

Animating a Sudoku solver

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Level 4 Project — November 18, 2015

Abstract

Sudoku is a popular puzzle played all over the world. It consists of filling in a 9x9 grid such that every row, column and 3x3 sub-grids have different digits from 1 to 9. Solving the puzzle will make use of the all-different algorithm from Constraint Programming for which an implementation will be provided. Finally, the program will animate all the steps done by the algorithm.

Education Use Consent

| I hereby give my perm | ission for this project to be shown to other University of Glasgow students and to be |
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| distributed in an electro | nic format. Please note that you are under no obligation to sign this declaration, but |
| doing so would help fu | ture students. |
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Introduction

1.1 Aims

Teaching constraint programming, need for visualisations

1.2 Background

1.3 Sudoku

Sudoku is a puzzle

1.4 Constraint Programming

Every Constraint Satisfaction Problem (CSP) has variables with domains and constraints associated to them. To start solving Sudoku puzzles, we have to come up with a model representation of the problem. For the purpose of this exercise, there will be 81 variables representing the 9x9 cells in the Sudoku grid, each with an initial domain of integers from 1 to 9. As all the initial puzzles start with a partially completed grid, values that are known will be reflected as variables with an initial domain of one digit, the known value. Such variables will therefore keep their initial domain throughout the algorithm, as no other values are present in their domain.

The constraints for a Sudoku puzzle can be expressed in natural language as such: digits may not appear twice in the same row, column or 3x3 subregions. The rules of the puzzle highlight the need of thinking in terms of 9 rows, 9 columns and 9 subregions, each of them having digits from 1 to 9 with no repetition. We can create 36 not equals constraints between the 9 variables of each row, column, subregions times 27 such regions for a total of 972 constraints between two variables. This would be highly inefficient, hence the need for a better approach. A better approach is to use 27 all-different constraints, one for each row, column and subregions. Each constraint contains 9 variables that should take different values. As each variable is part of a row, column and subregion at the same time, it will therefore be involved in 3 constraints.

Initial puzzles are always partially completed with enough values that by the end of the game there will be a single unique solution to the puzzle.

We get to the solution by running the alldifferent algorithm on rows, columns and subregions to remove values from the domain of the variables. Initially, the known values given at the start of the puzzle will be removed from the domains of possible values in the variables that are part of the same row, column or subregion. As the algorithm progresses, we will eventually remove enough values from the domain of a variable until we are left with a single value that represents the correct answer for that position. Once a value is found, there is a waterfall-like effect across the puzzle as we propagate the knowledge across the grid. The propagation takes place across all the variables that are in the same row, column and subregion with the one just solved. This will keep the 3 constraints in a consistent state, as all the variables will have valid remaining domains not containing any already solved values.

The algorithm is intelligent in a sense that it mimics the thinking of humans when solving a Sudoku puzzle. At some stages of the game we have enough information to know that two particular values will fill two cells, but no information about which value corresponds to which cell. This problem is what makes some Sudoku puzzles so hard to solve.

Humans and the alldifferent algorithm are smart enough to use this information to their advantage and remove the two values from the domains of the adjacent variables. By using this technique, we can ensure the progress of the algorithm towards a solution, even though we still dont know which two variables take which two values. It is good to note that although humans may choose a guessing approach to continue towards a solution, the alldifferent algorithm performs no guessing and works only with the information about the current state of the puzzle.

1.5 The Alldifferent Constraint

Review Regins paper and explain how constraint works [2]

Example of its usage and how it works (maybe 5 variables with domains)

Algorithms for AllDiff

2.1 Ford Fulkerson

A flow represents a directed graph with special nodes called the source and the sink. Source nodes have outgoing edges to inner nodes, while sink nodes have incoming edges from inner nodes of the graph. The problem requires sending flows, for example water through a pipe, from the source to the sink, while conserving the equilibrium.

Flows need to get from the source to the sink by following the directed edges that often have different maximum capacity constraints, such as the diameter of a water pipe. Given a flow f in graph G, we build a residual directed graph G_f . Forward edges in the graph havent reached their maximum capacity, while backward edges reached their upper bound.

