# Final

### April 22, 2025

## 0.1 Merge Sort

Merge Sort is an efficient, general-purpose, and comparison-based divide-and-conquer sorting algorithm. It's efficient and general purpose in that in it's worst case, it runs in  $O(n \log n)$  time, on par with the best of the classic sorting algorithms; comparison-based in that elements are directly compared to insert into the sorted array; and divide-and-conquer in that algorithm recursively sorts subarrays, effectively breaking up the problem into subproblems until a base case is reached.

The divide-and-conquer nature of this algorithm makes it a suitable candidate for parallel processing, further improving on the efficiency of such an algorithm. The default sorting algorithm for Python, TimSort, actually uses a hybrid of merge sort and insertion sort. Beyond sorting, Merge Sort has applications in polynomial multiplication, genomic sequencing, and comparative ranking systems.

## 0.2 Implementation

```
[1]: import numpy as np
     # General merge sort algorithm, recursively calls itself until the base case is
      ⇒reached.
     def mergeSort(array):
      n = len(array)
       # base case of only one element in the array:
       if n \le 1:
         return array
       # setting the midpoint:
       if n\%2 == 0:
           mid = int(n/2)
       else:
           mid = int(n/2)+1
       # splitting the array down the middle and recursively sorting the halves:
       left = mergeSort(array[0:mid])
       right = mergeSort(array[mid:n])
       # merging and returning the sorted halves of the array:
       return merge(left, right)
```

```
# Helper function for merging two already sorted halves of the array:
def merge(left, right):
  # initializing the resulting sorted array and the indices of interest:
 result = []
 i = 0
 j = 0
  # looping through the elements of both arrays to compare and place in sortedu
 ⇔array:
 while i < len(left) and j < len(right):</pre>
    if left[i] <= right[j]:</pre>
      result.append(left[i])
      i += 1
    else:
      result.append(right[j])
      j += 1
  \# the remaining elements can be assumed to be greater than all the elements \sqcup
 already placed in the array, and will only come from one half of the array.
  # looping through the remaining elements to clean up and add to sorted array:
  while i < len(left):
    result.append(left[i])
    i += 1
 while j < len(right):</pre>
    result.append(right[j])
    j += 1
  # resulting array is fully sorted.
  return result
```

```
Unsorted: [2, 7, 6, 3, 3, 7, 3, 8, 8, 9]
Sorted: [2, 3, 3, 3, 6, 7, 7, 8, 8, 9]

Unsorted: [7, 2, 1, 8, 3, 4, 8, 2, 8, 6, 6, 7, 5, 2, 5, 3, 3, 9, 1, 2]
Sorted: [1, 1, 2, 2, 2, 2, 3, 3, 3, 4, 5, 5, 6, 6, 7, 7, 8, 8, 8, 9]

Unsorted: [3, 8, 7, 1, 2, 4, 5, 9, 4, 4, 6, 9, 8, 2, 1, 4, 9, 6, 4, 7, 1, 1, 5, 4, 8, 8, 7, 4, 7, 5, 8, 6, 1, 1, 1, 2, 4, 6, 4, 7]
Sorted: [1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 3, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 5, 5, 5, 6, 6, 6, 6, 7, 7, 7, 7, 8, 8, 8, 8, 8, 9, 9, 9]
```

### 0.3 Experimental Analysis

- Measure the running time of the algorithm on different input sizes and compare it to its theoretical time complexity.
- Measure the peak memory usage of the algorithm and analyze how memory requirements scale with input size.
- Prepare visualizations, such as graphs or tables, to illustrate your experimental results.

```
[35]: import time
      import matplotlib.pyplot as plt
      import tracemalloc
      # Measures the peak memory allocation and run time for the given input size:
      def analysis(input_size):
          # Generating the test case with the given input size
          ex = np.random.randint(1, 100, int(input_size))
          # Setting a timer
          start_time = time.time()
          # Tracking memory allocation
          tracemalloc.start()
          # Calling the algorithm with the test case
          mergeSort(ex)
          # Extracting insights from memory allocation tracker
          current, peak = tracemalloc.get traced memory()
          tracemalloc.stop()
          # Return the peak memory allocation and run time of test case
          return peak / 10**3, time.time()-start_time # return in MB
      # Initiate arrays for storing run times and memory usage, as well as input \Box
       ⇔sizes for analysis
```

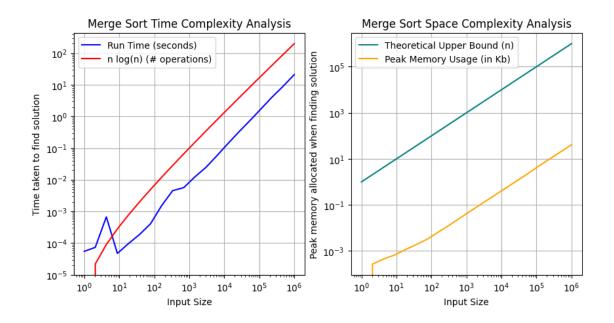
```
times = []
memory_usages = []
input_sizes = np.logspace(0, 6, 20)

# Looping through input sizes to measure each one
for input_size in input_sizes:
   mem, time_elapsed = peak_memory(input_size)
   times.append(time_elapsed)
   memory_usages.append(mem)
```

```
[34]: fig, (ax1, ax2) = plt.subplots(1, 2)
      fig.set_figwidth(10)
      # Plotting running time against theoretical upper bound
      ax1.loglog(input_sizes, times, 'b-', label='Run Time (seconds)')
      nlogn = np.log2(input_sizes) * input_sizes * 10**-5
      ax1.loglog(input_sizes, nlogn, label='n log(n) (# operations)', color='red')
      # Plotting memory usage against thoeretical upper bound
      ax2.loglog(input_sizes, input_sizes, color='teal', label='Theoretical Upper_

→Bound (n)')
      ax2.loglog(input_sizes, memory_usages, label='Peak Memory Usage (in Kb)', u
       ⇔color='orange')
      # Add labels and title for first plot
      ax1.set_xlabel("Input Size")
      ax1.set_ylabel("Time taken to find solution")
      ax1.set_title("Merge Sort Time Complexity Analysis")
      ax1.grid(True)
      ax1.legend()
      # Add labels and title for second plot
      ax2.set_xlabel("Input Size")
      ax2.set_ylabel("Peak memory allocated when finding solution")
      ax2.set_title("Merge Sort Space Complexity Analysis")
      ax2.grid(True)
      ax2.legend()
```

[34]: <matplotlib.legend.Legend at 0x11efdfe00>



## 0.4 Key Takeaways

From this project we learned that merge sort is a very useful and reliable sorting algorithm. The divide-and-conquer structure gurantees that the time complexity will always be  $O(n \log n)$  no matter what input it takes. Merge sort is very useful for very large datasets that need to be sorted. This algorithm does have some trade offs such as the tempory space it uses to sort the subproblems or its recursive nature. However, its guranteed time complexity and ease of being scaled for larger and larger inputs makes it an extremly useful and powerful tool to sort data.

#### 0.5 Deliverables

- Source Code: A well-documented implementation of the algorithm, including a README file describing how to run the code and reproduce results. 2
- Report: A 2–3 page report summarizing the algorithm, its implementation, experimental analysis, and key takeaways.
- Presentation Slides: A professional slide deck to accompany your presentation. This project is an opportunity for you to apply theoretical knowledge, practice your coding skills, and analyze the performance of algorithms in real-world scenarios. It's your chance to dive into the fascinating world of algorithms and showcase your creativity and problem-solving skills. We look forward to seeing your innovations!