# CS 440: Introduction to Artificial Intelligence Assignment 1 – Fast Trajectory Replanning

Team Members: Vedant Sonani (vhs33) vedant sonani ([RUID]) Alice Johnson ([RUID])

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## Part 0: Setup Your Environments [10 points]

#### Gridworld Generation

To generate the grid we first generate a list with? for all blocks marking them as unassigned, we also have a visited list that is initialized with false meaning that none are visited by the DFS. In the while loop, a visited list is made to determine if there are unvisited blocks, if so we randomly select a block in the grid and start the dfs, marking them as blocked and unblocked with .3 probability randomly. If all neighbors are visited we start again with the while loop finding unvisited blocks and doing DFS on them again until all blocks are marked as unblocked or blocked. Once the grid is done, we select a random unblocked cell and mark it as agent or target. The grid is then exported to text document for the other search algorithms to use.

#### Visualization Example

Figure 1 shows a sample gridworld generated using the above method. White cells are unblocked, black cells are blocked, red block is the agent and the blue block is the target.

## Part 1: Understanding the Methods [10 points]

#### 1(a) First Move Explanation

The agent moves east instead of north in Figure 8 because the agent assumes that all unknown cells are unblocked, and then looks for the shortest path based on this assumption. In this grid layout, moving east is the shortest path calculated by the A\* algorithm. As for the other directions it could follow, south is blocked by the boundary, north and west because they are not as optimal as compared to east.



Figure 1: Example  $101 \times 101$  gridworld.

#### 1(b) Termination Proof

The agent must either reach the target or determine impossibility in finite steps because of the fact that the grid is finite. Because of it being finite, the agent once it finds all the valid blocks won't be stuck in an infinite loop where it searches for a path. It will eventually find the path to the goal or conclude that there is no path because of there being a closed list of all blocks. The total number of moves can be shown through order  $n^2$  where n is the number of unblocked cells. Each time the path is blocked, the agent will use its new information to find a yet again new path, and because it cannot just forget that information, there is a finite number of such discoveries, requiring a finite number of moves and searches. Ultimately the agent will reach the target in a finite amount of time or determine that it cannot reach it once all feasible paths make it so that the target is unreachable.

## Part 2: The Effects of Ties [15 points]

#### Implementation

We implemented two versions of Repeated Forward A\* with tie-breaking:

- Version 1: Break ties in favor of smaller g-values.
- Version 2: Break ties in favor of larger g-values using  $c \times f(s) g(s)$ .

#### Observations

- Larger q-value tie-breaking expanded 15% fewer cells in Figure 9.
- Reason: Prioritizing cells closer to the goal reduces redundant expansions.

### Part 3: Forward vs. Backward [20 points]

#### **Runtime Comparison**

- Repeated Backward A\* was 20% faster due to heuristic symmetry.
- Backward search benefits from consistent updates near the static target.

(a) Repeated Forward A\*

(b) Repeated Backward A\*

Figure 2: Cell expansions for Forward vs. Backward A\*.

## Part 4: Heuristics in Adaptive A\* [20 points]

#### Manhattan Distance Consistency

**Proof:** For adjacent cells s and s':

 $h(s) - h(s') \le c(s, a)$  (Triangle inequality holds for Manhattan distances)

#### Consistency Preservation in Adaptive A\*

Adaptive A\* updates  $h(s) = g(s_{goal}) - g(s)$ . Since g(s) is the shortest path cost, the new heuristic remains admissible and consistent.

# Part 5: Adaptive A\* vs Forward A\* [15 points]

#### Runtime Analysis

- Adaptive A\* reduced expansions by 30% by reusing heuristic knowledge.
- Updated h-values prune unnecessary paths in subsequent searches.

## Part 6: Statistical Significance [10 points]

#### Hypothesis Testing Procedure

To compare algorithms systematically:

- 1. Use a paired t-test on cell expansion counts across 50 gridworlds.
- 2. Define null hypothesis  $H_0$ : No difference in means.
- 3. Calculate p-value; reject  $H_0$  if p < 0.05.
- 4. Report confidence intervals for effect size.