

# Computational Intelligence

## Unit # 10

## Tabu Search

- Tabu search uses common-sense ideas to enable the search process to escape from a local optimum.
- It uses a local search procedure to find a local optimum.
- A key strategy of tabu search is that it then continues the search by allowing *non-improving moves* to the best solutions in the neighborhood of the local optimum.
- The danger with this approach is that after moving away from a local optimum, the process will cycle right back on the same local optimum.

## Tabu List

- To avoid this, a tabu search temporarily forbids moves that would return to (or perhaps towards) a solution recently visited.
- A **tabu list** records these forbidden moves, which are referred to as tabu moves.
- The only exception to forbidding such a move is if it is found that a tabu move actually is better than the best feasible solution found so far.
- This use of *memory* to guide the search by using tabu lists to record some of the recent history of the search is a distinctive feature of tabu search.

## Sub-Tour Reversal in TSP

- A sub-tour reversal adjusts the sequence of cities visited in the current trial solution by selecting a subsequence of the cities and simply reversing the order in which that subsequences of cities is visited.
- The subsequence being reversed can consist of as few as two cities, but also can have more.
- The sub-tour reversal is an example of a local improvement procedure.

## Outline of a Basic Tabu Search Algorithm

- Initialization
  - Start with a feasible initial trial solution.
- Iteration
  - Use an appropriate local search procedure to define the feasible moves into the local neighborhood of the current trial solution.
  - Eliminate from consideration any move on the current tabu list unless that move would result in a better solution than the best trial solution found so far.
  - Determine which of the remaining moves provides the best solution and adapt it regardless of whether it is better or worse than the current trial solution.
  - Update the tabu list. If the tabu list is already full, delete the oldest member of the tabu list.
- Stopping Rule
  - Same as used for other algorithms discussed in this course

## Example Revisited

	A	B	C	D	E	F	G
A	0	12	10				12
B	12		8	12			
C	10	8		11	3		9
D		12	11		11	10	
E			3	11		6	7
F				10	6		9
G	12		9		7	9	

$$A - B - C - D - E - F - G - A : 12 + 8 + 11 + 11 + 6 + 9 + 12 = 69$$

## Working of Tabu Search

- Initial solution
  - A – B – C – D – E – F – G – A : Distance = 69
  - Tabu list: Blank at this point
- Iteration 1: Reverse C – D
  - Deleted Links: B – C, D – E
  - Added Links: B – D, C – E
  - Tabu List: B – D, C – E
  - A – B – D – C – E – F – G – A : Distance = 65
- Iteration 2: Reverse C – E – F
  - Deleted Links: D – C, F – G
  - Added Links: D – F, C – G
  - Tabu List: B – D, C – E, D – F, C – G
  - A – B – D – F – E – C – G – A : Distance = 64

## Working of Tabu Search (Cont'd)

- Iteration 3: Reverse C – G
  - Deleted Links: E – C, G – A
  - Added Links: E – G, C – A
  - Tabu List: ~~B – D, C – E~~, D – F, C – G, E – G, C – A
  - A – B – D – F – E – C – G – A : Distance = 66
- The process continues in a similar fashion until one of the stopping condition is satisfied.

## Simulated Annealing

- Developed in 1983, SA is another widely used metaheuristic that enables the search process to escape from a local optimum.
- The name and inspiration come from annealing in metallurgy, a technique involving heating and controlled cooling of a material to increase the size of its crystals and reduce their defects.
- The heat causes the atoms to become unstuck from their initial positions (a local minimum of the internal energy) and wander randomly through states of higher energy; the slow cooling gives them more chances of finding configurations with lower internal energy than the initial one.

## Analogy

- By analogy with this physical process, each step of the SA algorithm replaces the current solution by a random "nearby" solution, chosen with a probability that depends both on the difference between the corresponding function values and also on a global parameter  $T$  (called the *temperature*), that is gradually decreased during the process. The dependency is such that the current solution changes almost randomly when  $T$  is large, but increasingly "downhill" as  $T$  goes to zero.

