



Inspire...Educate...Transform.

## **Simulations, Evolutionary searches**

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Why do we need samples

# **SAMPLING STRATEGY**



# Learn to sample

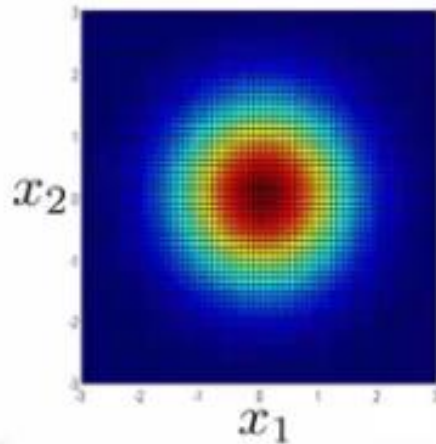
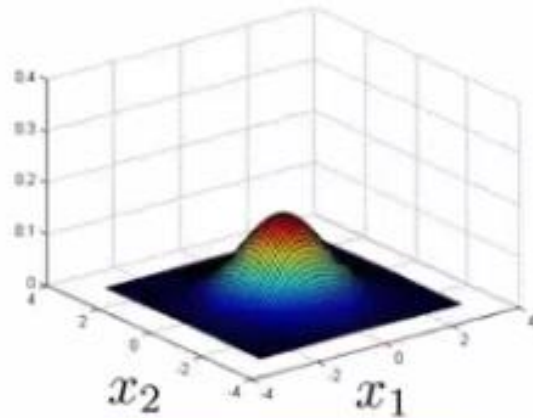
- Given a distribution, let us learn to sample from it
- First, let us understand why and then learn the strategy
- Then check how good this process is

# Measuring central tendencies

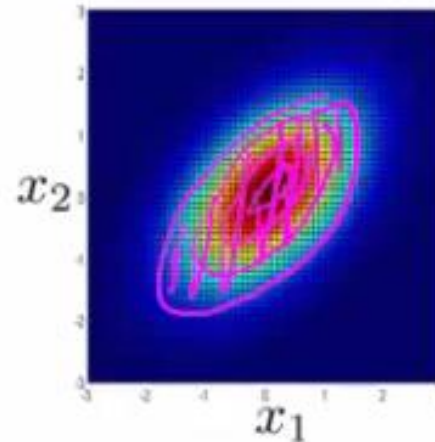
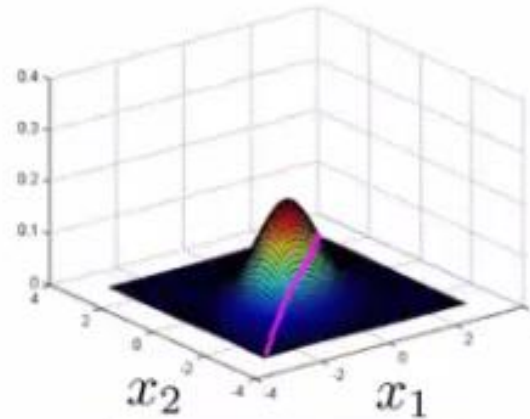
- Computing mean median and mode is a lot of calculus for a continuous distribution
  - $\Sigma$  must be replaced by  $\int$  sign.
  - $Mean = \int xp(x)dx$
  - $Median = \int_0^M p(x)dx = \int_M^k p(x)dx = \frac{1}{2}$
  - $Mode$  is where  $\frac{d(p(x))}{dx} = 0$

# Even normal distribution becomes complex

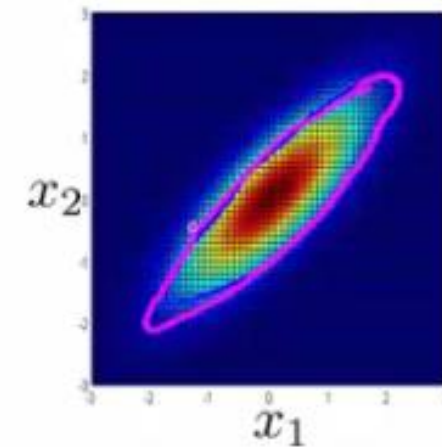
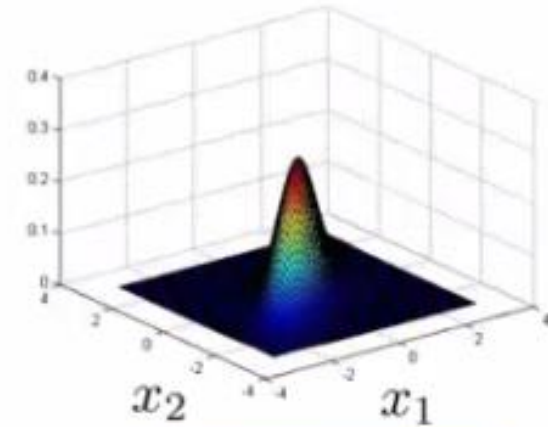
$$\mu = \begin{bmatrix} 0 \\ 0 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$$



$$\mu = \begin{bmatrix} 0 \\ 0 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1 & 0.5 \\ 0.5 & 1 \end{bmatrix}$$



$$\mu = \begin{bmatrix} 0 \\ 0 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1 & 0.8 \\ 0.8 & 1 \end{bmatrix}$$



# Multivariate Gaussian

*Bell curve in multivariate case =*

$$2\pi^{\frac{-N}{2}} |\Sigma|^{\frac{-1}{2}} e^{\left(\frac{-1}{2}(X-\mu)^T \Sigma^{-1} (X-\mu)\right)}$$

$$\mu = \frac{1}{m} \sum_{i=1}^m x^{(i)}$$

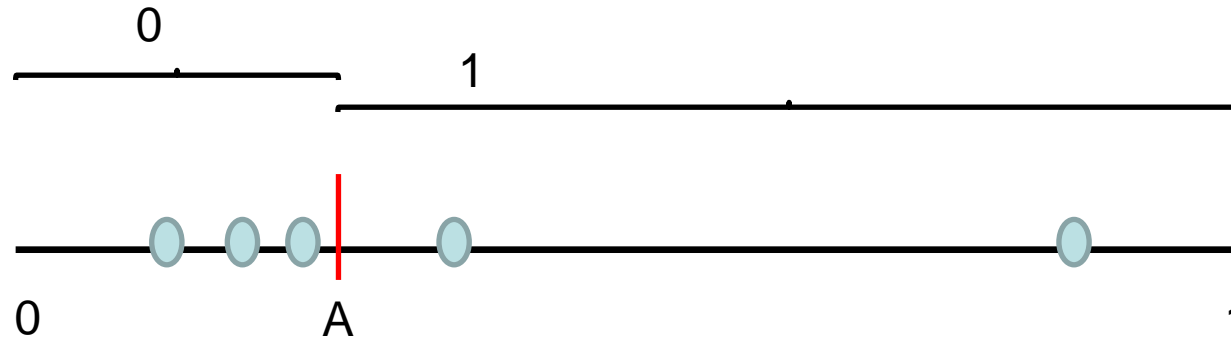
$$\Sigma = \frac{1}{m} \sum_{i=1}^m (x^{(i)} - \mu)(x^{(i)} - \mu)^T$$

# Estimating from samples

- Let us have IID (identically, independently distributed samples) of a distribution
- Then, mean is

$$\frac{1}{M} \sum_{m=1}^M x[m]$$

# Creating samples from a discrete distribution



$$P(0) = A \text{ and } P(1) = 1 - A$$

Randomly generate a number between 0 and 1; If it is less than  $A$ , sample is 0 else it is 1

Sample = (0,1,0,0,1)



# Discretizing a continuous variable

- Equal bin, equal frequency
- With class variable, sort and decide where to cut (cut wherever the class variable changes its values)
- Many such algorithms exist

Age	25	32	34	35	35	37	37	38
Loan	0	1	1	0	0	0	1	1

# Sampling based estimations

$$T_D = \frac{1}{M} \sum_{m=1}^M X[m]$$

Hoeffding Bound:

$$P_D(T_D \notin \underbrace{[p - \epsilon, p + \epsilon]}) \leq 2e^{-2M\epsilon^2}$$

The probability of getting the sample estimate outside the range of  $\mathcal{E}$  is given by right hand side. So, this is the probability of a bad estimate.

It goes down exponentially with sample size and tolerance.

For additive bound  $\epsilon$  on error with probability  $> 1-\delta$ :

$$M \geq \frac{\ln(2/\delta)}{2\epsilon^2}$$

It turns out that we do need a large number of samples in most cases.

