

Branch and bound – number of selected children

Heuristics – Case: Fruit fly

Authors: Mercyllyn Wiemer (10749306), Shan Shan Huang (10768793) & Marwa Ahmed (10747141)

Goal of experiment:

In this experiment we change the number of selected children in the branch and bound (BnB): depth-first search which prunes using breakpoints. It will show what the influence of this constrain is on the run time and the numbers of inversions made to find the solution.

Methods:

The BnB depth-first search with breakpoints searches for a path from a given genome to a solution genome. The path consists of reversions of gene blocks in a genome. From the given root genome, every possible reversion is generated: the children. Depending on the number of selected children, one, two, or three children with the least breakpoints are explored first. The upper bound is updated, every time a solution is found.

Results:

Table1. Branch and bound: Influence of the number of selected children on the runtime and number of inversions needed to get to the solution

| No of selected children (fruit flies) | Number of inversions to solution | Runtime |
|---------------------------------------|----------------------------------|--------------------|
| 1 | 17 | 0.4 sec |
| 2 | 13 | 144 sec / 2.6 min. |
| 3 | - | > 10min |

Branch and Bound: breakpoints

genome fruitfly: [23, 1, 2, 11, 24, 22, 19, 6, 10, 7, 25, 20, 5, 8, 18, 12, 13, 14, 15, 16, 17, 21, 3, 4, 9]
Solution found in generation: 17, but still searching for a better solution

Best solution found in generation: 17

Path to solution:

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swap: 0 [23, 1, 2, 11, 24, 22, 19, 6, 10, 7, 25, 20, 5, 8, 18, 12, 13, 14, 15, 16, 17, 21, 3, 4, 9]
swap: 1 [23, 1, 2, 11, 24, 22, 19, 20, 25, 7, 10, 6, 5, 8, 18, 12, 13, 14, 15, 16, 17, 21, 3, 4, 9]
swap: 2 [23, 1, 2, 11, 24, 22, 19, 20, 25, 7, 10, 6, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 3 [11, 2, 1, 23, 24, 22, 19, 20, 25, 7, 10, 6, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 4 [7, 25, 20, 19, 22, 24, 23, 1, 2, 11, 10, 6, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 5 [10, 11, 2, 1, 23, 24, 22, 19, 20, 25, 7, 6, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 6 [10, 11, 12, 13, 14, 15, 16, 17, 21, 3, 4, 5, 6, 7, 25, 20, 19, 22, 24, 23, 1, 2, 18, 8, 9]
swap: 7 [10, 11, 12, 13, 14, 15, 16, 17, 25, 7, 6, 5, 4, 3, 21, 20, 19, 22, 24, 23, 1, 2, 18, 8, 9]
swap: 8 [10, 11, 12, 13, 14, 15, 16, 17, 22, 19, 20, 21, 3, 4, 5, 6, 7, 25, 24, 23, 1, 2, 18, 8, 9]
swap: 9 [10, 11, 12, 13, 14, 15, 16, 17, 18, 2, 1, 23, 24, 25, 7, 6, 5, 4, 3, 21, 20, 19, 22, 8, 9]
swap: 10 [10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 3, 4, 5, 6, 7, 25, 24, 23, 1, 2, 22, 8, 9]
swap: 11 [10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 2, 1, 23, 24, 25, 7, 6, 5, 4, 3, 8, 9]
swap: 12 [10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 2, 1, 23, 24, 25, 3, 4, 5, 6, 7, 8, 9]
swap: 13 [10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 25, 24, 23, 1, 2, 3, 4, 5, 6, 7, 8, 9]
swap: 14 [10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 1, 2, 3, 4, 5, 6, 7, 8, 9]
swap: 15 [25, 24, 23, 22, 21, 20, 19, 18, 17, 16, 15, 14, 13, 12, 11, 10, 1, 2, 3, 4, 5, 6, 7, 8, 9]
swap: 16 [25, 24, 23, 22, 21, 20, 19, 18, 17, 16, 15, 14, 13, 12, 11, 10, 9, 8, 7, 6, 5, 4, 3, 2, 1]
swap: 17 [1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25]
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runtime: 0.38631903028792564

Figure 1. Result of BnB depth-first search when selecting one child.

