

#### Genetic Algorithm and Direct Search Toolbox

V2.3 (R2008a)

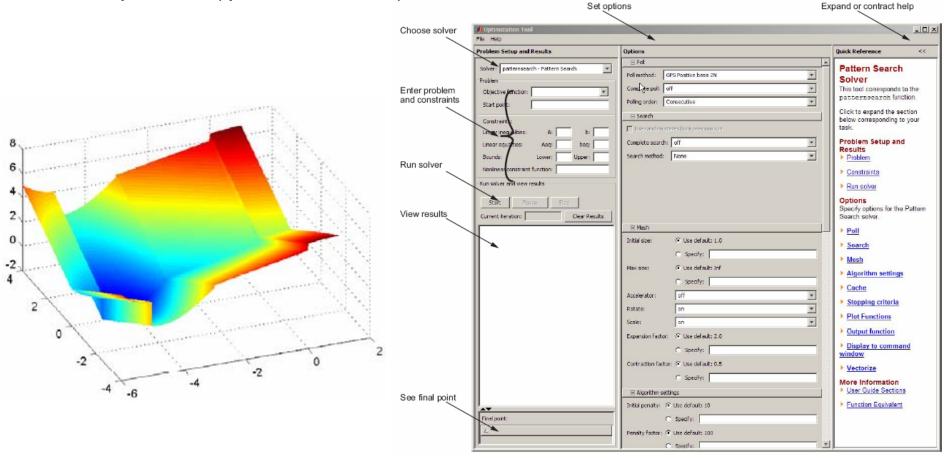
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## **Direct Search**

- method for solving optimization problems that does not require any information about the gradient of the objective function
- searches a set of points around the current point, looking for one where the value of the objective function is lower than the value at the current point
- solves problems for which the objective function is not differentiable, stochastic, or even continuous
- generalized pattern search (GPS) algorithm
- mesh adaptive search (MADS) algorithm

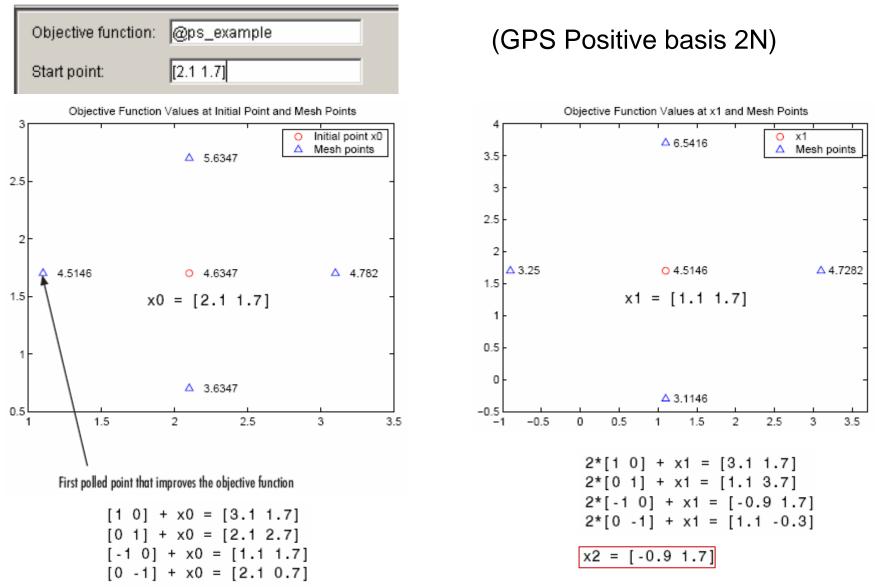
### **DS: Function Call**

- [x fval exitflag output] =patternsearch(@objfun, x0, A, b, Aeq, beq, lb, ub, nonlcon, options)
- optimtool('patternsearch')



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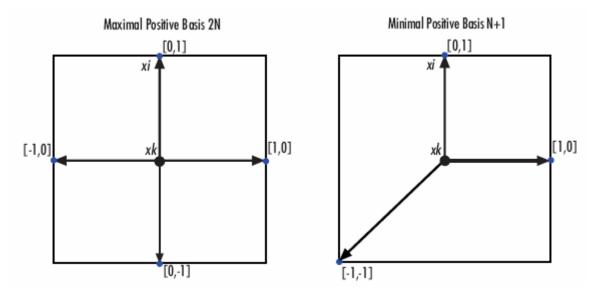
#### How Pattern Search Works