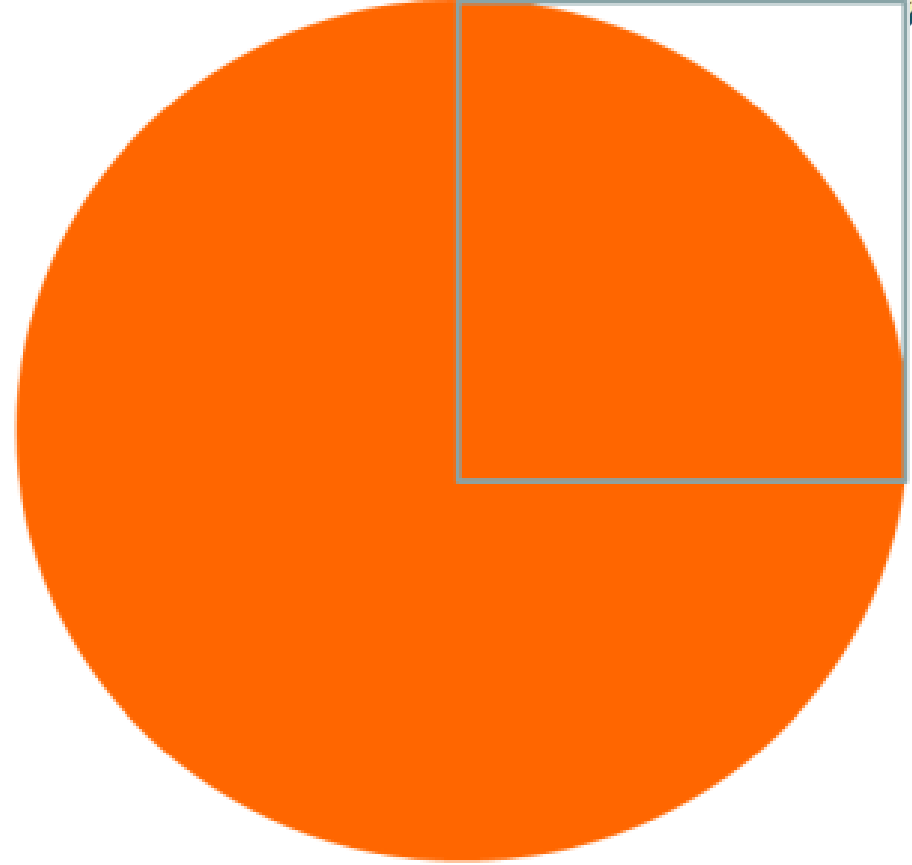


Computing central tendencies

## **APPLICATIONS OF SAMPLING:**

# Monte Carlo Simulations

Value of Pi

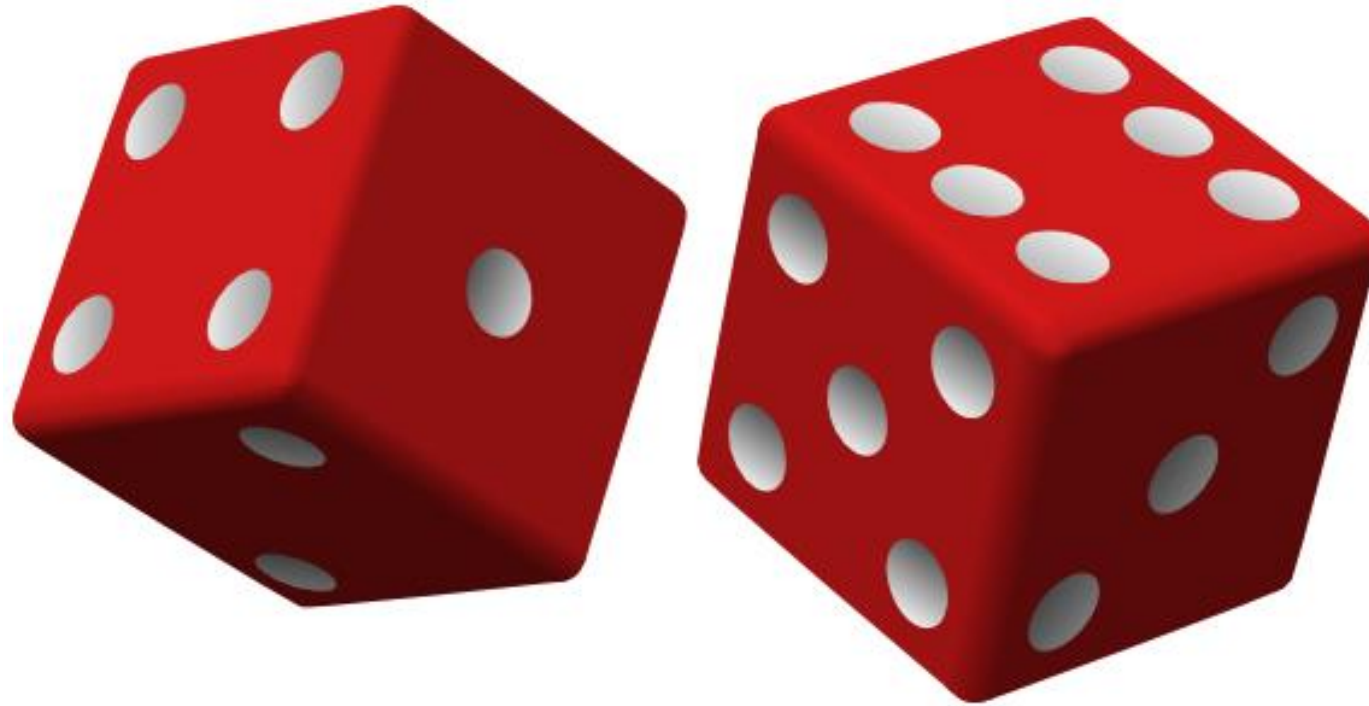




# When to stop?

- Define a small tolerance  $\delta$  delta
- If the difference between  $n^{\text{th}}$  iteration and  $n+1^{\text{th}}$  iteration is less than delta, stop.  
Else, continue

# Probability



# Capital markets



Outcome	Green	Red	White
1	0.8	0.06	0.9
2	0.9	0.2	1
3	1.05	1	1
4	1.1	3	1
5	1.2	3	1
6	1.4	3	1.1





# **GENERATING AND UNDERSTANDING BEHAVIOR OF TEXT, IMAGES, AUDIO**



# Transition probabilities

- Process
  - Learn the distribution from a large corpus
  - Collate counted probabilities as a matrix (columns, current state and rows future state)
  - Text, images, music etc.

# Learning distributions



	a	b	c	d	e	f	g	h	i
A	0.08	0.09							
B									

Larger corporuses yield better descriptions

# Character generation

- Random
  - No use
- First order (character-Character transitions)
  - t I amy, vin. id wht omanly heay atuss n macon  
aresethe hired boutwhe t, tl, ad torurest t plur I wit  
hengamind tarer-plarody thishand.
- Second order (Previous 2 characters, di-gram)
  - Ther I the heingoind of-pleat, blur it dwere wing  
waske hat trooss. Yout lar on wassing, an sit."  
"Yould," "I that vide was nots ther.

# Applications of Markov chains



- 3<sup>rd</sup> order
  - I has them the saw the secorrow. And wintails on my my ent, thinks, fore voyager lanated the been elsed helder was of him a very free bottlemarkable,
- 4 gram
  - His heard." "Exactly he very glad trouble, and by Hopkins! That it on of the who difficentralia. He rushed likely?" "Blood night that.

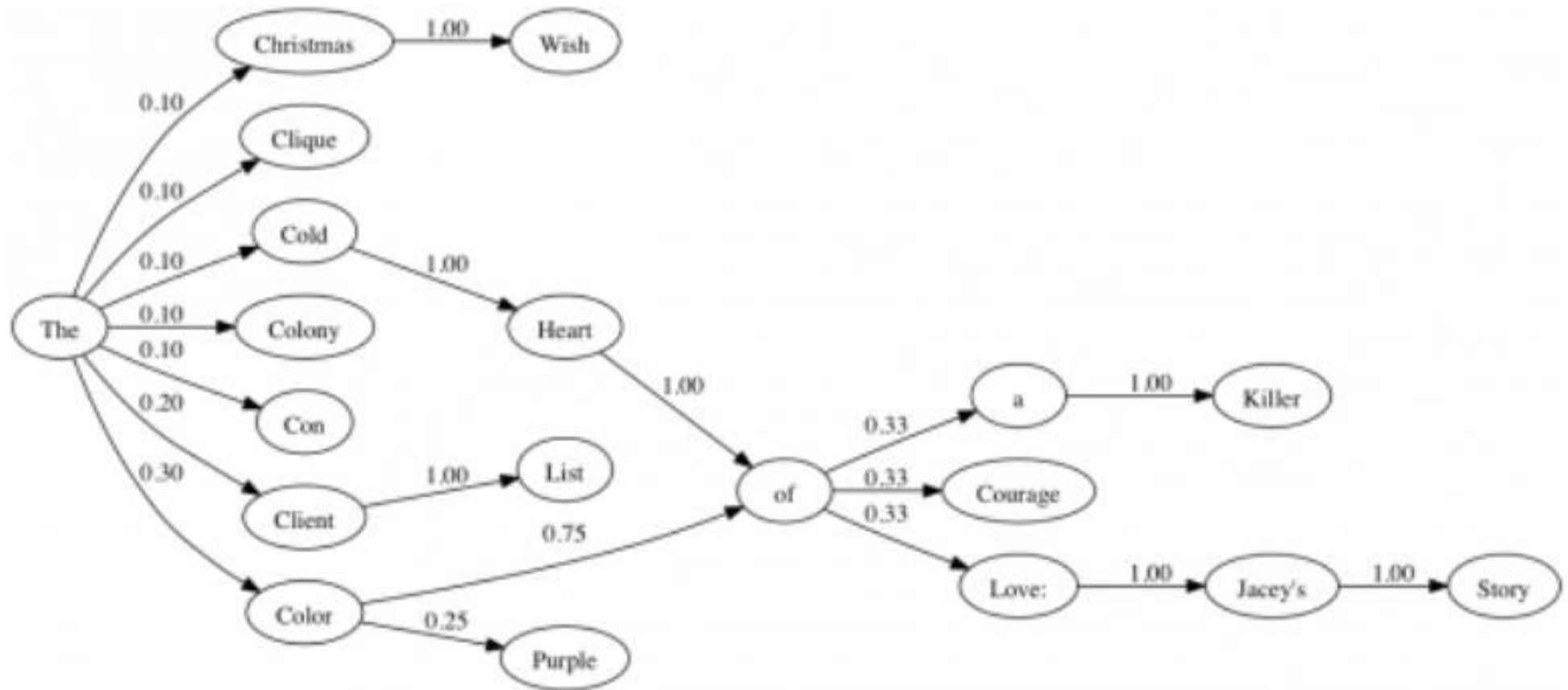
# Text generation

- First order word markov chains
  - Construct a matrix with transition probabilities of all word pairs and the probability of succeeding
  - Start random and select the words obeying the distribution



# Let us generate new movie titles

- <http://www.soliantconsulting.com/blog/2013/02/title-generator-using-markov-chains>
  - The Christmas Wish, The Client, The Client List, The Clique, The Cold Heart of a Killer, The Colony, The Color of Courage, The Color of Love: Jacey's Story, The Color Purple, The Con
  - Constructing a list of unique words that appear in these titles: "The", "Christmas", "Wish", "Client", "List", "Clique", "Cold", "Heart", "of", "a", "Killer", "Colony", "Color", "Courage", "Love:", "Jacey's", "Story", "Purple", "Con".