Ford-Fulkerson algorithm uses the residual graph to achieve a maximum flow in the graph. The algorithm looks for augmenting paths from source S to sink T in a breadth first-search manner. Once a path p is found, we send the weight that will make the bottleneck edge in p fill its maximum capacity. Then, we add (subtract) the weight of each forward (backward) edge in p. Forward edges that reach their maximum capacity turn into backward edges. This results in flow f that has a greater value than before. Berges theorem guarantees that once all the augmenting paths have been successively visited in the algorithm, the end result is a maximum flow of the graph [?].

The maximum matching problem is a special case of the maximum flow problem. For the purpose of our example, there will be only one source S that has outgoing edges to the nodes in the left partition L of the bipartite graph and one sink T with incoming edges from the right partition R of the graph. Nodes in L are connected through edges to nodes in R. A matching TODO. A maximum matching TODO. To find a maximum matching for a Graph G, we add a source and sink nodes and set all edges to a maximum capacity of one. We then run the Ford-Fulkerson algorithm on the graph, by finding all the augmenting paths from free nodes in L to free nodes in R.

Having a maximum matching for our graph, the next step in the all different algorithm is to turn the previously undirected graph, into a directed one. The matches resulted from the Ford Fulkerson algorithm are assigned a direction from the 1 to 9 values to the corresponding cells of the Sudoku row. The edges that remain unused in the matching are given a direction from left to right (i.e. from the Sudoku cells of the row to the 1-9 values).