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# DS: Terminology

• Pattern: direction



- Mesh: step size (expansion/contraction factor)
- Polling: finds a point whose objective function value is less than that of the current point

### DS: Options (1)

– options = psoptimset(@patternsearch)

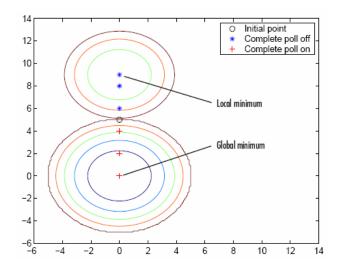
options =

TolMesh: 1.0000e-006 TolCon: 1.0000e-006 TolX: 1.0000e-006 TolFun: 1.0000e-006 TolBind: 1.0000e-003 MaxIter: '100\*numberofvariables' MaxFunEvals: '2000\*numberofvariables' TimeLimit: Inf MeshContraction: 0.5000 MeshExpansion: 2 MeshAccelerator: 'off' MeshRotate: 'on' InitialMeshSize: 1 ScaleMesh: 'on' MaxMeshSize: Inf InitialPenalty: 10 PenaltyFactor: 100

PollMethod: 'gpspositivebasis2n' CompletePoll: 'off' PollingOrder: 'consecutive' SearchMethod: [] CompleteSearch: 'off' Display: 'final' OutputFcns: [] PlotFcns: [] PlotFcns: [] PlotInterval: 1 Cache: 'off' CacheSize: 10000 CacheTol: 2.2204e-016 Vectorized: 'off' UseParallel: 'never'

# DS: Options (2)

- Poll Method
  - GPS/MADS + Positive basis 2N/Np1
- Complete Poll: Off/On
- Using a Search Method
- Mesh Expansion and Contraction
- Mesh Accelerator
  - When the mesh size is below a certain value
- Using Cache
  - Store a history, eliminate redundant computations
- Setting Tolerances for the Solver
  - Mesh, X, function, nonlinear constraint, bind
- Constrained Minimization Using pattern search



## Genetic Algorithm

- method for solving both constrained and unconstrained optimization problems that is based on natural selection, the process that drives biological evolution
- solves a variety of optimization problems that are not well suited for standard optimization algorithms, including problems in which the objective function is discontinuous, nondifferentiable, stochastic, or highly nonlinear

Classical Algorithm	Genetic Algorithm
Generates a single point at each iteration. The sequence of points approaches an optimal solution.	Generates a population of points at each iteration. The best point in the population approaches an optimal solution.
Selects the next point in the sequence by a deterministic computation.	Selects the next population by computation which uses random number generators.

#### GA: Function Call

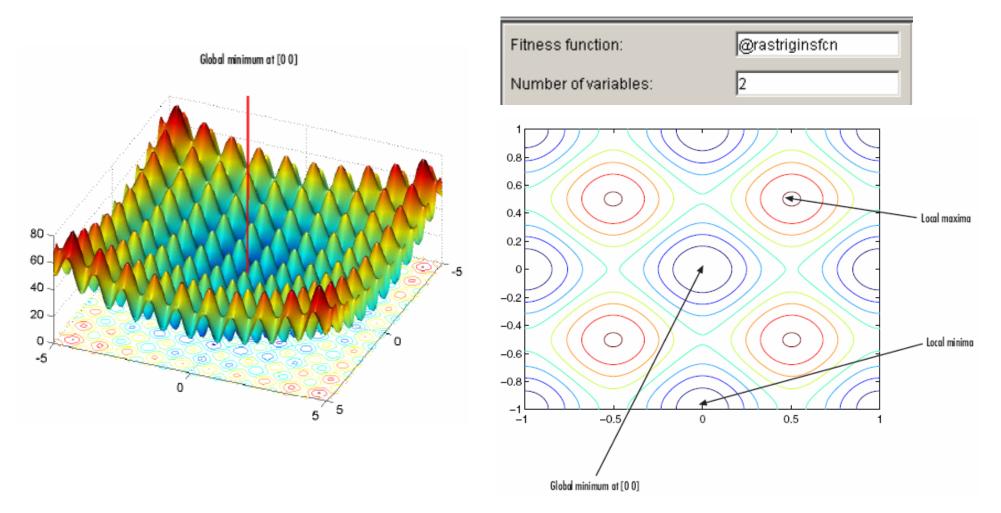
- [x fval exitflag output population scores] = ga(@fitnessfun, nvars, A, b, Aeq, beq, lb, ub, nonlcon, options)
- optimtool('ga')