Probability diagram



# Process



- We can start with the word “The” and for each new word desired, we’ll choose from the distribution which next node to move to accordingly.
- Some possible generated strings include new titles “The Color of a Killer” or “The Cold Heart of Love: Jacey’s Story”.
- With more data, a more complicated graph can be constructed, and more interesting and varied strings can be generated as a result. You should use as much data in your corpus as you can find!
- Small tweaks like setting a minimum length, choosing to start with start words and end with end words might help



# Markov Chain in Text

- [http://en.wikipedia.org/wiki/Mark\\_V\\_Shaney](http://en.wikipedia.org/wiki/Mark_V_Shaney)
- Finding who the author of a work is
  - Compare the transitional probabilities of unknown work and known works

# Music: Monte Carlo



rand\_music.wav

# First order transitions



## 1st-order matrix

Note	A	C#	E♭
A	0.1	0.6	0.3
C#	0.25	0.05	0.7
E♭	0.7	0.3	0

# Markov second order

2nd-order matrix

Note	A	D	G
AA	0.18	0.6	0.22
AD	0.5	0.5	0
AG	0.15	0.75	0.1
DD	0	0	1
DA	0.25	0	0.75
DG	0.9	0.1	0
GG	0.4	0.4	0.2
GA	0.5	0.25	0.25
GD	1	0	0

# Music\_Markov Chain



first\_score.wav

<https://ssodelta.wordpress.com/2014/03/20/generating-mozart-from-markov-chains/>

<https://medium.com/@omgimanerd/generating-music-using-markov-chains-40c3f3f46405#.2ydm6na5>

<http://www.cs.uml.edu/ecg/uploads/Alfall11/SimoneHill.FinalPaper.MarkovMelodyGenerator.pdf>



- HMM models
  - Learn transition probabilities of parts of speech using 2<sup>nd</sup> order
  - Gene sequences



# **AN INTERESTING PROPERTY OF TRANSITION PROBABILITY MATRICES**



# Understanding Markov Chain

- Let us say, my class has the following properties
  - 90% of the interested students will remain interested; 10% doze off
  - 50% of the dozing students get interested; remaining stay as it is



# What is the steady state?

- Let us start with 100 interested and 0 boring
- Work out what happens after 5 periods

# Using matrices to solve the problem



- Transition matrix
  - Each columns: one state transition
- Transition matrix \* current distribution =  
Next distribution

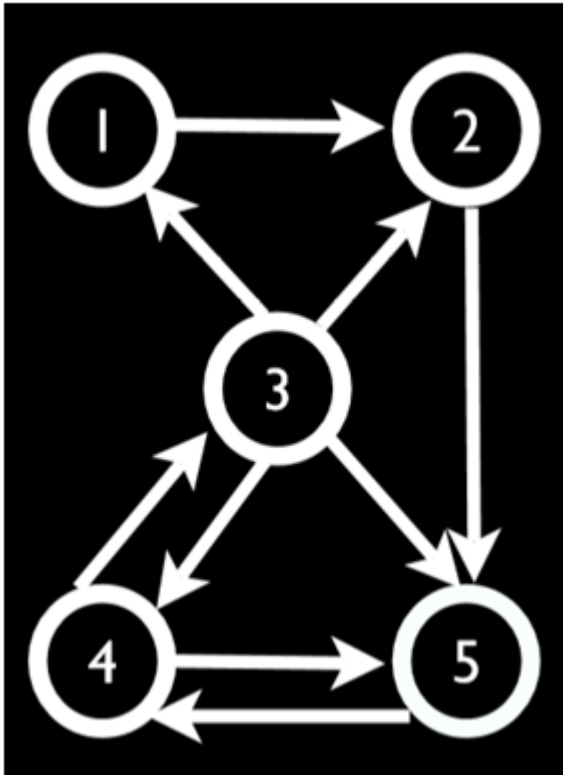


# This is a Markov chain

- Finite states
- Fixed transition probabilities
- From one state, I should be able to go to any state
- Column stochastic matrices

# Other applications

- Google Page rank



$$\begin{bmatrix} 0 & 0 & 1/4 & 0 & 0 \\ 1 & 0 & 1/4 & 0 & 0 \\ 0 & 0 & 0 & 1/2 & 0 \\ 0 & 0 & 1/4 & 0 & 1 \\ 0 & 1 & 1/4 & 1/2 & 0 \end{bmatrix}$$

$$\begin{bmatrix} 1/20 \\ 5/20 \\ 1/10 \\ 5/20 \\ 7/20 \end{bmatrix} = \begin{bmatrix} 0 & 0 & 1/4 & 0 & 0 \\ 1 & 0 & 1/4 & 0 & 0 \\ 0 & 0 & 0 & 1/2 & 0 \\ 0 & 0 & 1/4 & 0 & 1 \\ 0 & 1 & 1/4 & 1/2 & 0 \end{bmatrix} \begin{bmatrix} 1/5 \\ 1/5 \\ 1/5 \\ 1/5 \\ 1/5 \end{bmatrix}$$

$$r^{(t+1)} = H r^{(t)}$$

	Iteration 0	Iteration 1	Iteration 2	Page Rank
$P_1$	1/5	1/20	1/40	5
$P_2$	1/5	5/20	3/40	4
$P_3$	1/5	1/10	5/40	3
$P_4$	1/5	5/20	15/40	2
$P_5$	1/5	7/20	16/40	1