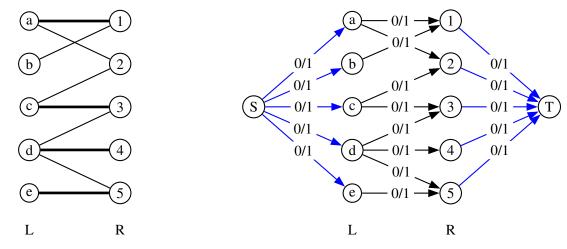


Fig. 2.1: Initial graph

Fig. 2.2: S&T nodes added

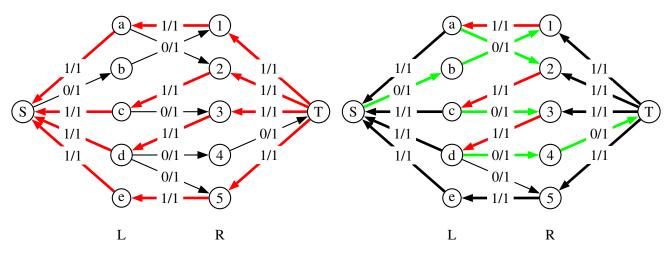


Fig. 2.3: Greedy match

Fig. 2.4: Longer augmenting path

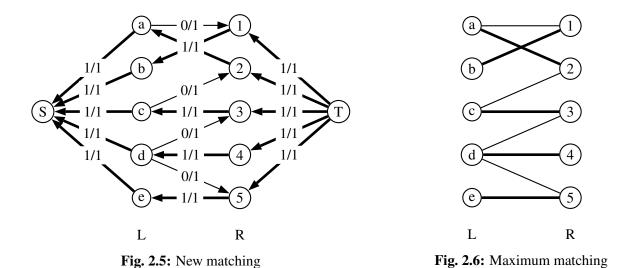


Fig. 2.7: FordFulkerson algorithm

```
Algorithm 1: Ford Fulkerson
```

```
1 void FordFulkerson(Graph G)
2 begin
3
        Global int capacity[][] \leftarrow getCapacity(G)
        Global int n \leftarrow |V(G)|
4
        Global int source \leftarrow 0
5
        Global int sink \leftarrow n-1
6
       Global pred \leftarrow new int[n]
7
        Global visited \leftarrow new boolean[n]
8
        Global Q \leftarrow new \, \mathbf{Queue}()
9
        while bfs(G) do
10
            update()
11
12 boolean bfs(Graph G)
13 begin
       clear(Q)
14
        fill(pred, -1)
15
        fill(visited, false)
16
        enqueue(source, Q)
17
        visited[source] \leftarrow true
18
       while (\neg isEmpty(Q)) do
19
            int v \leftarrow dequeue(Q)
20
            if v = sink then return true
21
            for w \leftarrow 0 to n do
22
                if \neg visited[w] and capacity[v][w] > 0 then
23
                     pred[w] \leftarrow v
24
                     enqueue(w, Q)
25
                     visited[w] \leftarrow true
26
        return false
27
28 void update()
29 begin
       int f \leftarrow minCost()
30
       \mathbf{int} \; v \leftarrow sink
31
32
        while pred[v] \neq -1 do
            int u \leftarrow pred[v]
33
            capacity[u][v] \leftarrow capacity[u][v] - f
34
            capacity[v][u] \leftarrow capacity[v][u] + f
35
            v \leftarrow u
36
37 int minCost()
38 begin
       \mathbf{int} \; minCost \leftarrow \infty
39
       \mathbf{int} \; v \leftarrow sink
40
        while pred[v] \neq -1 do
41
            minCost \leftarrow minimum(minCost, capacity[pred[v]][v])
42
            v \leftarrow pred[v]
43
        {f return}\ minCost
44
```

Ford Fulkerson's algorithm for finding a maximum flow in a graph is shown in algorithm 1. [1]. The Ford Fulkerson procedure on line 1 takes a graph G as an argument and represents the start of the algorithm. The graph passed as an argument holds information about the capacity of each edge. Line 3 stores the capacity of the graph G in a global two-dimensional integer array. The capacity of an edge between vertices u, v is stored in capacity[u][v]. Line 4 declares an integer n used to store the number of vertices in the given graph. Lines 5 and 6 declare two integers called source and sink representing references to the first and the last vertices in the given graph.

In line 7 we declare a vertex-indexed array pred that is used to store references to the previous vertex discovered in the breadth-first search tree. Line 8 declares a vertex-indexed array visited used to keep track if a vertex has already been visited in the breadth-first search procedure. Line 9 declares a queue of integers Q used to store vertices in the order they are visited.

The algorithm consists of repeated calls to the breadth-first search procedure (line 10). In line 11, we call the update procedure to update the capacities of the edges after finding augmenting paths.

The breadth-first search procedure starts at line 12 and takes a graph G as an argument. In lines 14 to 16, the queue Q, pred and visited array are reset to have default values. The breadth-first search calls always start from the first vertex, representing the source. This vertex is enqueued on queue Q in line 17 and is marked as visited in line 18.

Lines 19 to 26 contain a loop performing changes on the queue Q that is used to store an augmenting path. The loop starts on line 20 by dequeuing a vertex from the queue Q and storing it in v. Line 21 forces the breadth-first search procedure to return true when we reach the last vertex in the graph, the sink. Lines 22 to 26 contain a loop over all the vertices w in the graph. In line 23 we check to see if the discovered vertex w hasn't been already visited and we consider it if it still has available capacity. Line 24 stores the index value of w's predecessor in the augmenting path. Line 25 enqueues the discovered vertex w on the queue Q and marks it as visited in line 26. Finally, the breadth-first search procedure returns false when no more augmenting paths are found.

The update procedure at line 28 is used to update the capacities of the edges in the graph to reflect the updated flow. Line 30 declares an integer f used to store the amount of flow we are adding along the found path. The value for f is calculated in the minCost procedure at line 37. Line 31 declares an integer v, initially representing the sink which is the last vertex in the graph. Lines 32 to 35 contain a loop that updates the capacities of the edges in the augmenting path. Line 33 uses the integer u to store the index of a vertex that is the predecessor of vertex v along the path. Lines 34 and 35 updates the flow along the edge from u to v and v to v by the found difference. Finally, v is updated on line 36 to take the value of v such that the while loop continues to change the capacities of the remaining edges along the path.