	Setoptions		Expand or contract help
Choose solver	->> Optimization Tool		×
Choose solver	File Help		<b>X</b>
	Problem Setup and Results	Options	Quick Reference <<
	Solver: ga - Genetic Algorithm	Population type: Double Vector	Genetic Algorithm
Enter problem and constraints	Fitness function:	Population size:  C Use default: 20 C Specifiv:	This tool corresponds to the ga function.
	Constraints: Linear inclusities: A: b:	Creation function: Use constraint dependent default	Click to expand the section below corresponding to your task.
	Linear equalities: Aeq: beq: Bounds: Lower: Upper:	Initial population:  Use dafault:  Specfy:	Problem Setup and Results Problem
	Noninea constraint function:	Initial scores: ( Use default: []	<u>Constraints</u>
Run solver	-Run solver and view results	C Specfy:	Run solver
	Use random states from previous run Start Pause Stop	Initial range: @ Use default: [0;1] C Specify:	Options Specify options for the Genetic Algorithm solver.
	Current iteration: Clear Results	Fitness scaling	Population
View results		Scaing function: Rank	• Fitness scaling
	$\mathbf{k}$		<u>Selection</u>
			Reproduction
		Selection	• Mutation
		Selection Function: Stochastic uniform	<u>Crossover</u>
			<u>Migration</u>
See final point			Algorithm settings
		Elte count: C Use default: 2	Hybrid function
		C Specify:	Stopping criteria
	Final point:	Crossover fraction: C Use default: 0.8	Plot Functions
	<u></u>	C Specify:	Output function
		Mutation	Display to command window
		Mutation function: Use constraint dependent default	• <u>Vectorize</u>
	I	*	More Information

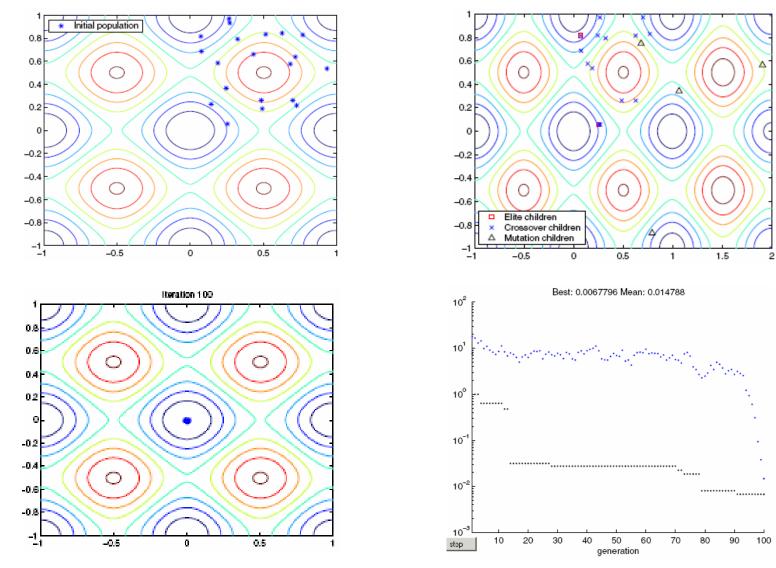
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#### Rastrigin's Function

$$Ras(\mathbf{x}) = 20 + x_1^2 + x_2^2 - 10(\cos 2\pi x_1 + \cos 2\pi x_2)$$



#### **Iteration History**



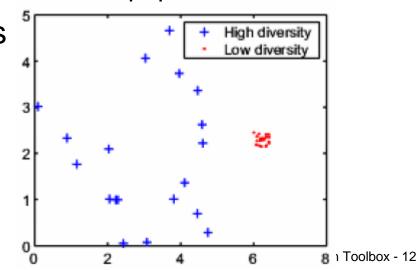
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# GA: Terminology

- Fitness function: objective function
- Individual: genome (genes)
  - any point to which you can apply the fitness function
- Population: an array of individuals
- Generation: Each successive population
- Diversity