Guided random searches

# **CAN SAMPLING BE USED TO SEARCH FOR MAXIMA AND MINIMA?**

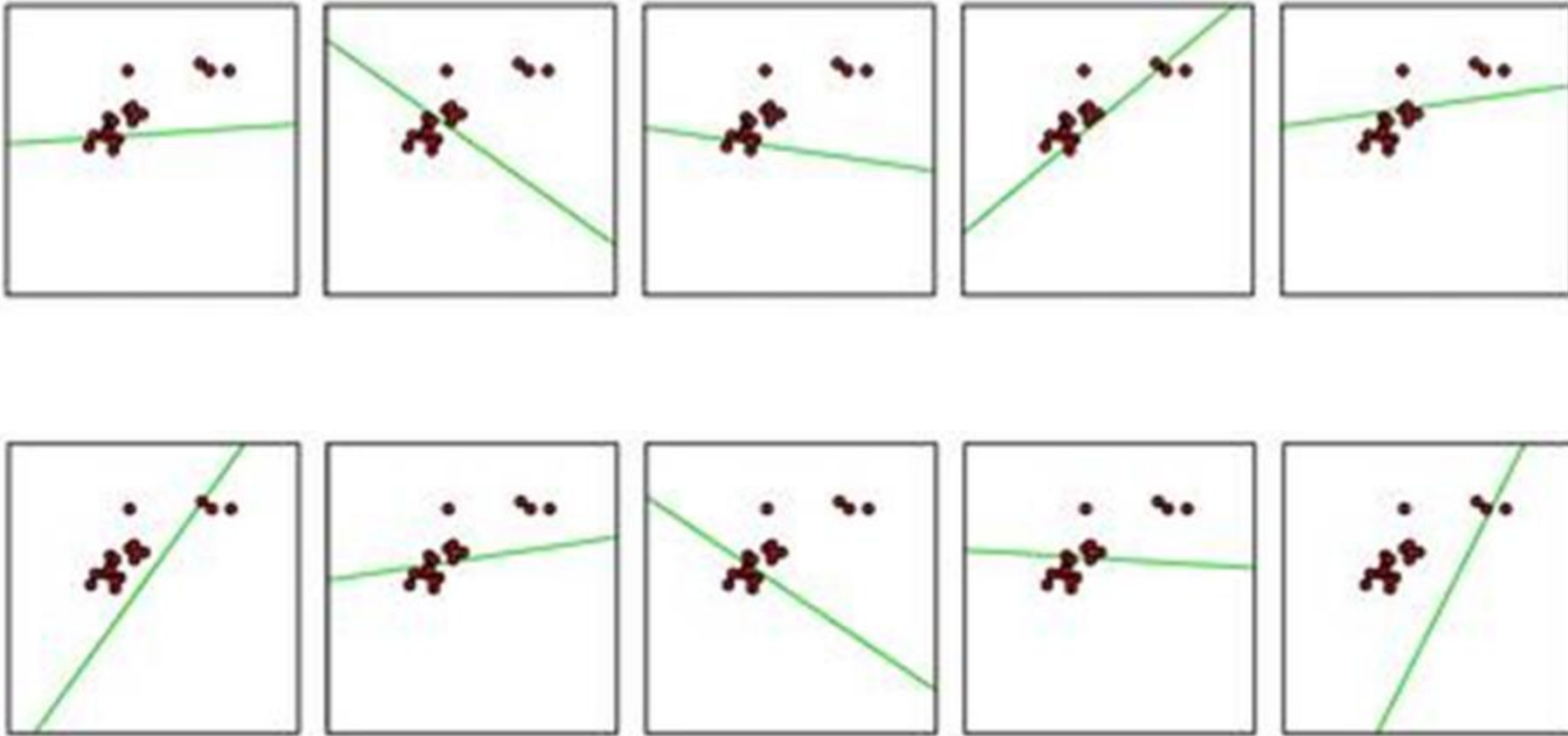




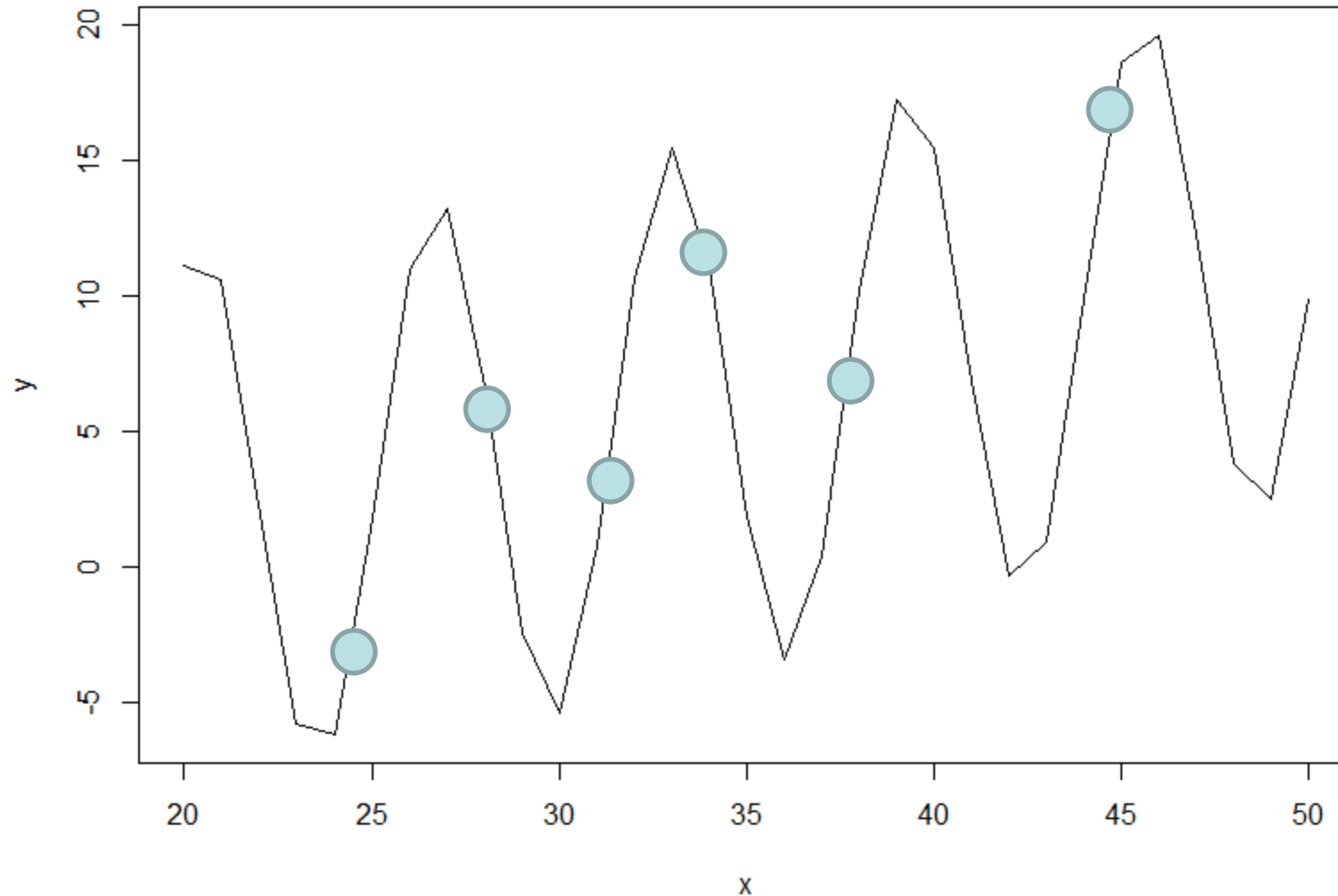
# What Sampling and MonteCarlo are not

- They are random searches
- But, not optimization methods
- Can you do linear regression with Montecarlo?

# Linear regression using Monte Carlo



# A complex function and we need to find minima





# A possible approach

- Do a million simulations and pick the most optimal solution
- Not guaranteed that we get anywhere close to the solution
- What to do?
  - Have memory



Genetic algorithms

# **GUIDED RANDOM SEARCHES**

# What does survival of the fittest mean



# Summary of biology



- Not all members survive until reproduction. Only a few learn to earn lunch. Others are lunch!
- The strong ones then need to attract and mate other members of the species that survived.
- The offspring generation hence is normally better than the parents generation
- Over generations, species develop

# Can a similar theory be applied to optimization



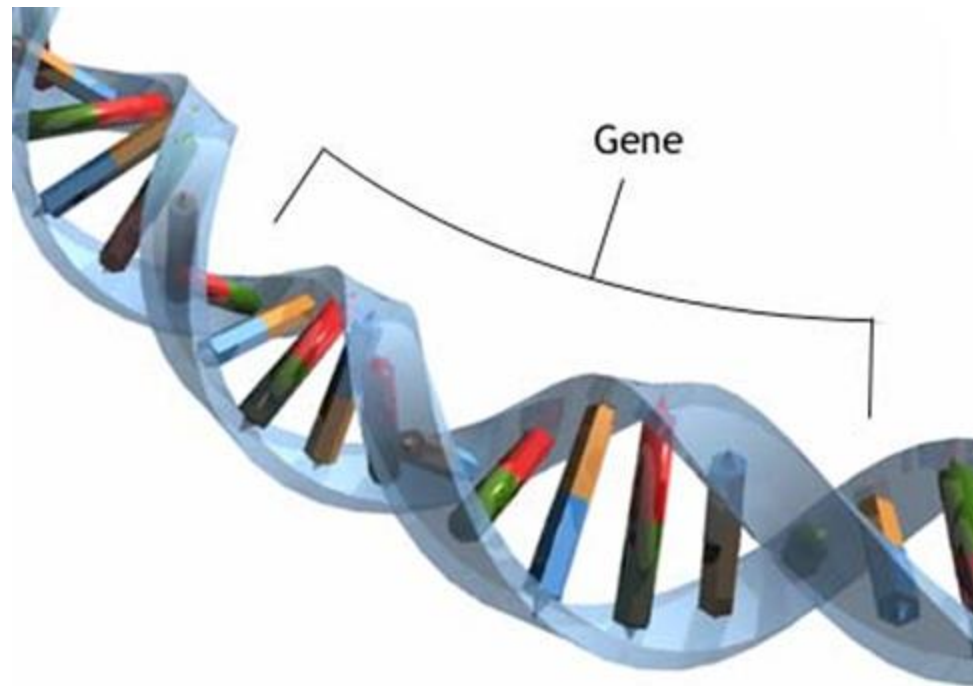
- Start with random solutions
- Kill the weak ones (memory!)
- Reproduce from the good ones
- Will the newer set be more optimal than its parents?



# Gene



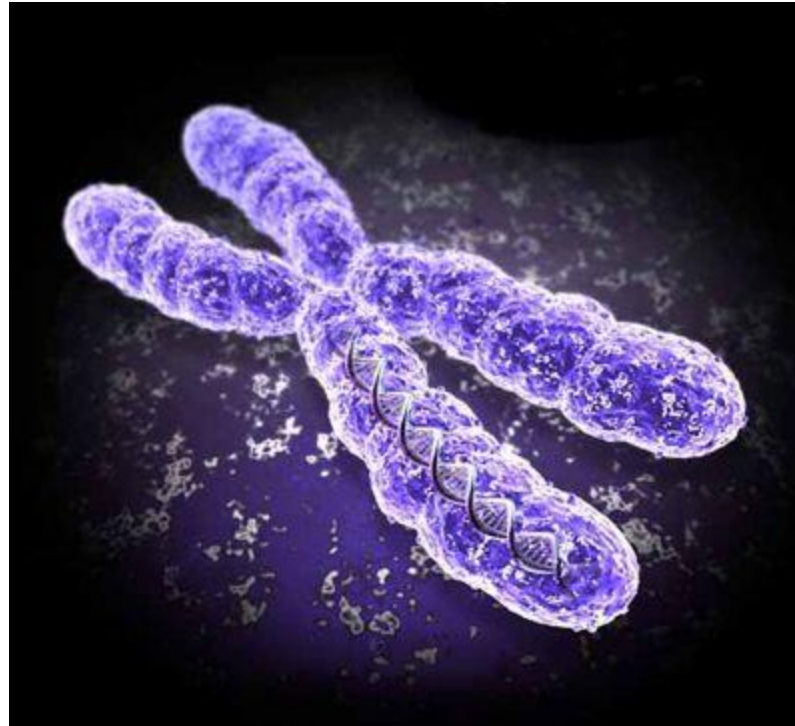
Gene: The reproducible building block of chromosome



# Beg, Borrow or Steel



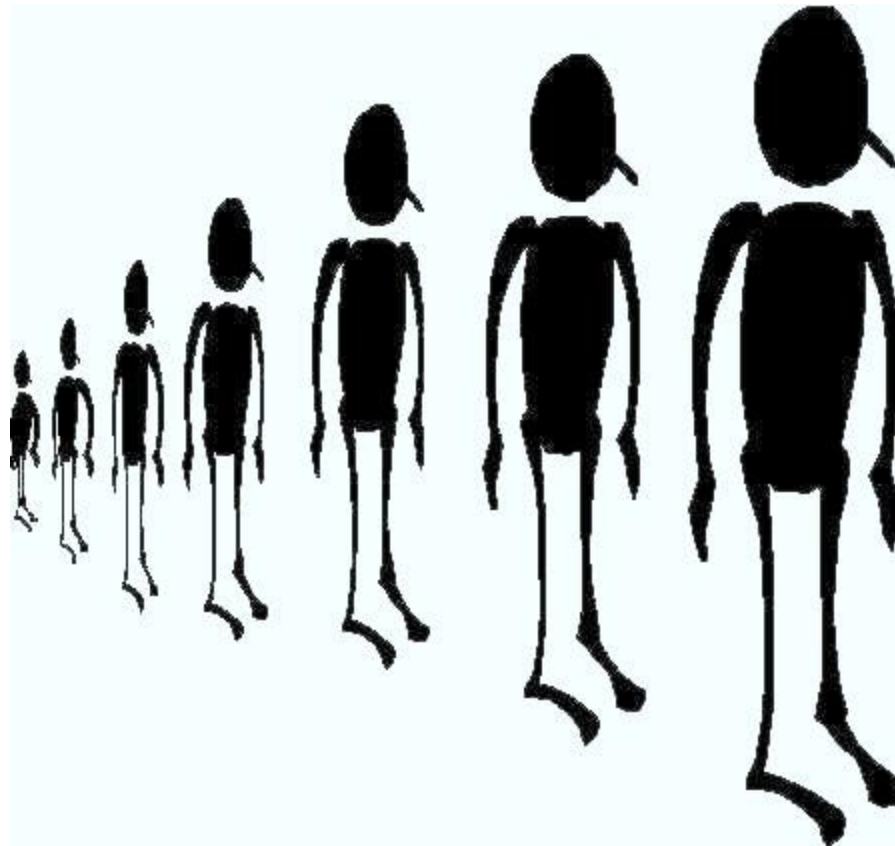
- Chromosome: A possible solution



# Population



- A set of chromosomes



# Generation

- A population derived from the fittest chromosomes of the previous population



# Reproduction, Mating and Mutation



- Create a new solution from the old fitter solutions



```
{  
    initialize population;  
    evaluate population;  
    while TerminationCriteriaNotSatisfied  
    {  
        select parents for reproduction;  
        perform recombination and mutation;  
        evaluate population;  
    }  
}
```

# So, GA at a snapshot



- Create a schema to create possible solutions in a bit/byte format
- Start with a number of random solutions
- Identify the best ones
- Create new ones from them using some exchange mechanism
- Continue with the last two steps until a solution is found



# Some unique differentiators

- GAs work with a coding of the parameters and not the parameters themselves
- GAs search from a population of points and not a single point
- GAs work with the objective function and not the derivatives
- GAs transition based on random rules and not on deterministic rules





# OPERATIONAL ASPECTS



# Innovations: Creating a coding scheme

- Knap Sack
- Polynomial fit
- TSP
- Portfolio allocation
  
- Define objective function and solution

# Population



Chromosomes could be:

- Bit strings
- Real numbers
- Permutations of element
- Lists of rules
- Program elements
- ... any data structure ...