The minCost procedure at line 37 is used to find the amount of flow to be sent along the new augmenting path from the sink to the source. The value is equal to the minimum capacity that is still available along the edges of the path. Line 39 declares an integer minCost used to represent the current value of the minimum cost, initialized to a big value. Line 40 declares an integer variable v initially used to store a reference to the sink vertex. Lines 41 to 43 contain a loop over the edges along the path, starting from the sink toward the source. Line 42 updates the value of minCost if the current edge in the loop has a smaller capacity left than what was previously discovered. The minimum procedure returns the minimum of two given integers. Line 43 updates the value of integer v to its predecessor so the loop can continue along the path until it reaches the source. At the end of the minCost procedure, we return the value of the currently found minCost variable.

2.2 Tarjan algorithm

A strongly connected component SCC of a directed graph G is a maximal set of vertices such that every vertex is reachable from every other vertex by following the directed edges.

There are four kinds of edges in a depth-first traversal: tree edges, forward edges, back edges and cross edges. Tree edges are edges between a vertex and vertices that were not previously visited. Forward edges are from ancestor vertices to descendant vertices that were already visited during the search. Back edges link descendant vertices back to already visited ancestor vertices. Lastly, cross edges represent edges between two vertices that are neither an ancestor nor a descendant for each other, such as the edges between different components of the graph.

For each vertex in a graph G, the algorithm keeps track of two properties. The first property we remember is the index of the node, its consecutive order of discovery during the depth-first search, a value that will remain constant throughout the algorithm. The second property associated with all the nodes is named low and keeps track of the lowest (oldest) ancestor reachable from their position in the graph. As each node of the graph gets discovered in a depth-first search manner, the initial value that low gets assigned is the same as the order the node was discovered. The value for low may get changed during the search if an ancestor for the node is discovered by following a back edge. After the recursive depth-first search finishes visiting all of a nodes neighbours, the nodes value for low gets updated to the lowest index of its oldest reachable ancestor. If the value for low of a node stays the same even after visiting all of its neighbours, then it means it is the root of a strongly connected component or its an individual vertex that will be a component on its own. [3]

For the purpose of finding SCCs, every time a node is discovered in the search, the node is put on a stack S. After returning from the recursive DFS calls on all of the adjacent neighbours of vertex, if both index and low have the same value, it means weve finished discovering a SCC. The vertices in the newly discovered strongly connected component will be popped from the stack S until we reach the current vertex representing the root of the SCC. Vertices that are part of other SCCs are still left on the stack as they have different roots represented by the value of low.

Figure 2.8 TODO shows a bipartite directed graph with 18 vertices. There are 9 backward edges that represent the maximum matching found using the Ford-Fulkerson algorithm in section TODO. The rest of the possible edges are assigned a forward direction. This graph is similar to the graphs that will be generated to represent a row, column or one of the nine sub-grids of the Sudoku puzzle. For example, the left partition of the graph will contain the nine variables of a Sudoku row, and the right partition will contain the digits from 1 to 9. The edges will then link each variable of the row to values that are still available in their domain.

The Table 2.1 TODO contains the output after running the Tarjans algorithm on the graph in figure 2.8 TODO. As previously discussed, vertices are visited in a depth-first search manner, in this case starting from the first vertex. Each node has two properties associated to it: its order of discovery or *inded*, and its oldest reachable ancestor. Initially, both properties take the value of the index, but as the algorithm progresses the values for low may become lower after visiting all of the neighbours of the node and finding the oldest (lowest) reachable ancestor. All vertices are pushed on a stack in the order they are visited. Once a back edge is found in the graph, vertices that are currently on the stack get popped including the root of the newly discovered SCC. Note that vertices may still remain on the stack as they will be part of different components that have other roots. As each vertex is made part of a component, its value for *low* gets updated to a large value to mark it as already part of a component such that it will not become the root of another component in the future.

The graph in figure 2.9 TODO shows the 5 strongly connected components identified after running Tarjans algorithm on the initial graph in figure 2.8 TODO. Vertices that have the same background are part of the same component; meaning that one vertex can reach every other vertex in the component by following the directed edges. Note that some vertices make a component on their own as there is no way to reach them again in a cycle.

