EE496: COMPUTATIONAL INTELLINGENCE

EA02: N-QUEEN: AN EA SOLUTION

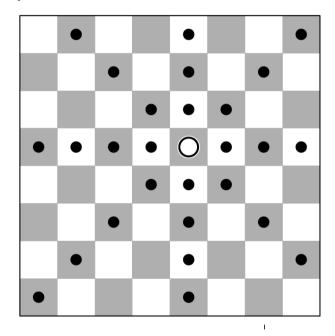
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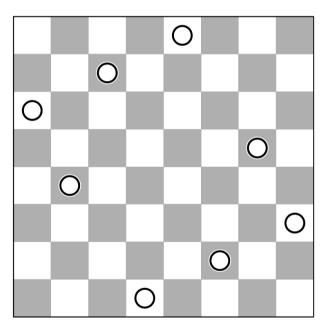
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The n-Queens Problem

place n queens onto a n x n chessboard in such a way that no row (rank), no column (file) and no diagonal contains more than one queen or: place queens in such a way that no queen is in the way of another queen







A solution for n-queen problem

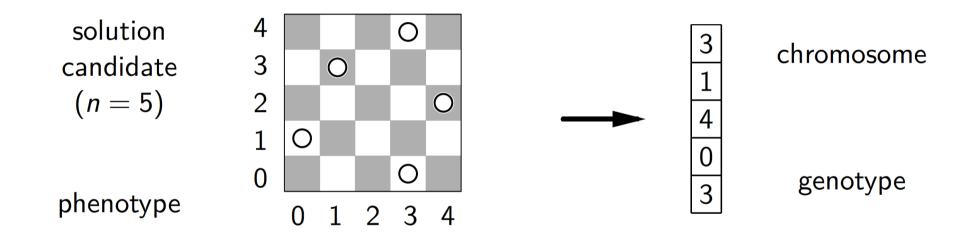
The n-Queens Problem

Backtracking Solution of the n-Queens Problem

- 1. place queens row-by-row from bottom to up (or column by column from left to right)
- 2. consider each row as follows:
 - place one queen in a rank sequentially from left to right onto the squares of the board
 - for each placement: check if queen collides with queens in lower rows
 - if not, work on next row recursively
 - afterwards: shift queen one sqare rightwards
- 3. return solution if queen is placed in top row with no collision

EA: Encoding

- Representation: 1 solution candidate = 1 chromosome with n genes
- each gene: one row of the board with n possibles alleles
- value of the gene: position of the queen in corresponding rank



solution candidates with > 1 queen each rank not permitted ⇒ smaller search space

EA: data structure

- data type for 1 chromosome, which stores the fitness
- data type for 1 chromosome with buffer for "intermediate population" and flag for the best individual

```
/* --- an individual --- */
typedef struct {
    int fitness;
                         /* fitness (number of collisions) */
                         /* number of genes (number of rows) */
    int n;
                         /* genes (queen positions in rows) */
    int genes[n];
                         /* (individual) */
} IND;
                         /* --- a population --- */
typedef struct {
                         /* number of individuals */
    int size;
                         /* vector of individuals */
    IND **inds;
    IND **buf;
                         /* buffer for individuals */
    IND *best;
                         /* best individual */
                         /* (population) */
} POP;
```

EA: main loop

shows basic form of an EA:

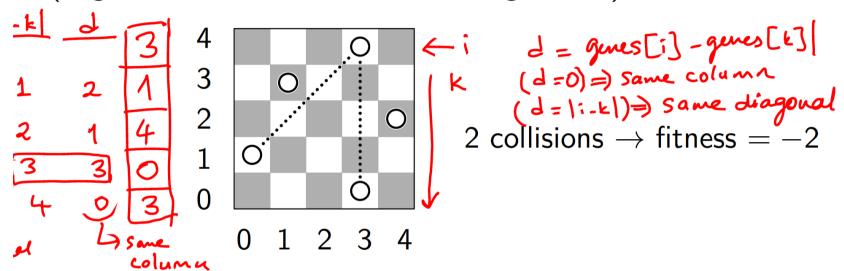
```
pop_init(pop);
                                  /* initialize the population */
while ((pop_eval(pop) < 0)
                                 /* while no solution found and */
                                 /* not all generations computed */
&& (gencnt >= 0)) {
    pop_select(pop, tmsize, elitist); /* select individuals (natural selection) */
    pop_cross (pop, frac); /* do crossover (parental sel. & crossover */
    pop_mutate(pop, prob); /* and mutate individuals */
parameters:
gencht: maximum amount of remaining generations
tmsize: size of tournament selection
elitist: indicates, if best individual will always be taken
        fraction of individuals, which will be submitted by cross-over
frac:
prob:
        mutation probability
```

EA: Initialize

```
create random series of n numbers from \{0, 1, ..., n-1\}
/* 0 < drand() < 1, so 0 <= (int) n * drand() < n-1 */
void ind init (IND *ind)
                                 /* --- initialize an individual */
                                 /* loop variable */
    int i;
    for (i = ind->n; --i >= 0; ) /* initialize the genes randomly */
        ind->genes[i] = (int)(ind->n *drand());
    ind->fitness = 1;
                       /* fitness is not known yet */
                                 /* ind init() */
void pop init (POP *pop)
                                 /* --- initialize a population */
                                 /* loop variable */
int i;
for (i = pop->size; --i >= 0;)
ind_init(pop->inds[i]);
                                 /* initialize all individuals */
} /* pop_init() */
```

EA Evaluation

ullet fitness: negated number of columns and diagonals with ≥ 1 que (negated number due to maximizing fitness)