(0101 ... 1100)

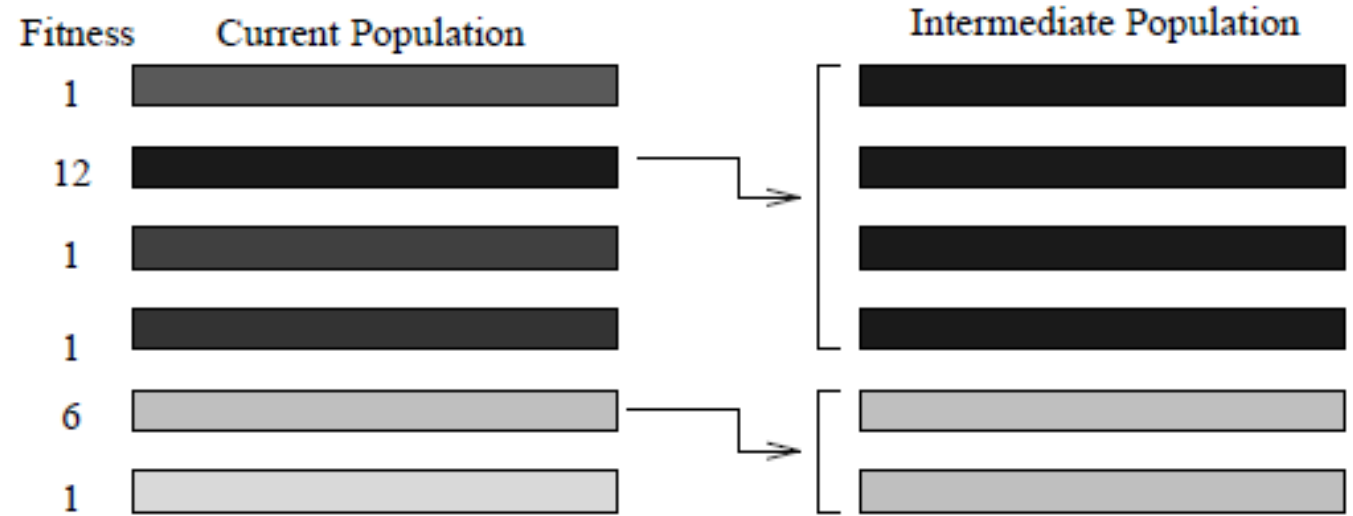
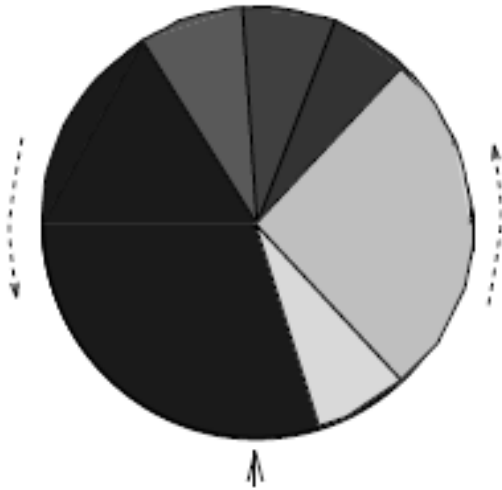
(43.2 -33.1 ... 0.0 89.2)

(E11 E3 E7 ... E1 E15)

(R1 R2 R3 ... R22 R23)

(genetic programming)

# How do we decide mating pool



*Figure 2. Selection application*



# Mutation or mating?

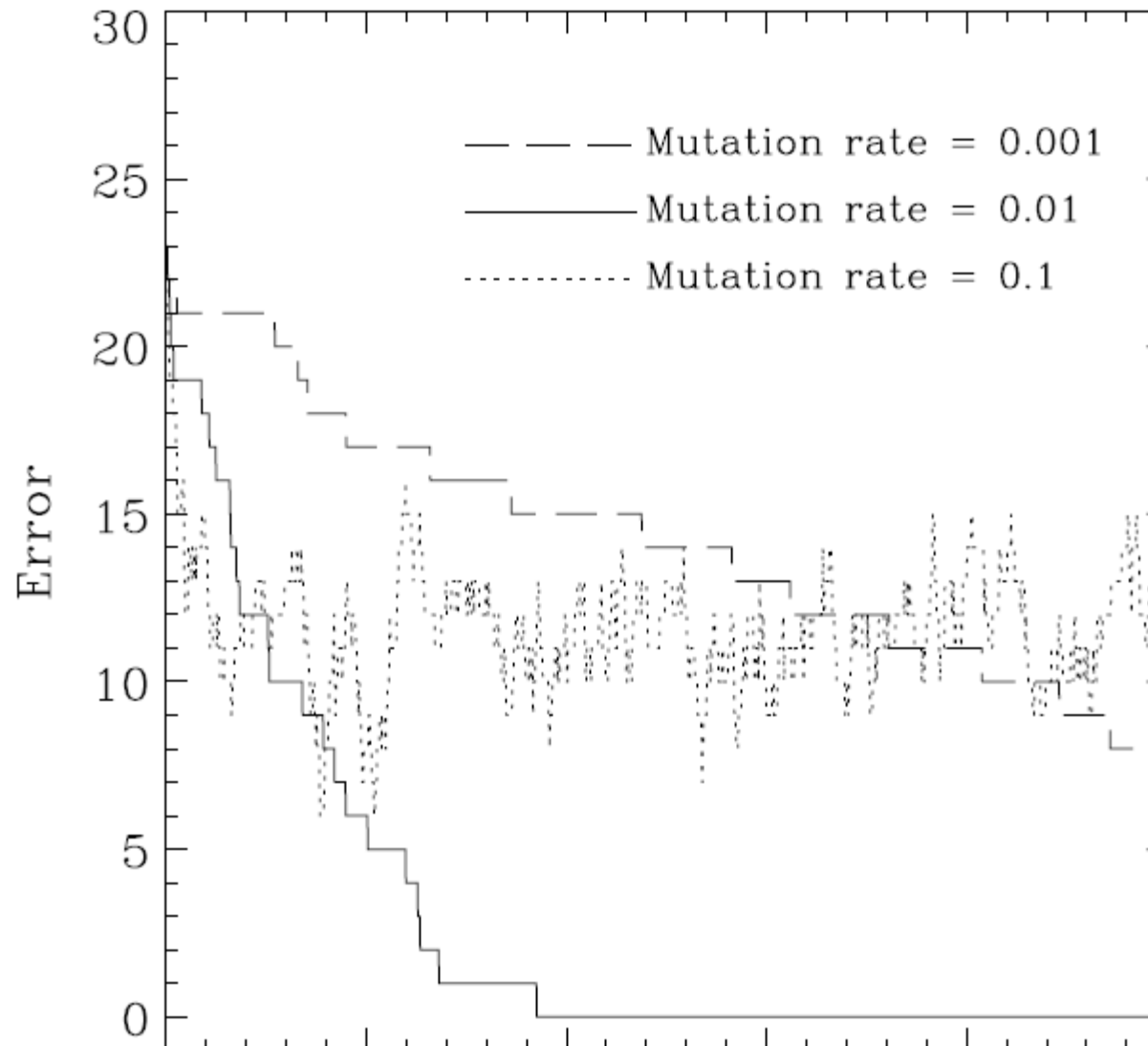
- A low probability of mutation
- Come down with generation



# Non uniform mutation

- Initially mutate heavily
- As the number of generations increase mutate less

# What about mutation?



# How do we determine reproduction

- Most common: Cross over
  - npoint crossover:  $n$  crossover points are randomly selected and the segments of the parents, defined by them, are exchanged for generating the offspring.
  - uniform crossover: the values of each gene in the offspring are determined by the uniform random choice of the values of this gene in the parents.



# For real coding

## Arithmetical crossover (Michalewicz, 1992)

Two offspring,  $H_k = (h_1^k, \dots, h_i^k, \dots, h_n^k)$   $k = 1, 2$ , are generated, where  $h_i^1 = \lambda c_i^1 + (1 - \lambda)c_i^2$  and  $h_i^2 = \lambda c_i^2 + (1 - \lambda)c_i^1$ .  $\lambda$  is a constant (uniform arithmetical crossover) or varies with regard to the number of generations made (non-uniform arithmetical crossover).

- Flat crossover: An offspring,  $H$  is generated, where  $h_i$  is a randomly (uniformly) chosen value of the interval  $[c1_i; c2_i]$