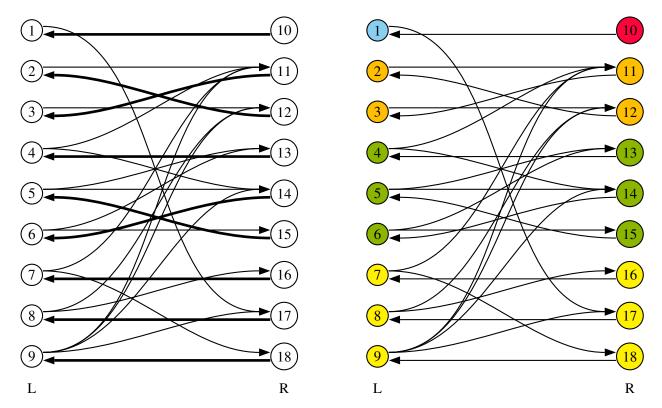


Fig. 2.8: Initial graph

Fig. 2.9: Maximum matching TODO RENAME

| node | index | low | stack | new component | color |
|------|-------|-----|---|-----------------------|-------|
| 1 | 1 | 1 | [1] | | |
| 17 | 2 | 2 | [1, 17] | | |
| 8 | 3 | 2 | [1, 17, 8] | | |
| 12 | 4 | 4 | [1, 17, 8, 12] | | |
| 2 | 5 | 4 | [1, 17, 8, 12, 2] | | |
| 11 | 6 | 4 | [1, 17, 8, 12, 2, 11] | | |
| 3 | 7 | 4 | [1, 17, 8, 12, 2, 11, 3] | | |
| | | | [1, 17, 8] | [3, 11, 2, 12] | |
| 16 | 8 | 2 | [1, 17, 8, 16] | | |
| 7 | 9 | 2 | [1, 17, 8, 16, 7] | | |
| 18 | 10 | 2 | [1, 17, 8, 16, 7, 18] | | |
| 9 | 11 | 2 | [1, 17, 8, 16, 7, 18, 9] | | |
| 14 | 12 | 12 | [1, 17, 8, 16, 7, 18, 9, 14] | | |
| 6 | 13 | 12 | [1, 17, 8, 16, 7, 18, 9, 14, 6] | | |
| 13 | 14 | 12 | [1, 17, 8, 16, 7, 18, 9, 14, 6, 13] | | |
| 4 | 15 | 12 | [1, 17, 8, 16, 7, 18, 9, 14, 6, 13, 4] | | |
| 15 | 16 | 12 | [1, 17, 8, 16, 7, 18, 9, 14, 6, 13, 4, 15] | | |
| 5 | 17 | 12 | [1, 17, 8, 16, 7, 18, 9, 14, 6, 13, 4, 15, 5] | | |
| | | | [1, 17, 8, 16, 7, 18, 9] | [5, 15, 4, 13, 6, 14] | |
| | | | [1] | [9, 18, 7, 16, 8, 17] | |
| | | | | [1] | |
| 10 | 18 | 18 | [10] | | |
| | | | | [10] | |

Table 2.1: Steps Tarjan's algorithm TODO

Algorithm 2: Tarjan's strongly connected components

```
1 void Tarjan(Graph G)
2 begin
        Global int index \leftarrow 0
3
        Global int components \leftarrow 0
 4
        Global int n \leftarrow |V(G)|
5
        Global S \leftarrow new \operatorname{Stack}()
 6
        Global stacked \leftarrow new boolean[n]
 7
        Global id \leftarrow new \operatorname{int}[n]
8
        Global low \leftarrow new int[n]
        for u \in V(G) do
10
            if \neg stacked[u] then dfs(u, G)
11
12 void dfs(int u, Graph G)
13 begin
14
       push(u, S)
        stacked[u] \leftarrow true
15
        low[u] \leftarrow index
16
        index \leftarrow index + 1
17
        int min \leftarrow low[u]
18
        for v \in N(u,G) do
19
            if \neg stacked[v] then dfs(v,G)
20
            min \leftarrow minimum(low[v], min)
21
        low[u] \leftarrow minimum(low[u], min)
22
        integer v
23
        repeat
24
            v \leftarrow pop(S)
25
            id[v] \leftarrow components
26
            low[v] \leftarrow n
27
        until v \neq u
28
        components \leftarrow components + 1
29
```

