

1 GRASP

- Input: $n \in N$, $h \in H$, minHours, maxHours, maxConsec, maxPresence
- Output: $W_{n,h}, \text{cost}$
- Procedure Construct
 - (1) Initialize $W_{n,h} = 0$, $\forall n \in N, \forall h \in H$
 - (2) for n in N
 - (3) $H' = H$
 - (3) ComputeGreedyCost(H')
 - (4) $\text{order1} = \text{sort}(H', 'gc', \text{asc})$
 - (5) $\text{min}_{gc} = \text{the gc of of first}(\text{order1})$, $\text{max}_{gc} = \text{the gc of the last}(\text{order1})$
 - (6) $\text{RCL} \leq \text{min}_{gc} + \alpha(\text{max}_{gc} - \text{min}_{gc})$
 - (7) Randomly select an hour (h) from RCL, set $W_{n,h} = 1$, remove(h)
 - (8) if complete(All $gc_h \geq 0$) then **break**
 - (9) while k ≤ max
 - (10) get the h_{start} & h_{end} of the working hours
 - (11) if $\text{Sum}(w_{n,h}) \geq \text{min}$ & $h_{end} - h_{start} + 1 == \text{maxPresence}$ then **break**
 - (12) $\text{order2} = \text{sort}(H'[h_{start}-2, h_{end} + 2], 'gc', \text{asc})$
 - (13) if satisfy maxConsec
 - (14) $h' = \text{first}(\text{order2})$, set $W_{n,h'} = 1$
 - (15) remove(h')
 - (16) $\text{cost} = \text{getNumberOfWorkingNurses}$
 - (17) return $W_{n,h}$, cost

End Construct
- Procedure Local Search
 - (1) $W''_{n,h} = W_{n,h}$
 - (2) for n ∈ NumberofWorkingNurse
 - (3) $W'_{n,h} = W''_{n,h}$
 - (4) for n' ∈ NumberofWorkingNurse \ n
 - (5) Set $W'_{n,h} = 0, W'_{n',h} = 0, \forall h \in H$
 - (6) if $\text{SumofWorkinghours}(\forall h \in H) \leq \text{Demand}_h - 2$ then **continue**
 - (7) elseif CanNotGenerateFeasibleW' then **continue**
 - (8) Generate a Feasible $W'_{n,h}$
 - (9) $W''_{n',h} = W'_{n',h}, W''_{n,h} = W'_{n,h}$
 - (10) cost = 1

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(11)      break
(12)      Update NumberofWorkingNurse
(13)  $W_{n,h}=W''_{n,h}$ 
(14) return  $W_{n,h}$ , cost
End Local Search

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- The function of greedy cost is as below:

$$gc = Current_h - Demand_h$$

2 BRKGA

The length of chromosome: $N \times H$

- Input: population, data
- Procedure Decoder
 - (1) for each chromosome in population
 - (2) for each gene in chromosome
 - (3) if gene ≥ 0.5 then sol=1
 - (4) else sol=0
 - (5) if satisfy min/max/maxConsec/MaxPresence/CannotRest2hours
 - (6) solution=append(sol)
 - (7) fitness=the Number of Rows with any gene ≥ 0.5
 - (8) else solution=null, fitness= ∞
 - (9) Return (chromosome,solution, fitness)
 End decoder