# Cross over



- Hello World
- Knap Sack
- Polynomial fit
- TSP
- Portfolio allocation



# Inversion or Mutation

- Flip
- Random generation
- An unexpected operation



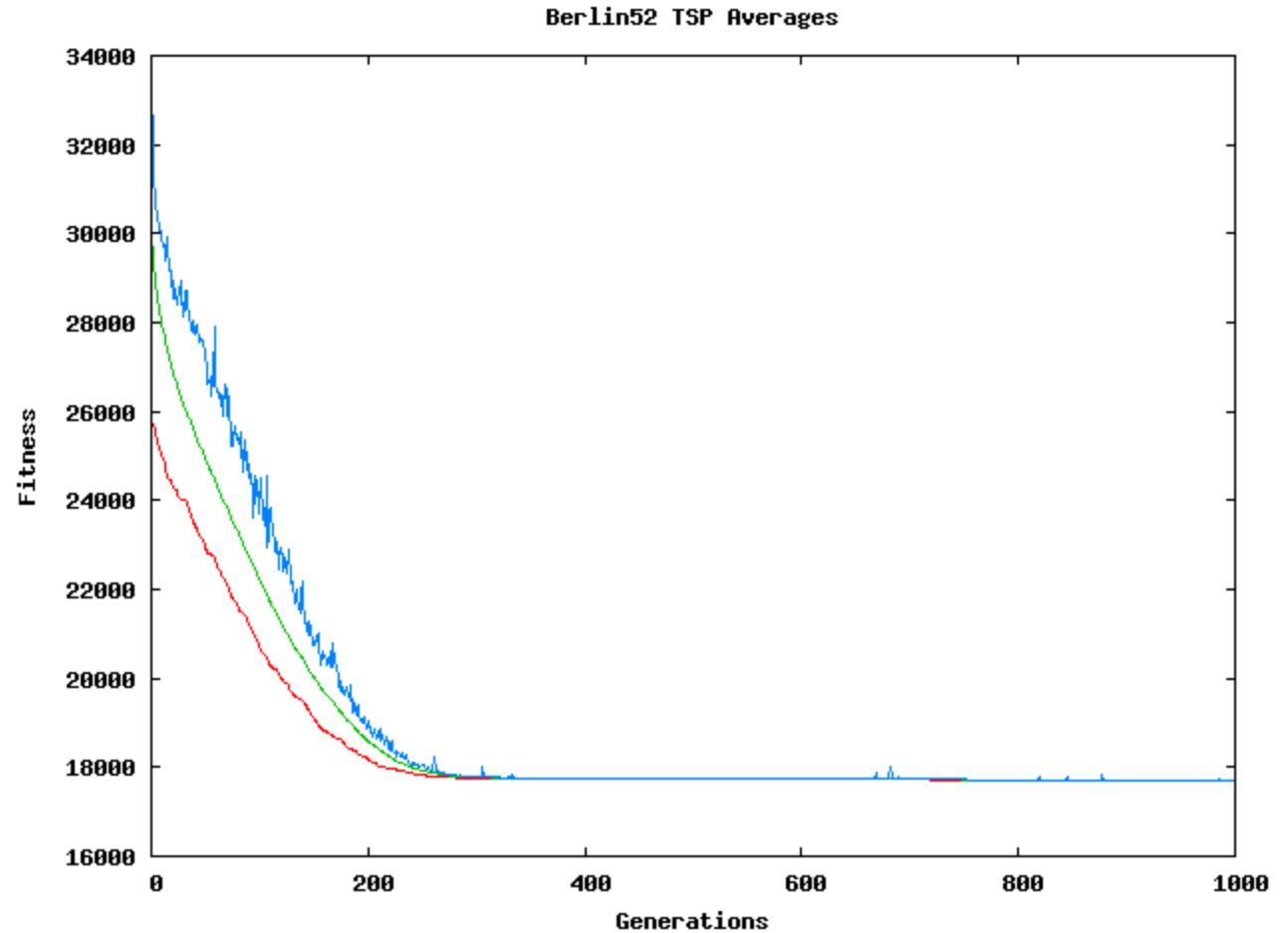
# Mutation scheme

- Hello World
- Knap Sack
- Polynomial fit
- TSP
- Portfolio allocation

# Convergence of a GA



When to stop?





# EXAMPLES

# A Simple Example



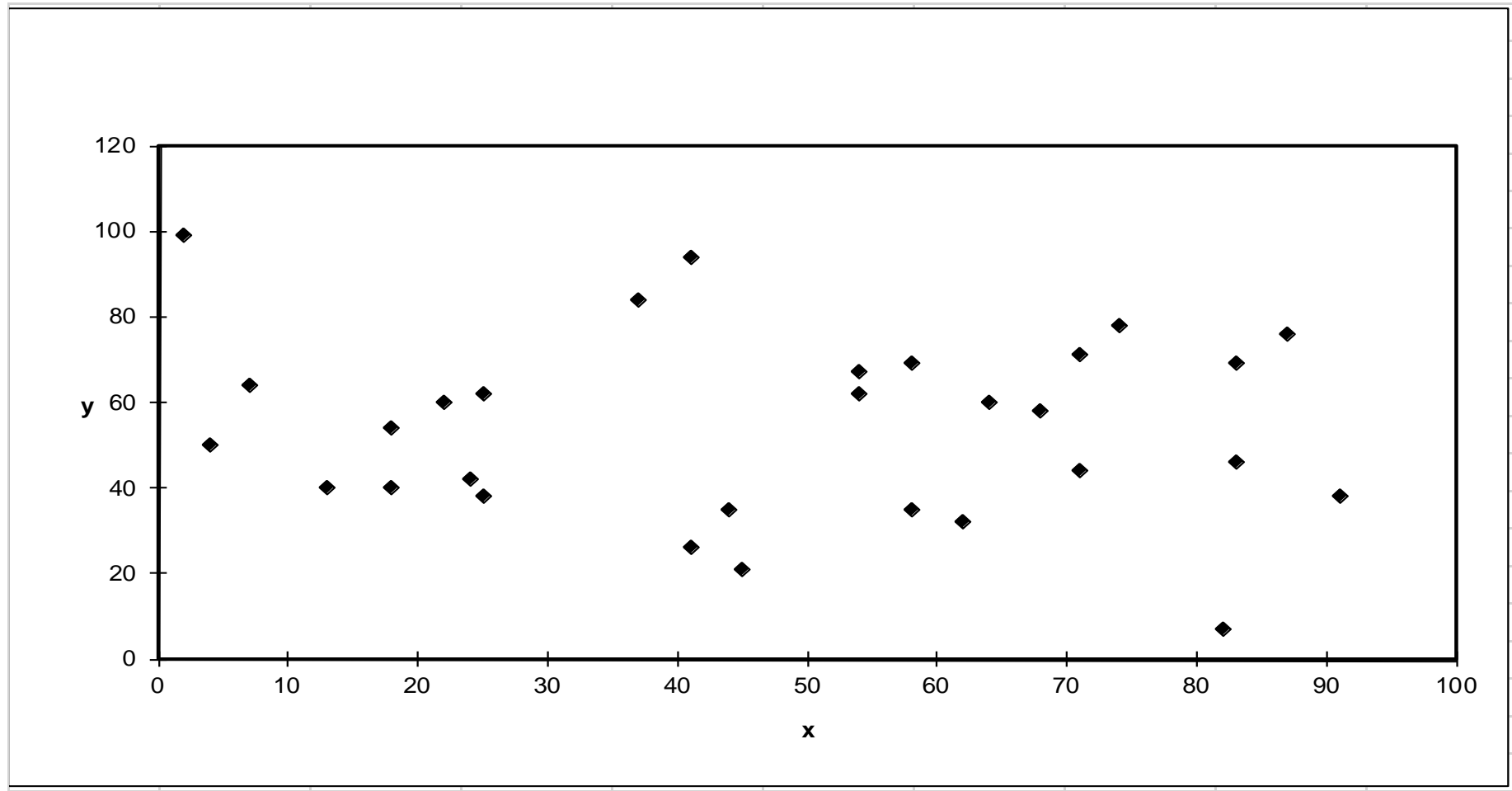
The Traveling Salesman Problem:

Find a tour of a given set of cities so that

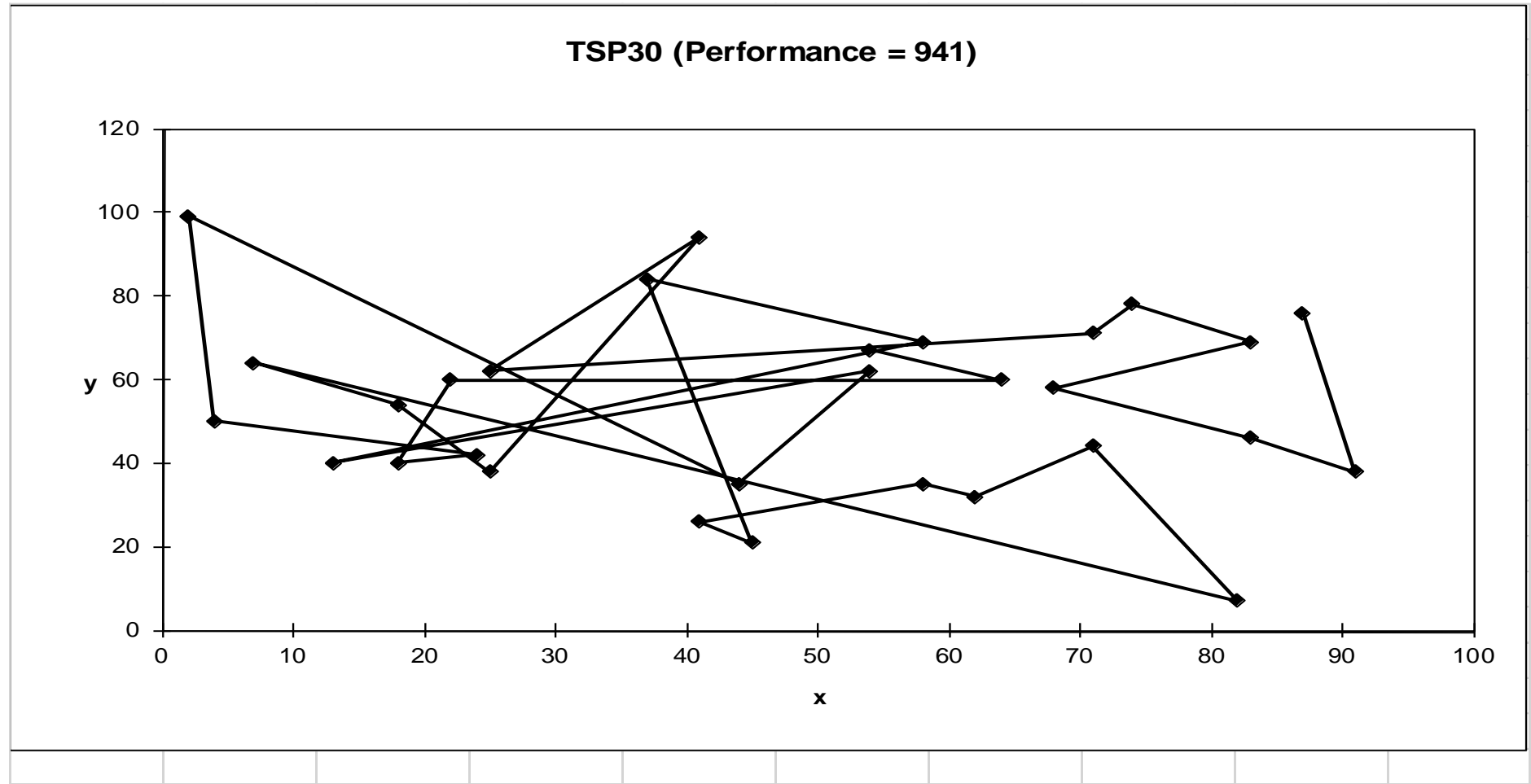
- each city is visited only once
- the total distance traveled is minimized