Tarjan's algorithm for finding strongly connected components is shown in algorithm 2. The Tarjan proceduce in line 1 takes a Graph G as an argument and represents the start of the algorithm. The global integer index declared in line 3 stores the current order in the depth-first search, and gets incremented each time a vertex is discovered. The integer components in line 4 keeps track of how many components were identified in the given graph. The integer n in line 5 represents the number of vertices in the given graph.

In line 6 we introduce a Stack data structure that will hold all the vertices that are part of the same strongly connected component. Line 7 contains a declaration for the stacked vertex-indexed array of booleans used to keep track if a vertex is present or not on the stack S.

An integer vertex-indexed array id is declared in line 8 to store the id of the strongly connected component of each vertex. Line 9 declares a vertex-indexed array low used to store the topmost reachable vertex in the depth-first search tree through a back edge. Lines 10 and 11 contain a for loop that calls a procedure for performing depth first search on every vertex in the given graph that is not currently stacked.

Line 12 declares a recursive procedure for performing a depth-first search starting from vertex u in graph G. The procedure starts by pushing the vertex u on the stack S (line 14) and then updating the value in the stacked array (line 15) to reflect the change. Line 16 sets the vertex's v value for low to index as every vertex assumes to have

no ancestors until a back edge to one is found. In line 17, the value for index is incremented by one to keep count of the current number of discovered vertices. Line 18 declares an integer min used to hold information about the oldest, lowest ancestor that will be discovered later in the search and is initialized to take the same value of the current vertex.

Lines 19 to 21 loop through every vertex v that is a neighbour of vertex u. The procedure N(u,G) on line 19 returns a list of neighbouring vertices of a given vertex, not including the given vertex. If the discovered neighbour v is not currently on the stack S, then we proceed to make a recursive call to the depth-first search function on the found vertex.

After the recursive call finishes processing the sub-tree of the neighbouring vertex v, min gets updated on line 21 to store the topmost reachable vertex that could be the same, or even higher in the tree if a higher back-edge from v was discovered.

Line 22 updates the topmost reachable vertex of vertex u to reflect any changes after processing all the neighbouring vertices.

Line 23 declares an integer v used to store a vertex that for the repeat-until loop in lines 24 to 28. The loop begins by popping vertices out of the stack s on line 25. Each vertex v found on the stack is assigned the current component id. Line 26 assigns the last leaf node of the graph as a topmost reachable vertex of vertex v in order to mark it as already part of a component. The repeat-until loop in line 24 to 28 is runs until reaching the root of the component.

The depth-first search procedure finishes on line 29 by updating the total number of strongly connected components discovered.

The final step in the all-different algorithm uses the knowledge gained after finding the strongly connected component in the graph using Tarjans algorithm. For this step, we are going to be interested only the edges that link two vertices in different strongly connected component. Figure 2.10 TODO shows a graph highlighting the edges we are interested in. It is important to remember that we chose a graph with 18 vertices to highlight how the all-different algorithm will run on the 9 variables of a Sudoku row. The variables of the Sudoku row are shown in the left L partition of the graph and are assigned letters from A-I, each having edges to vertices in the right R partition of the graph representing digits in the domain of the variable. The direction of the edges that are left indicate two actions. Forward edges (coloured in red) indicate that the digits from the right R partition should be removed from the domain of the variables in the L partition to achieve consistency TODO. Backward edges (coloured in blue) indicate finding the solution to a variable, in this case the first variable R of a selected Sudoku row will certainly take the value of 1 as all other options were removed from its domain.