- average distance between individuals in a population

- Fitness value and best fitness
- Parent and children



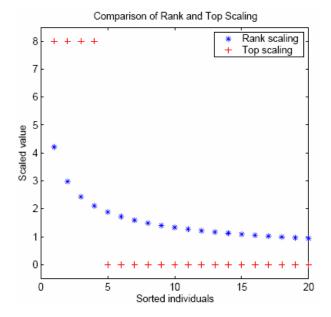
## GA: Options (1)

```
– options = gaoptimset(@ga)
```

```
options =
                                            InitialPopulation: []
       PopulationType: 'doubleVector'
                                                InitialScores: []
          PopInitRange: [2x1 double]
                                               InitialPenalty: 10
       PopulationSize: 20
                                                PenaltyFactor: 100
            EliteCount: 2
                                                 PlotInterval: 1
    CrossoverFraction: 0.8000
                                                  CreationFcn: @gacreationuniform
        ParetoFraction: []
                                            FitnessScalingFcn: @fitscalingrank
    MigrationDirection: 'forward'
                                                 SelectionFcn: @selectionstochunif
    MigrationInterval: 20
                                                 CrossoverFcn: @crossoverscattered
    MigrationFraction: 0.2000
                                                  MutationFcn: {[1x1 function_handle] [1] [1]}
           Generations: 100
                                           DistanceMeasureFcn: []
             TimeLimit: Inf
                                                    HybridFcn: []
          FitnessLimit: - Inf
         StallGenLimit: 50
                                                      Display: 'final'
                                                     PlotFcns: []
        StallTimeLimit: Inf
                                                   OutputFcns: []
                TolFun: 1.0000e-006
                                                   Vectorized: 'off'
                TolCon: 1.0000e-006
                                                  UseParallel: 'never'
```

## GA: Options (2)

- Population Diversity
  - Setting the initial range / population size
- Fitness Scaling
  - Rank, Proportional, Top, Shift linear
- Selection
  - Stochastic uniform, Remainder, Uniform, Roulette, Tournament

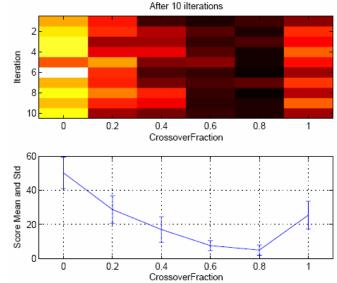


## GA: Options (3)

- Reproduction Options
  - Elite count, Crossover fraction
  - >>deterministicstudy
- Crossover

Population size	20	Children
Elite count	2	2
Crossover fraction	0.8	0.8*18→14
Mutation		4

 Scattered, Single point, Two point, Intermediate, Heuristic, Arithmetic

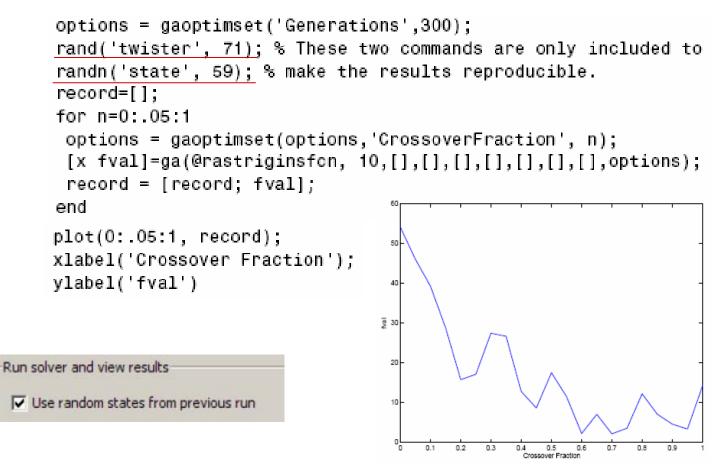


# GA: Options (4)

- Mutation
  - Mutation function: Gaussian, Uniform, Adaptive feasible
  - Scale: controls the standard deviation of the mutation
  - Shrink: controls the rate at which the average amount of mutation decreases
- Global vs. Local minima
  - Increase the initial range
- Using a Hybrid Function
  - optimization function that runs after the genetic algorithm terminates in order to improve the value of the fitness function
  - uses the final point from the genetic algorithm as its initial point
- Setting the Maximum Number of Generations
- Vectorizing the Fitness Function
  - 'Vectorized' On (compute the fitness for all individuals at once)
  - tic; ga(@fitnessfun, nvars); toc

#### **Reproducing Results**

run the genetic algorithm with different settings for
 Crossover fraction to see which one gives the best results