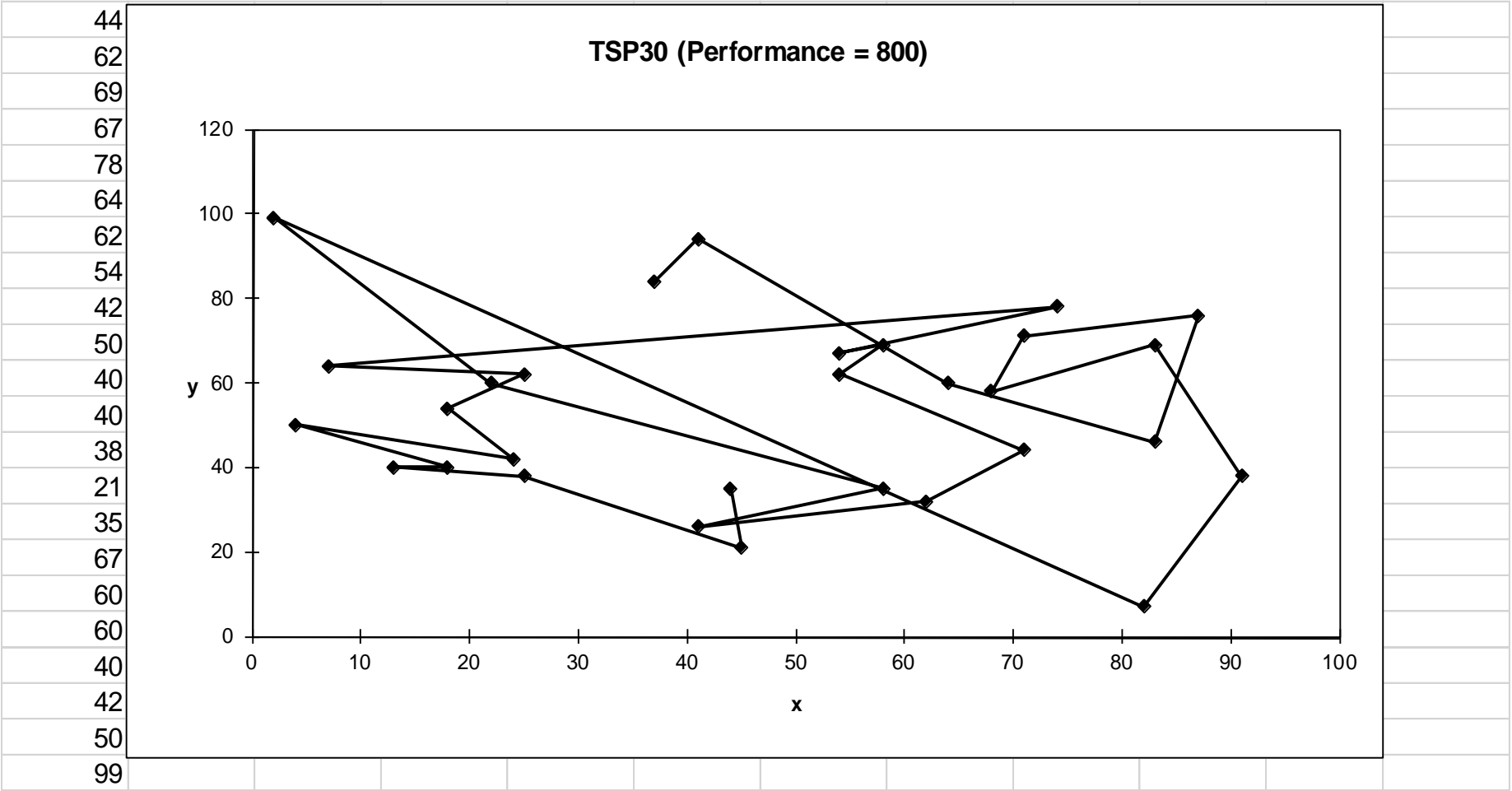
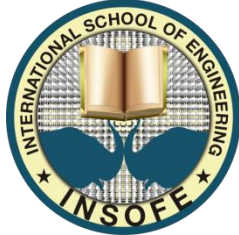
# TSP Example: 30 Cities



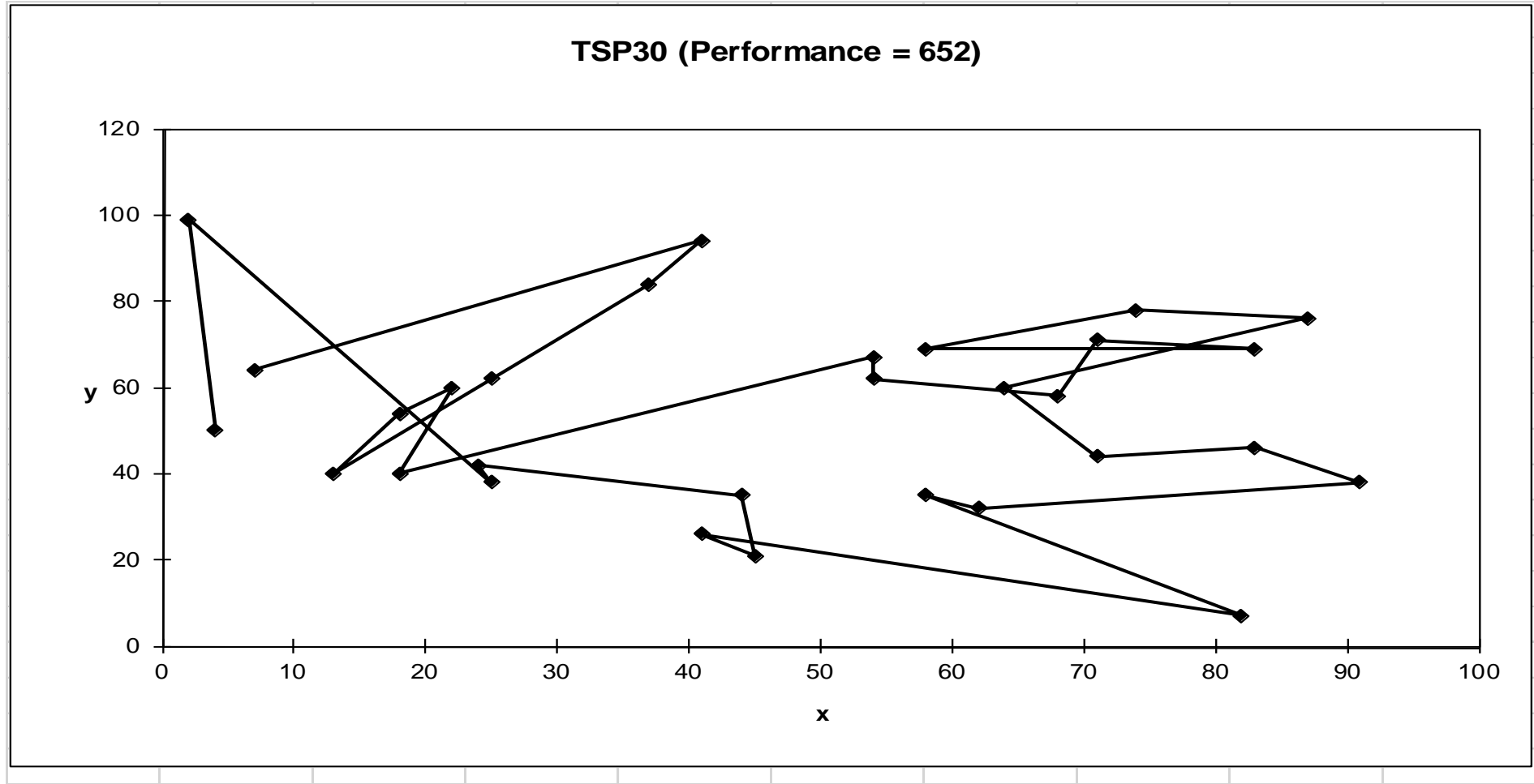
# Solution <sub>i</sub> (Distance = 941)



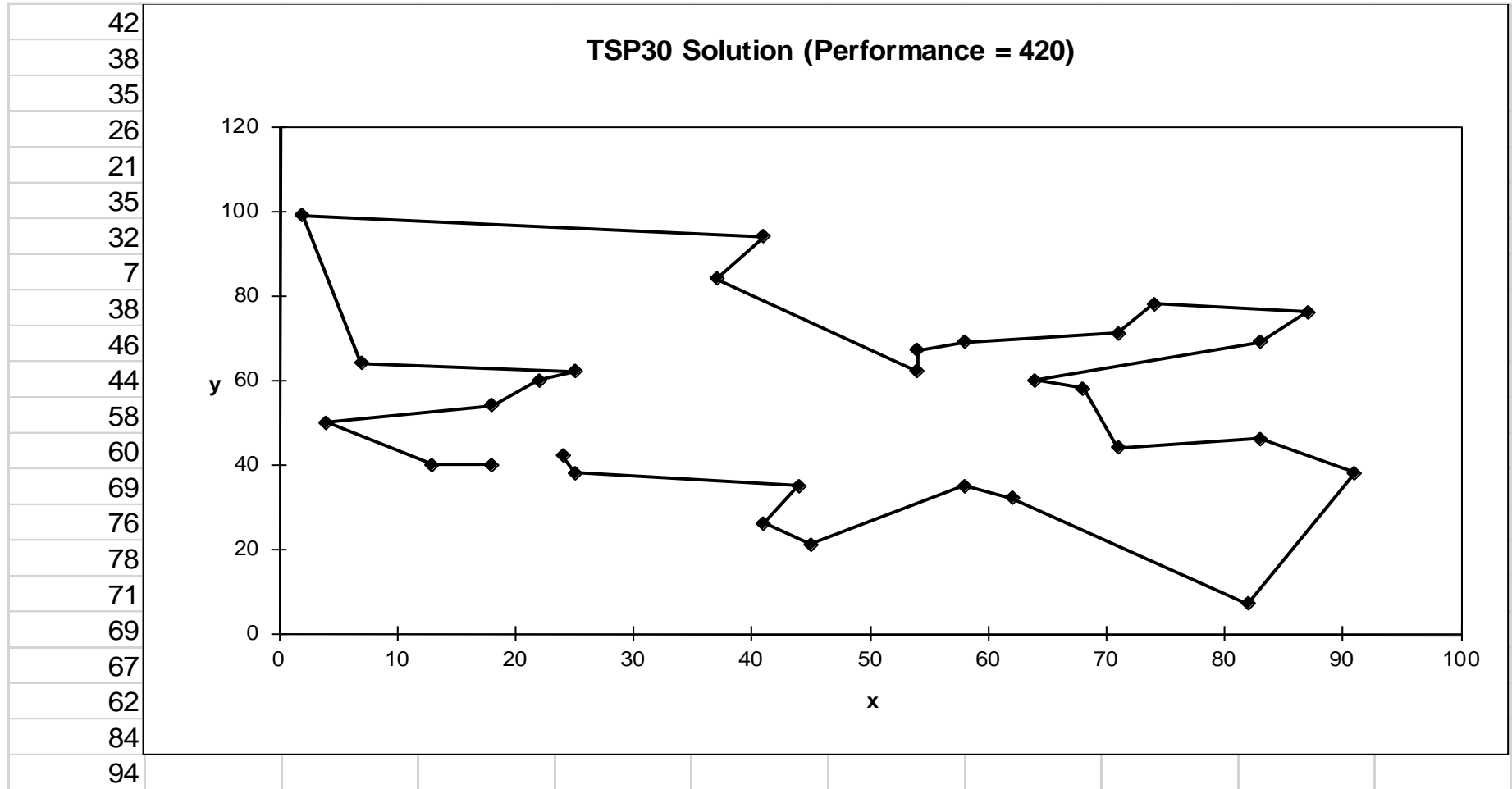
# Solution <sub>j</sub> (Distance = 800)