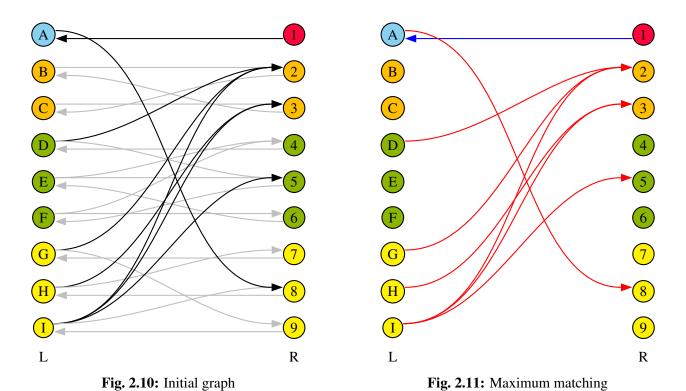


Fig. 2.12: Tarjan's algorithm next steps TODO: rename

AllDiff Demo

implementation and usage

Explain how demo is used, what it shows, etc (NOT implementation)

Implementation

The Graphical User Interface (GUI) is inside a JLayeredPane which is a special swing component that allows components inside it to overlap according to their specified depth. In my implementation, the backmost layer contains a JPanel responsible for the drawn edges in the graph. The layer above contains the rest of the GUI represented by the Sudoku puzzle on the left hand side of the screen, and the vertices of the graph on the right hand side of the screen. By using a layered panel, the edges of the graph can be drawn behind the vertices.

The Sudoku Grid present on the left hand side of the screen is made up of 9 by 9 Sudoku Cells positioned at specific locations. When a new Sudoku Cell instance in made, its ith row and jth column number are passed as arguments to the constructor. Using the information related to its location in the grid, it is now possible to render the border of each cell. Initially all the top, left, bottom and right borders have a width of 1px. To represent the 3 by 3 sub-grids, specific borders are set to have a width of 2px. In the end, the 9 by 9 grid is given an overall thick border by specifying a width of 4px to the cells that are in the first and last columns and rows. All Sudoku Cells contain a JLabel with text retrieved from the Model denoting which possible values are still available for that particular cell.

Not visible to the human eye, a new Sudoku grid instance is drawn on top of the existing one, this time with all borders set to a width of 1px. The Sudoku Cells inside this second grid contain the same information about the state of the puzzle retrieved from the Model. Additional information such as a custom background will be applied to these cells as the algorithm progresses. Cells inside the second grid become visible once the user makes a selection. Once visible, the cells inside the selection are animated into their position inside the graph on the right hand side of the screen. When the user performs deselects his choice, the nice cells are animated back into their original position in the grid and become invisible again.

Particular attention was given to make the software thread safe. It is crucial that the application logic doesnt make the Graphical User Interface unresponsive. To achieve this, the program will run on two threads, one main thread responsible with all the processing need to be done by the program, and the Event Dispatching Thread.

The Event Dispatching Thread (EDT) is a background thread used to invoke Swing methods that change the GUI and listen to events associated to components. Most Swing components are not thread safe, therefore bugs such as race conditions could unexpectedly arise if it was to perform the actions from normal threads. The EDT acts as a queue of events that are performed sequentially.

Throughout the application, Timer objects are used to schedule repeated actions such as the fading out of the components or movement of the cells. The model contains a two-dimensional ArrayList of Timer objects that acts as a queue of animations. Timers on the same level start running at the same time until they all finish. Once all the timers on a level finish, the next level in the two-dimensional ArrayList is started, allowing us to queue a movement of a row after all the cells finish fading out.

Conclusion and Future Work

We have shown how to implement the all-different constraint.

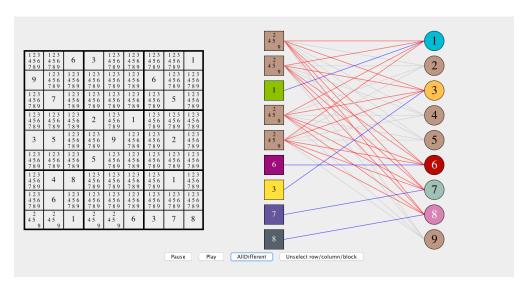


Fig. 5.1: The state of the program after the all-different algorithm finishes on a row, in this case, the last one.