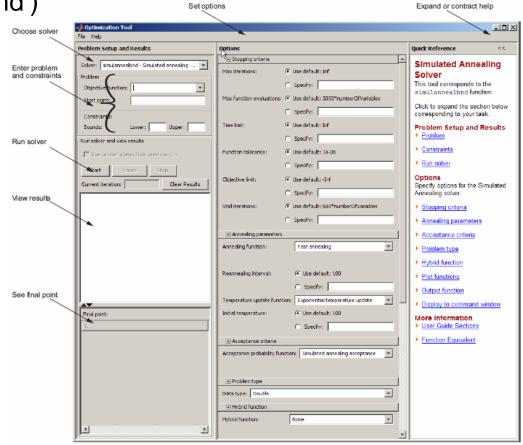
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## Simulated Annealing and Threshold Acceptance

- method for solving unconstrained and bound-constrained optimization problems
- The method models the physical process of heating a material and then slowly lowering the temperature to decrease defects, thus minimizing the system energy
- Threshold acceptance
  - Instead of accepting new points that raise the objective with a certain probability, it accepts all new points below a fixed threshold (faster than SA)
  - The threshold is then systematically lowered, just as the temperature is lowered in an annealing schedule

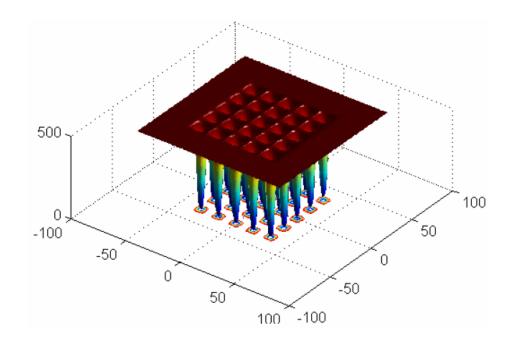
## Function Call

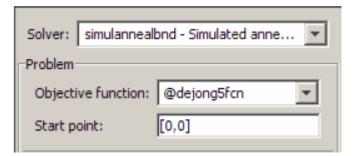
- [x fval exitflag output] = simulannealbnd(@objfun, x0, lb, ub, options)
- [x, fval exitflag output] = threshacceptbnd(@objfun, x0, lb, ub, options)
- optimtool('simulannealbnd')



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#### De Jong's Fifth Function





fun = @dejong5fcn;
[x fval] = simulannealbnd(fun, [0 0])

x =
 -31.9779 -31.9595
fval =
 0.9980

# Terminology

- Objective function
- **Temperature:** InitialTemperature, TemperatureFcn
  - It determines the probability of accepting a worse solution at any step and is used to limit the extent of the search in a given dimension
- Annealing schedule: TemperatureFcn
  - The rate by which the temperature is decreased
  - The slower the rate of decrease, the better the chances are of finding an optimal solution, but the longer the run time
- **Reannealing:** ReannealInterval
  - Raises the temperature after a certain number of new points have been accepted, and starts the search again at the higher temperature
- Threshold acceptance
  - accepts a worse point if the objective function is raised by less than a fixed threshold

## SA: Options

- options=saoptimset('simulannealbnd')
- options=saoptimset('threshacceptbnd')

```
options =
          AnnealingFcn: @annealingfast
        TemperatureFcn: @temperatureexp
        AcceptanceFcn: @acceptancesa
                TolFun: 1.0000e-006
        StallIterLimit: '500*numberofvariables'
           MaxFunEvals: '3000*numberofvariables'
             TimeLimit: Inf
               MaxIter: Inf
       ObjectiveLimit: -Inf
               Display: 'final'
      DisplayInterval: 10
             HybridFcn: []
        HybridInterval: 'end'
              PlotFcns: []
          PlotInterval: 1
            OutputFons: []
    InitialTemperature: 100
     ReannealInterval: 100
              DataType: 'double'
```

#### **Nonlinear Constraint Solver**

- Augmented Lagrangian Pattern Search (ALPS) algorithm
- Augmented Lagrangian Genetic Algorithm (ALGA)

$$\begin{array}{l} \text{Minimize} \\ x \\ f(x) \\ \text{such that} \\ C_i(x) \leq 0, i = 1 \dots m \\ C_i(x) = 0, i = m + 1 \dots mt \\ Ax \leq b \\ A_{eq}x = beq \\ LB \leq x \leq UB \end{array}$$

$$\Theta(x,\lambda,s,\rho) = f(x) - \sum_{i=1}^m \lambda_i s_i \log(s_i - c_i(x)) + \sum_{i=m+1}^{mt} \lambda_i c_i(x) + \frac{\rho}{2} \sum_{i=m+1}^{mt} c_i(x)^2 + \frac{\rho}{2} \sum_{i=m+1}^{mt}$$

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