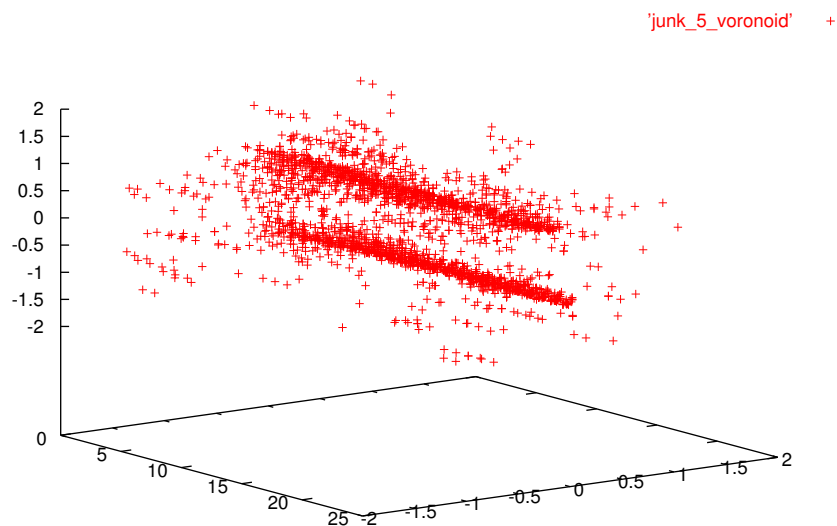
- During selection
 - randomly select parent 1
 - randomly select parent 2 from the Voronoi neighbours

Influence of Mating Restriction

- General pop. without mating restriction



- General pop. with mating restriction



- Diversity preserved indefinitely

Multi-Objective Testing

- Use seven function test-function set from Van Veldhuizen (1999): Pareto fronts
 - connected/unconnected in search space
 - connected/unconnected in cost space
 - convex/concave
