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# ***Tumor Detection in Brain MRI Images Using Image Segmentation and Flood Fill Algorithm (DFS)***

## **1. Algorithm Integration**

For this project, the primary algorithm implemented is **Flood Fill using Depth-First Search (DFS)**, designed to detect and isolate tumor regions in grayscale MRI brain images. The steps include:

- **Preprocessing:** Images are first converted to grayscale and thresholded to enhance contrast between normal and abnormal tissue.
- **Flood Fill with DFS:** The algorithm begins at a seed pixel, checking connected pixels (typically 4- or 8-directionally) for similarity in intensity values. DFS is applied using an explicit stack data structure (`std::stack`) rather than recursion, to give better control and prevent stack overflow on large regions.
- **Region Marking:** Tumor-like areas are colored or highlighted for visualization.

This approach allows precise traversal and segmentation of irregular regions, which is common in tumor boundaries.

## **2. Optimization & Efficiency Analysis**

To analyze performance and optimize, the following comparisons and observations were made:

- **DFS vs BFS:** DFS was chosen for its lower memory footprint in deep but narrow regions, which is common in organic, tumor-like patterns. BFS, while effective, can become memory-intensive as it stores a wider frontier at each level.
- **Stack Implementation:** Using a manual stack avoids system recursion limits, improving robustness and runtime performance in large images.

- **Threshold Tuning:** Adjustable intensity threshold values were tested to balance under-segmentation (missing parts of the tumor) and over-segmentation (including non-tumor tissue).
- **Time Complexity:** Approx.  $O(n)$ , where  $n$  is the number of pixels in the segmented region.
- **Space Complexity:**  $O(n)$  for visited pixels and the explicit stack.

Further experiments will include performance profiling on MRI images of different sizes and resolutions.

### 3. Edge Case Handling

Multiple edge cases have been tested or planned for testing:

- **Multiple Tumors:** The algorithm can detect multiple disconnected regions by applying DFS from multiple seed points.
- **Tumor at the Image Border:** Ensures the algorithm correctly detects regions even if they extend to image edges.
- **No Tumor Present:** Handled by setting a minimum region size; false positives are discarded.
- **Highly Noisy Images:** Noise filtering (e.g., Gaussian blur) is optionally applied before segmentation to improve accuracy.

Further enhancements are planned to automate seed point selection using histogram analysis or contours.

### 4. Implementation of Course Concepts

This project is strongly rooted in the **Data Structures and Algorithms (DSA)** concepts discussed in class:

- **Stacks:** Manual implementation of DFS using the stack data structure for iterative traversal.
- **Vectors:** Use of `std::vector` for dynamic storage of connected components and pixel positions.

- **Graph Traversal (DFS):** The MRI image is treated like a 2D grid graph where each pixel is a node and adjacency is defined by pixel connectivity.
- **Algorithm Design:** Custom thresholding and control flow to adapt DFS to a flood fill scenario.

The project directly applies classroom knowledge of graph traversal, memory management, and complexity analysis in a real-world application.

## Current Status and Next Steps

- **Completed:** Basic flood fill implementation using DFS, test cases for various input images, manual seed point detection.
- **In Progress:** Automatic seed selection, refining thresholding, benchmarking with other segmentation techniques (e.g., region growing, contour-based).
- **Next:** Implement GUI-based interaction using OpenCV, finalize performance comparison charts, write final report and documentation.