- if queens in 1 column/diagonal \geq 2: count each pair (easier to implement)
- fitness-function results immediately in termination criterion: Solution has (highest possible) fitness 0
- also: termination is guaranteed when maximal generation is reached

EA Evaluation

count collisions by computation on chromosomes:

```
int ind eval (IND *ind)
                             /* --- evaluate an individual */
                             /* loop variables */
 int i, k;
                             /* horizontal distance between queens */
 int d;
                             /* number of collisions */
 int n;
 if (ind->fitness <= 0) /* if the fitness is already known, */
   return ind->fitness; /* simply return it */
 for (n = 0, i = ind->n; --i > 0;)
   for (k = i; --k \ge 0;) { /* traverse all pairs of queens */
     d = abs(ind->genes[i] -ind->genes[k]);
     if ((d == 0) || (d == i-k)) n++;
              /* count the number of pairs of queens */
                     /* in the same column or diagonal */
 return ind->fitness = -n; /* return the number of collisions */
} /* ind eval() */
                      Lyif a is large, fitness is poor
```

EA Evaluation

- calculation of the fitness of all individuals of the population
- simultaneously: determination of best individual
- best individual fitness $0 \Rightarrow$ solution is found

EA: selection of individuals (for next generation)

tournament selection:

- consider tmsize arbitrarily chosen individuals
- best (of these) individual "wins" tournament and will be chosen
- the higher the fitness the better chance to get chosen

cubose initial best randomly

choose ind randomly

match between indix best

which ever wins is the new

sest

repeat thisize three

refure best

EA: selection of individuals (for next generation)

tournament selection:

- consider tmsize arbitrarily chosen individuals
- best (of these) individual "wins" tournament and will be chosen
- the higher the fitness the better chance to get chosen

EA: selection of individuals

- tournament selection for individuals of the next population generation
- perhaps best individuals will be applied (and not changed)

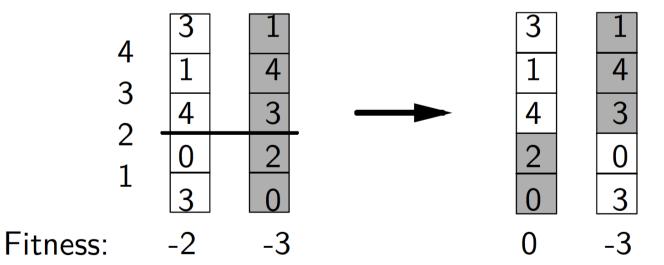
```
/* --- select individuals */
( Ang < dod spriedod)
    int i;
    /* loop variables */
    /* exchange buffer */

i = pop->size;
    /* select 'popsize' individuals */
    if (elitist)

/* preserve the best individual */
       ind_copy(pop->buf[--i], pop->best);
    while (--i >= 0) /* select (other) individuals */
       ind_copy(pop->buf[i], pop_tmsel(pop, tmsize)); kst 9
  p = pop->inds; pop->inds = pop->buf;
   pop->buf = p;
                      /* set selected individuals */
     pop->best = NULL; /* best individual is not known yet */
  } /* pop select() */
```

Crossover

- Exchange of a piece of the chromosomes between two individuals
- here: so called One-Point-Crossover
 - choose cutting line between two genes by random
 - change sequences of genes on one side of the cutting line
 - Example: choose cutting line 2



EA: Crossover

Exchange of pieces of the chromosomes between two individuals

```
void ind cross (IND *ind1, IND *ind2)
{
                       /* --- crossover of two chromosomes */
                       /* loop variable */
 int i;
                       /* gene index of crossover point */
 int k;
                       /* exchange buffer */
 int t;
 k = (int)(drand() *(ind1->n-1)) +1; /* choose a/crossover point */
 = ind1->genes[i];
  ind1->genes[i] = ind2->genes[i];
 ind2->genes[i] = t; /* exchange genes */
                     /* of the chromosomes */
 ind1->fitness = 1;  /* invalidate the fitness */
 ind2->fitness = 1; /* of the changed individuals */
} /* ind cross() */
```

EA: Crossover

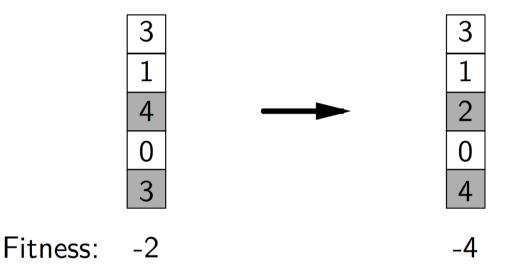
- certain rate of individuals is submitted by crossover
- include of both crossover-products in new population
- "parental individuals" are getting lost
- no crossover on best individual (if taken over)

crossover

inds

EA Mutataion

- replacement of randomly chosen genes (changing of alleles)
- perhaps number of replaced genes is chosen by random (number of replaced genes should be as small as possible)



- mutations are mostly damaging (decrease the fitness)
- non existing alleles can be created by mutation

- decide whether to continue mutating or not
- best individual (if taken over) won't be submitted to mutation

```
void ind mutate (IND *ind, double prob)
                                  /* --- mutate an individual */
    if (drand() >= prob) return; /* det. whether to change individual */
do ind->genes[(int)(ind->n *drand())] = (int)(ind->n *drand());
Loop (while (drand() < prob); /* randomly change random genes */
     } /* ind_mutate() */
   void pop_mutate (POP *pop, double prob)
                                  /* --- mutate a population */
                                  /* loop variable */
     int i;
       ind_mutate(pop->inds[i], prob); ) mutates some of individuals

* pop_mutate() * '
     for (i = pop->size -1; --i >= 0;)
   } /* pop_mutate() */ /* mutate individuals */
```