 - single/multiple Pareto curves
- Use test procedure/metrics from Van Veldhuizen (2000)
 - find true Pareto front using exhaustive search
 - run GA for 2^{16} cost function evaluations
 - 0.39% of space covered
 - generational distance

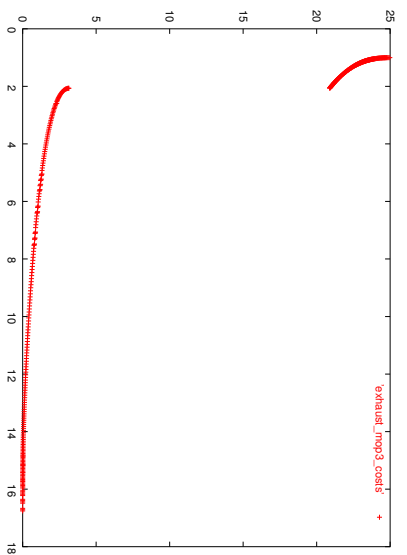
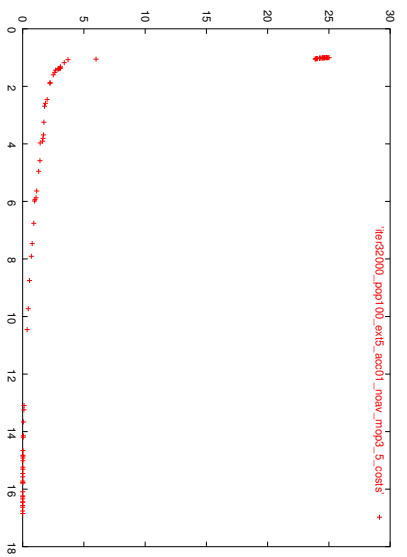
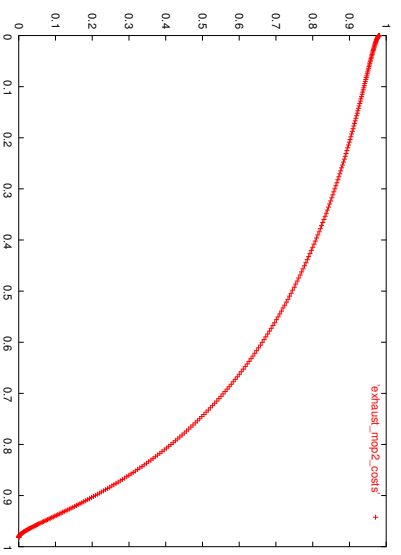
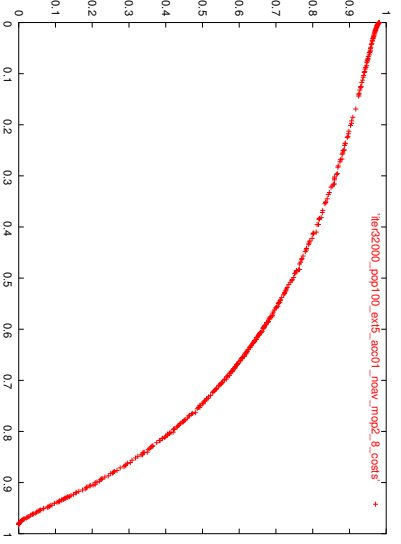
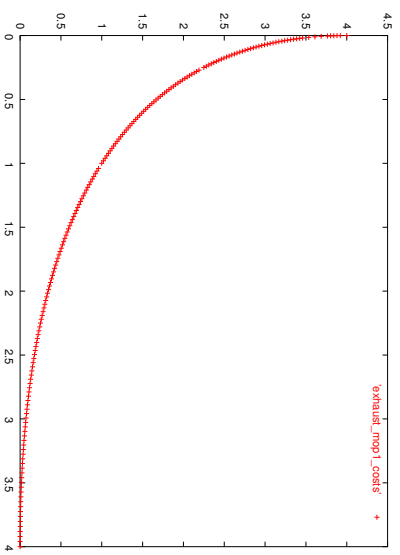
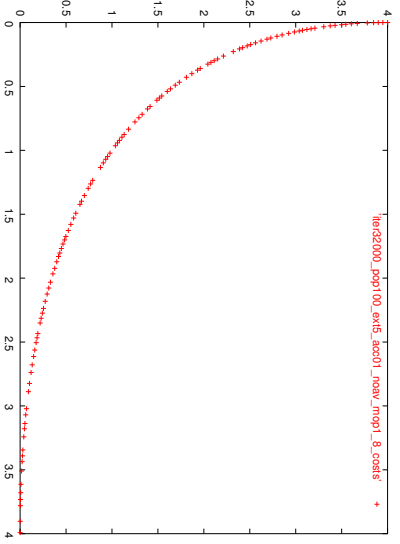
$$G = \frac{\sqrt{(\sum_{i=1}^n d_i^2)}}{n}$$