## Working of the Algorithm

- At each step, the SA heuristic considers some neighboring state  $s'$  of the current state  $s$ , and probabilistically decides between moving the system to state  $s'$  or staying in state  $s$ .
- These probabilities ultimately lead the system to move to states of lower energy.
- Typically this step is repeated until the system reaches a state that is good enough for the application, or until a given computation budget has been exhausted.

## Method

- SA's major advantage over other methods is its ability to avoid becoming trapped in local minima.
- The algorithm employs a random search which not only accepts changes that decrease the objective function  $f$ , but also some changes that increase it.
- The latter are accepted with a probability
  - $P = \exp(-\Delta f/T)$
  - Where  $\Delta f$  is the increase in  $f$  and  $T$  is a control parameter "temperate".

## Outline of Simulated Annealing Algorithm

- Initialization
  - Start with a feasible initial trial solution
- Iteration
  - Use the *selection rule* to select the next trial solution.
  - If none of the immediate neighbors of the current trial solution are accepted, the algorithm is terminated.
- Check the temperature schedule
  - When the desired number of iterations have been performed at the current value of  $T$ , decrease  $T$  to the next value in the temperature schedule and resume performing iterations at this next value.
- Stopping Rule
  - Same as used for other algorithms discussed in this course

## TSP Example: Temperature Schedule

- Five iterations are performed at each of the five values of  $T$  ( $T_1, T_2, T_3, T_4, T_5$ ) in turn, where
  - $T_1 = 0.2Z_c$  when  $Z_c$  is the objective function value for the initial trial solution
  - $T_2 = 0.5T_1$
  - $T_3 = 0.5T_2$
  - $T_4 = 0.5T_3$
  - $T_5 = 0.5T_4$

## TSP Example: Working

- Initial Trial
  - A – B – C – D – E – F – G – A,  $Z_c = 69$ ,  $T1=0.2 \times Z_c = 13.8$
- Second Iteration
  - Pick two random numbers to find the swapping slots. Suppose the sub-tour of cities C – D is reversed.
  - A – B – D – C – E – F – G – A,  $Z_n = 65$
- Third Iteration
  - Reverse C – E – F
  - A – B – D – F – E – C – G – A,  $Z_n = 64$
- Fourth Iteration
  - Reverse C – G
  - A – B – D – F – E – G – C – A,  $Z_n = 66$  and  $Z_c = 64$  implies  $\Delta f = -2$
  - Prob {acceptance} =  $\exp(-2/13.8) = 0.865$
  - If the next random number generated is less than 0.865, this candidate solution will be accepted as the next trial solution. Otherwise it will be rejected.

## Differential Evolution

- Differential evolution (DE) is a population-based search algorithm.
- The algorithm draws inspiration from the field of evolutionary computation, as it embeds implicit concepts of mutation, recombination and fitness-based selection to evolve good solutions to a problem of interest by manipulating a population of solution encodings.
- An individual in DE is generally comprised of a real-valued chromosome.



## Working of DE

- At the start of the algorithm, a population of  $N$ ,  $d$ -dimensional vectors  $X_j = (x_{j1}, x_{j2}, \dots, x_{jd})$ ,  $j = 1, \dots, N$ , each of which encode a solution, is randomly initialized and evaluated using a fitness function  $f$ .
- During the search process, each individual ( $j$ ) is iteratively refined.

## Three Major Steps

- The modification process has three steps:
  - Create a variant vector which encodes a solution, using randomly selected members of the population (mutation step).
  - Create a trial vector, by combining the variant vector with  $j$  (crossover step).
  - Perform a selection process to determine whether the newly-created trial vector replaces  $j$  in the population.