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Branch and Bound: breakpoints

genome fruitfly: [23, 1, 2, 11, 24, 22, 19, 6, 10, 7, 25, 20, 5, 8, 18, 12, 13, 14, 15, 16, 17, 21, 3, 4, 9]
Solution found in generation: 18, but still searching for a better solution
Solution found in generation: 17, but still searching for a better solution
Solution found in generation: 16, but still searching for a better solution
Solution found in generation: 15, but still searching for a better solution
Solution found in generation: 14, but still searching for a better solution
Solution found in generation: 13, but still searching for a better solution

Best solution found in generation: 13
Path to solution:
swap: 0 [23, 1, 2, 11, 24, 22, 19, 6, 10, 7, 25, 20, 5, 8, 18, 12, 13, 14, 15, 16, 17, 21, 3, 4, 9]
swap: 1 [23, 1, 2, 11, 24, 22, 19, 6, 10, 7, 25, 20, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 2 [24, 11, 2, 1, 23, 22, 19, 6, 10, 7, 25, 20, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 3 [24, 11, 2, 1, 23, 22, 19, 20, 25, 7, 10, 6, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 4 [1, 2, 11, 24, 23, 22, 19, 20, 25, 7, 10, 6, 5, 4, 3, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 5 [1, 2, 3, 4, 5, 6, 10, 7, 25, 20, 19, 22, 23, 24, 11, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 6 [1, 2, 3, 4, 5, 6, 7, 10, 25, 20, 19, 22, 23, 24, 11, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 7 [1, 2, 3, 4, 5, 6, 7, 24, 23, 22, 19, 20, 25, 10, 11, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 8 [1, 2, 3, 4, 5, 6, 7, 20, 19, 22, 23, 24, 25, 10, 11, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 9 [1, 2, 3, 4, 5, 6, 7, 20, 19, 11, 10, 25, 24, 23, 22, 21, 17, 16, 15, 14, 13, 12, 18, 8, 9]
swap: 10 [1, 2, 3, 4, 5, 6, 7, 20, 19, 11, 10, 9, 8, 18, 12, 13, 14, 15, 16, 17, 21, 22, 23, 24, 25]
swap: 11 [1, 2, 3, 4, 5, 6, 7, 17, 16, 15, 14, 13, 12, 18, 8, 9, 10, 11, 19, 20, 21, 22, 23, 24, 25]
swap: 12 [1, 2, 3, 4, 5, 6, 7, 17, 16, 15, 14, 13, 12, 11, 10, 9, 8, 18, 19, 20, 21, 22, 23, 24, 25]
swap: 13 [1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25]

runtime: 144.2186030793153

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Figure 2. Result of BnB depth-first search when selecting two children.

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Branch and Bound: breakpoints

genome fruitfly: [23, 1, 2, 11, 24, 22, 19, 6, 10, 7, 25, 20, 5, 8, 18, 12, 13, 14, 15, 16, 17, 21, 3, 4, 9]
Solution found in generation: 22, but still searching for a better solution
Solution found in generation: 21, but still searching for a better solution
Solution found in generation: 19, but still searching for a better solution
Solution found in generation: 18, but still searching for a better solution
Solution found in generation: 17, but still searching for a better solution

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Figure 3. Result of BnB depth-first search when selecting three children.

Discussion:

Selecting one child has a fast runtime of less than one second and finds a solution in 17 inversions (Figure 1). While selecting two children has a slightly longer runtime, runtime of 2.6 minutes, but finds a solution in just 13 inversions (Figure 2). Selecting three children, increased the runtime up to more than 10 minutes, and after 10 minutes it does not find a better solution than 17 inversions (Figure 3).

Conclusion:

Selecting two children works better than selecting one or three, when taking in consideration the number of inversions and runtime.