d_i = distance to true Pareto front

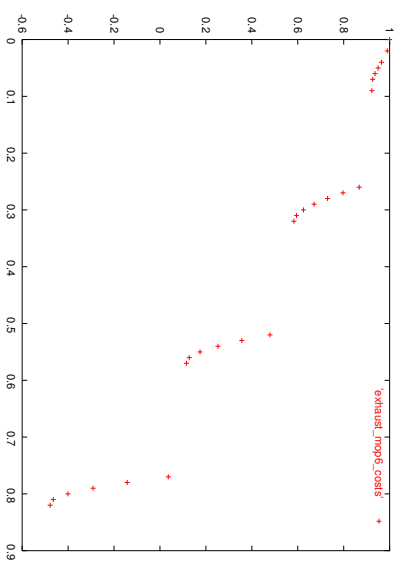
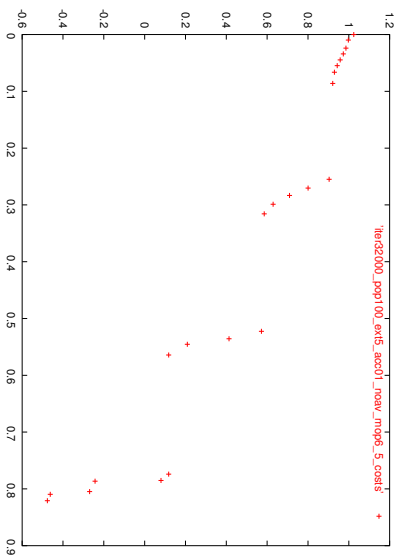
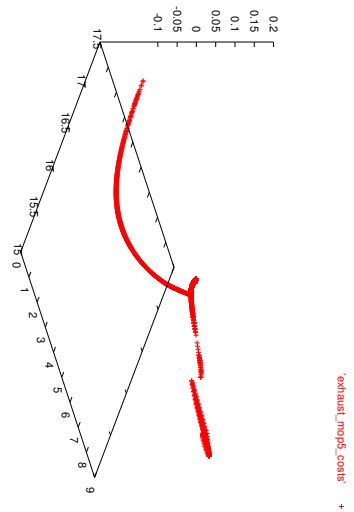
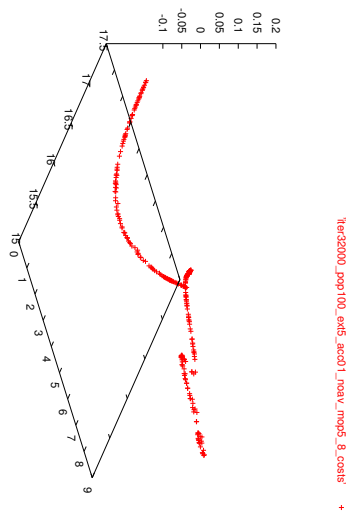
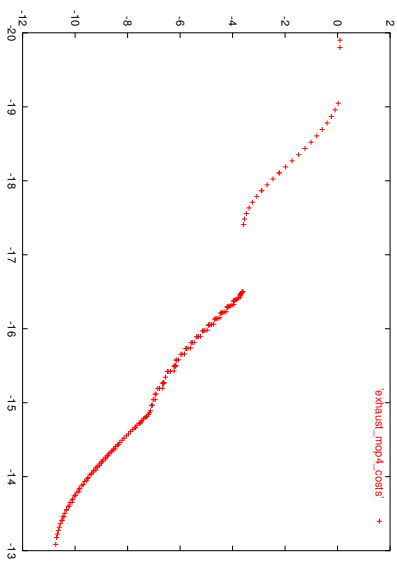
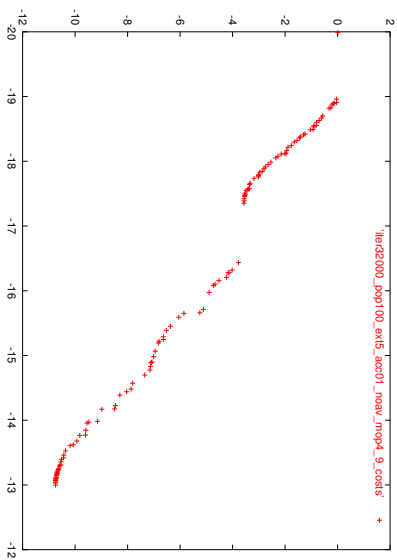
- ONVG = no. of solutions

Multi-Objective Results

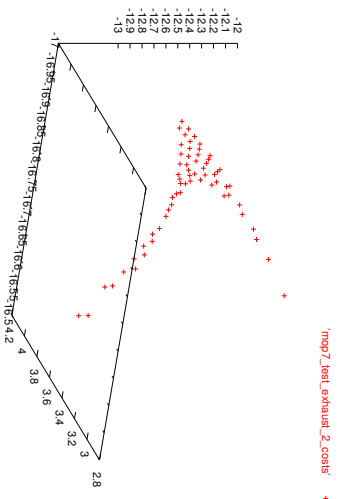
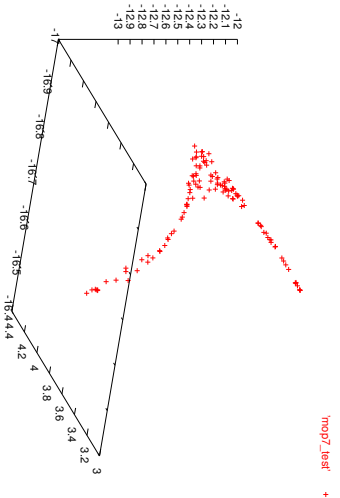
- Plots of GA results and exhaustive search in cost space