## Mutation

- Under the mutation operator, for each vector  $X_j(t)$  a variant vector  $V_j(t+1)$  is obtained:
  - $V_j(t + 1) = X_m(t) + F(X_k(t) - X_l(t))$
  - where  $k, l, m \in 1, \dots, N$  are randomly selected indices, and all the indices  $\neq j$  ( $X_m$  is referred to as the base vector, and  $X_k(t) - X_l(t)$  is referred to as a difference vector).

## Crossover

- Selecting the three indices randomly implies that all members of the current population have the same chance of being selected, and therefore influencing the creation of the difference vector.
- The difference between vectors  $X_k$  and  $X_l$  is multiplied by a scaling parameter  $F$  (typically  $F \in (0, 2]$ ).
- The scaling factor controls the amplification of the difference between  $X_k$  and  $X_l$ , and is used to avoid stagnation of the search process.

## Self-Scaling Mutation

- A notable attribute of the mutation step in DE is that it is self-scaling.
- The size/rate of mutation along each dimension stems solely from the location of the particles in the current population.
- The mutation step self-adapts as the population converges leading to a finer-grained search.
- In contrast, the mutation process in the canonical GA is typically based on draws from a separately defined (fixed) probability density function.

## Cross Over

- Following the creation of the variant vector, a trial vector  $U_j(t+1) = (uj_1, uj_2, \dots, uj_d)$  is obtained:

$$U_{jk}(t+1) = \begin{cases} V_{jk}(t+1), & \text{if } (rand \leq CR) \text{ or } (j = rnbr(ind)) ; \\ X_{jk}(t), & \text{if } (rand > CR) \text{ and } (j \neq rnbr(ind)). \end{cases}$$

- where  $k = 1, 2, \dots, d$ ,  $rand$  is a random number generated in the range  $(0,1)$ ,  $CR$  is the user-specified crossover constant from the range  $(0,1)$ , and  $rnbr(ind)$  is a randomly chosen index chosen from the range  $(1, 2, \dots, d)$ .
- The random index is used to ensure that the trial solution differs by at least one component from  $X_j(t)$ .

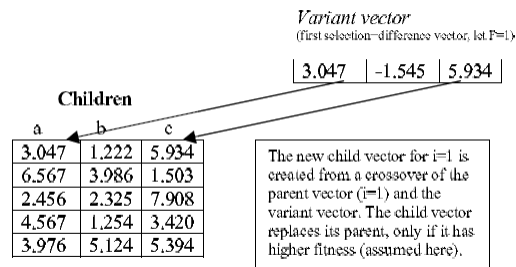
## Cross Over (Cont'd)

Index number	$X_i(t)$	$V_i(t+1)$		$U_i(t+1)$
1	a	q	$\text{rand}(1) > \text{CR}$	a
2	b	w	$\text{rand}(2) \leq \text{CR}$	w
3	c	e	$\text{rand}(3) \leq \text{CR}$	e
4	d	r	$4 = \text{mdbl}$	r

## Mutation and Cross Over

Parents				
a	b	c		
4.435	1.222	9.735	$i=1$	<i>First selection (<math>i_1</math>)</i>
6.567	3.986	1.503		4.567   1.254   3.420
2.456	2.325	7.908		<i>Second and third selections (<math>i_2</math> &amp; <math>i_3</math>)</i>
4.567	1.254	3.420		
3.976	5.124	5.394		2.456   2.325   7.908
				<i>Difference vector (<math>i_2 - i_3</math>)</i>
				-1.520   -2.799   2.514

## Mutation and Cross Over (Cont'd)



- The resulting trial (child) solution replaces its parent if it has higher fitness (a form of selection), otherwise the parent survives unchanged into the next iteration of the algorithm

## Summary of DE

- Initialize Population
- Evaluate fitness of population members
- Do
  - For each member of the population
    - Perform mutation operator by creating a variant vector with the help of three randomly selected individuals
    - Perform crossover operator by combining the variant vector with the individual. The resultant vector is called trial vector
    - Using a selection mechanism (Binary Tournament, etc.), decide if the trial vector replaces the original individual or not.
  - Next
- While (Not Finished)