# Solution $k$ (Distance = 652)



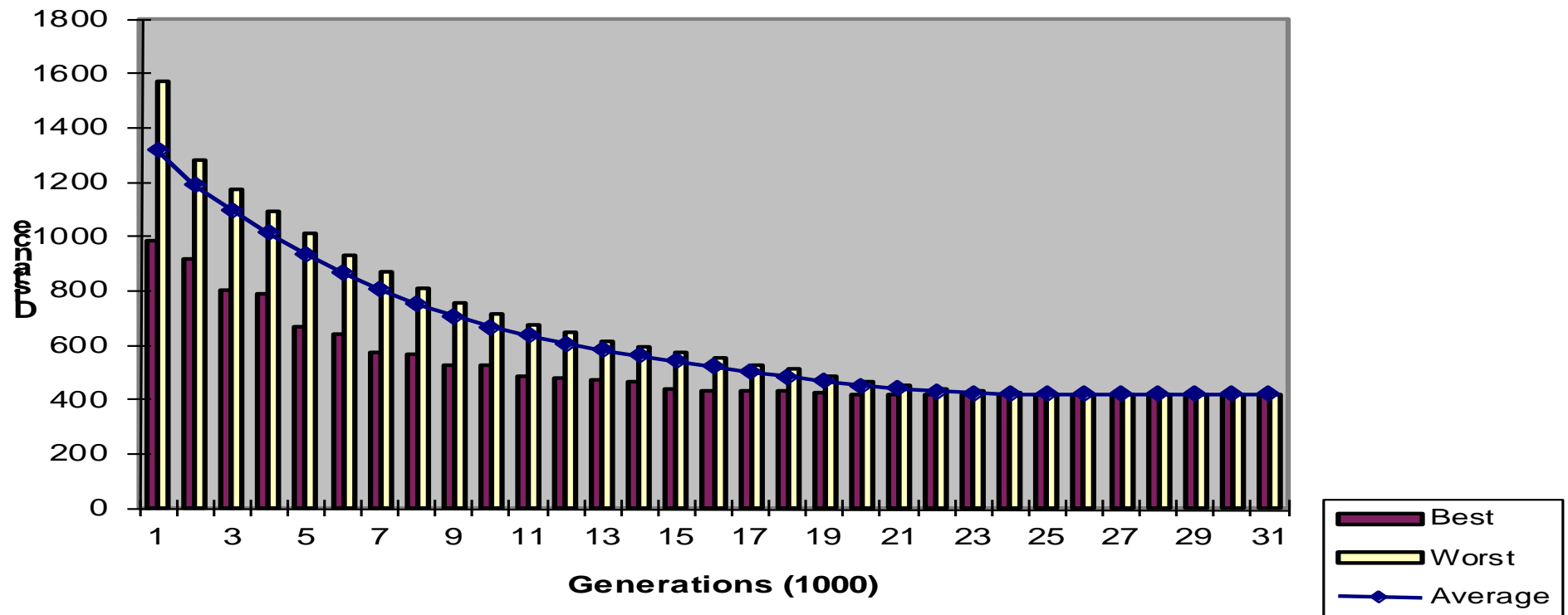
# Best Solution (Distance = 420)



# Overview of Performance



**TSP30 - Overview of Performance**



# GA and Decision trees

- Create many trees
- Define mating (interchanging branches) and mutation (replacing a node with another)
- Define objective function (accuracy)
- Pick parents and evolve!!

# Comparison

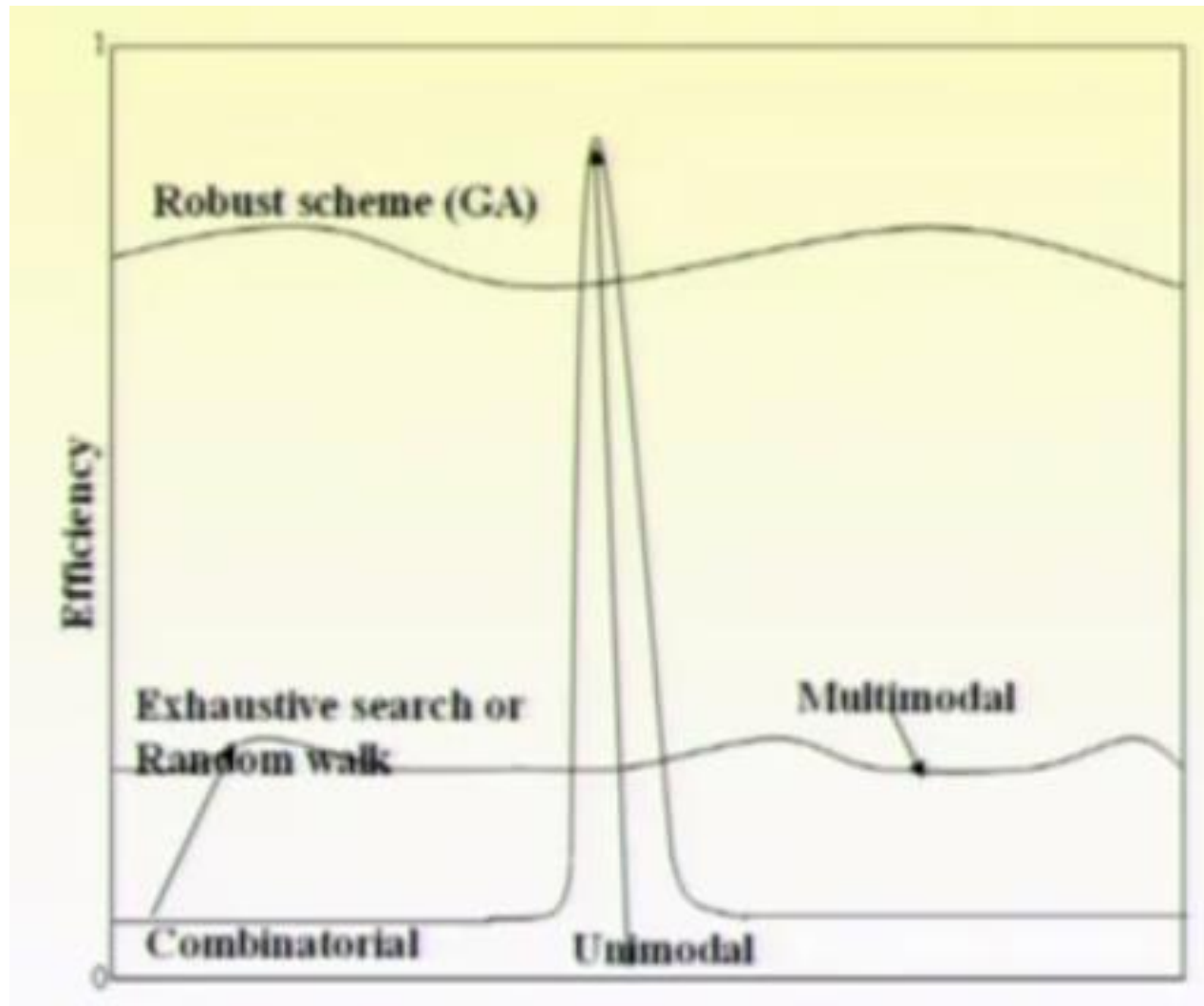
## Gradient descent and Traditional methods

- Require derivatives
- Gets stuck at local minima
- Quick where they work

## Genetic algorithms

- Substitute in the function
- Can yield global minima
- Time consuming





# Searching for optimal solution

- Enumerative search (search all possibilities): Works for very small data sets
- Random search: Works in cases where fluctuations are not much
- Calculus/Graphical driven search: Works well, fast but gets stuck at local minima
- Evolutionary search



## HYDERABAD

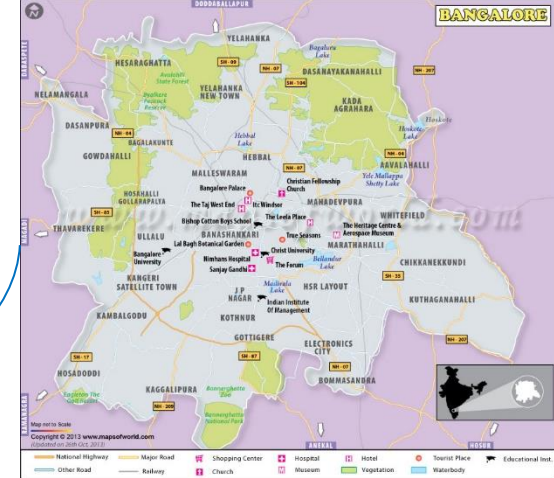
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