Multi-Objective Results

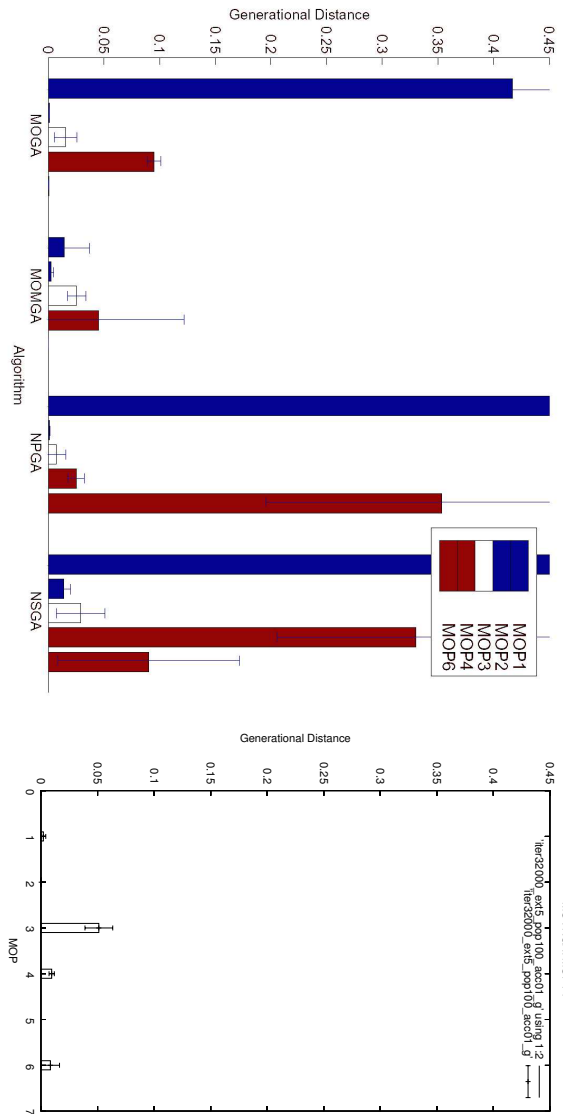


Multi-Objective Results

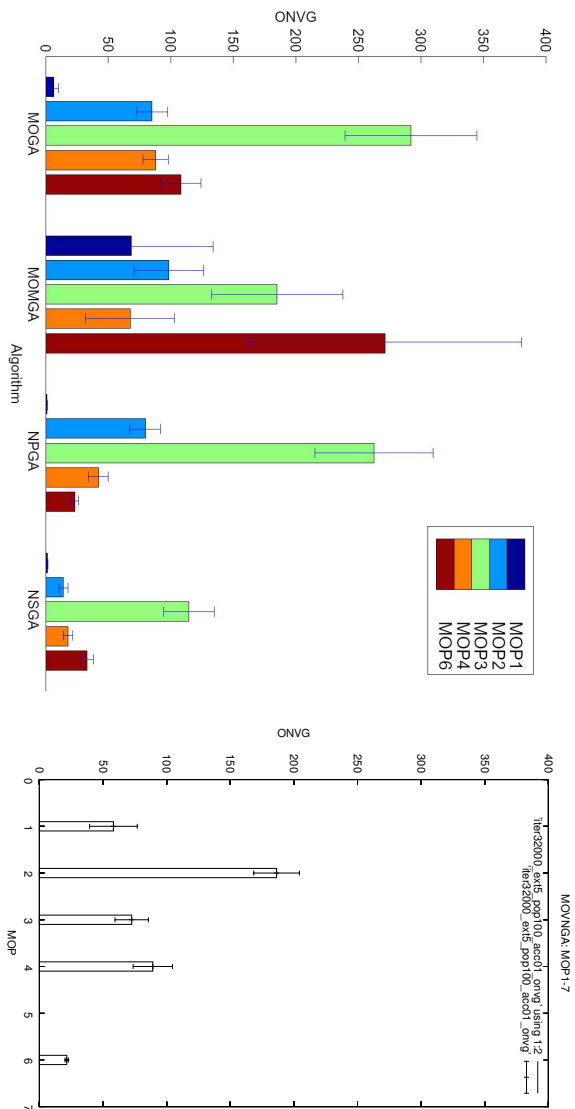


- Pareto front found for all 7 test functions
- Now compute G and ONVG
 - average over 10 GA runs for each test function
 - compare to GAs tested by Van Veldhuizen
MOGA (Fonaseca and Fleming, 1998)
MOMGA (Van Veldhuizen and Lamont, 2000)
NPGA (Horn et al, 1994)
NSGA (Srinivas and Deb, 1994)
 - all written to test aspects of GA theory
 - all highly cited
 - good competition

Multi-Objective Results



- Generational distance (lower is better)



- ONVG (higher is better)

Conclusions

- Novel GA
 - no fitness proportional reproduction
 - static extraction
 - random immigration
 - real-valued representation
 - self-adaptive mutation parameters
 - novel mating restriction (Voronoi neighbours)
- Testing
 - Van Veldhuizen’s test function collection and metrics
 - comparable performance to MOMGA

Further Work

- Progress
 - GA written / tested on simple cost functions
 - atrophy analysis software (almost) finished
 - patient data collected and quality controlled
 - 200 sets / 6 groups
- Future work
 - finish age correction
 - more self-adaptation testing
 - test GA on complex cost functions (TSP?)
 - relate back to GA theory
 - parallelise GA