Move this from here

Appendices

Appendix A

Running the Program

An example of running from the command line is as follows:

```
> javac *.java
> java Sudoku
```

This will open the application loaded with the hard Sudoku problem /herald20061222H.txt.

TODO: what about the Choco3 library? add it to path? remove it?

Appendix B

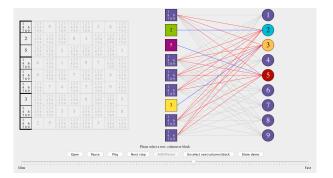
Proof of concept

The following sequence of steps will provide a visual proof of the all-different algorithm. The following figures capture the user running the all-different algorithm on 5 predetermined rows, columns or 3x3 sub-grids. For demonstration purposes, the Sudoku instance used is *lockedset.txt*.

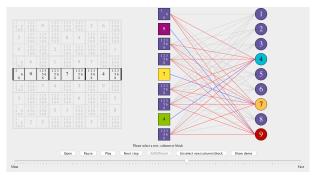
The sequence is as follows:

```
> Open lockedset.txt Sudoku instance
> Select the 1st column
> Run the all-different algorithm
> Unselect
> Select the 5th row
> Run the all-different algorithm
> Unselect
> Select the 6th row
> Run the all-different algorithm
> Unselect
> Select the 4th 3x3 sub-grid
> Run the all-different algorithm
> Unselect
> Select the 1st column
> Run the all-different algorithm
> Unselect
```

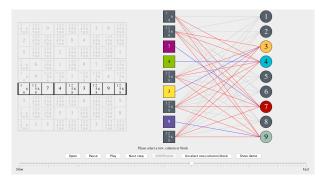
Alternatively, the user can press the *ShowDemo* button to run the same steps.



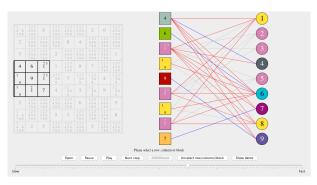




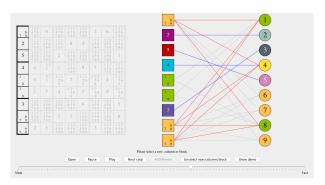
(b) 5th row after running the all-different algorithm



(c) 6th row after running the all-different algorithm



(d) 4th 3x3 sub-grid after running the all-different algorithm



(e) 1st column after running the all-different algorithm

Fig. B.1: Steps in the demo

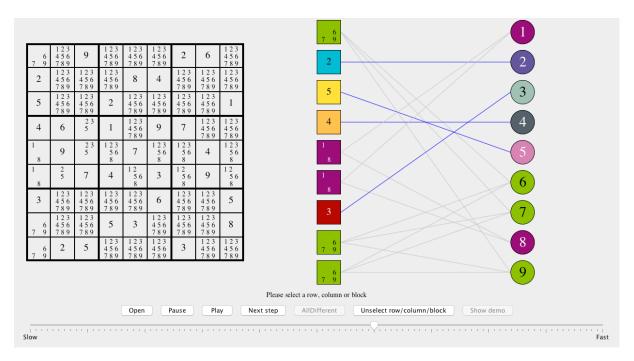


Fig. B.2: State of the program after running the demo

As seen in Fig. B.1, the all-different algorithm performs changes only to the selected 9 cells inside a row, column or 3x3 sub-grid.

The end result of following the above steps can be seen in Fig. B.2. This state mimics the human thinking when solving a Sudoku. In the 1st column there are two cells with a domain of 1, 8, but we don't know yet which cell will take which digit. What we do know though, is that the digits 1, 8 will be distributed in that two particular cells, therefore in the last step, after running the all-different algorithm on the 1st column we can observe that the digits 1, 8 disappear from the domains of the rest of the cells in the selection.

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