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Theoretical Run-time Analysis

Algorithm 1: Enumeration

```
cubicMSS(A[0...n-1], n) {
       maxSum = 0;
       sum = 0;
       for (i to n) { // starting index
               for(j = i to n) \{ // ending index \}
                       sum = 0;
                       for (k = i \text{ to } k \le j) \{ // \text{ compute sum from } i, j \}
                              sum += A[k]
                              If (sum > maxSum) {
                                      maxSum = sum;
                                      startIndex = i;
                                                             // keep this startIndex
                                      endIndex = j;
                                                             // keep this endIndex
                              }
                       }
               }
       }
       subArrSize = (endIndex - startIndex) + 1; // calculate sub array size
        maxSubArr[subArrSize];
                                             // make new array of appropriate size
       for (i to subArrSize) {
                                              // fill new sub array
               maxSubArr[i] = A[startIndex];
               increment startIndex;
       }
       return maxSum, maxSubArr;
                                             // return max subarray sum and subarray
}
```

This algorithm uses brute-force, trying every possible i, j pair and computing the sum of entries in A[i...j]. There are $O(n^2)$ pairs and it takes O(n) time to compute each sum of pairs.

```
O(n^2) * O(n) = O(n^3) run time
```

Algorithm 2: Better Enumeration

```
quadMSS(A[0...n-1], n) {
       maxSum = 0;
       sum = 0;
       for (i to n) {
                                     //O(n) iterations for outer loop (n possible start positions)
               sum = 0
              for (for i = i to n) {
                                     //O(n) iterations for inner loop (n possible end positions)
                      sum += A[i]; //O(1) constant time to update sum
                      if (sum > maxSum) {
                              maxSum = sum;
                              startIndex = i;
                                                    // keep this startIndex
                              endIndex = j;
                                                    // keep this endIndex
                      }
              }
       }
       subArrSize = (endIndex - startIndex) + 1; // calculate sub array size
       maxSubArr[subArrSize];
                                            // make new array of appropriate size
       for (i to subArrSize) {
                                            // fill new sub array
               maxSubArr[i] = A[startIndex];
               increment startIndex;
       }
       return maxSum, maxSubArr;
                                            // return max subarray sum and subarray
}
```

This algorithm also uses brute-force using iteration to go through every possible subarray until the max sum is found. Similar to algorithm 1, we have an outer for loop which loops through n possible start positions taking O(n) time and an inner for loop which loops through n possible end positions, also taking O(n) time. However rather than computing the sum from scratch every time, we keep the previous sum from A[i] to A[j-1] and just add the extra A[j] to that previous sum, eliminating the need for the third for loop as we saw in algorithm 1 and making the time to compute the sum a constant O(1) rather than O(n).

 $(O(n) \text{ outer for loop}) * (O(n) \text{ inner for loop}) * (O(1) \text{ updating sum}) = O(n^2) \text{ run time}$

```
findMaxSubArrCrossingMid(A[1...n-1], left, mid, right) {
        leftSum = - infinity;
        sum = 0:
        // find max subarray of the left half
        for (i = mid to left) { // mid down to left (decreasing indices)
                sum = sum + A[i];
                if (sum > leftSum) {
                         leftSum = sum;
                                         // starting index of max subarray
                         maxLeft = i:
                }
        rightSum = - infinity;
        sum = 0;
        // find max subarray of the right half
        for (j = mid + 1 to right) { // mid up to high (increasing indicies)
                sum = sum + A[j];
                if (sum > rightSum) {
                         rightSum = sum;
                         maxRight = j; // ending index of max subarray
                }
        }
        return(maxLeft, maxRight, leftSum + rightSum);
}
findMaxSubArr(A[0...n-1], left, right) {
        if (right == left) { //base case (only one element)
                return (left, right, A[left]);
        }
        else mid = (left + right) / 2; // calculate middle
        // return max subarray (recursive calls) in one of the following:
                // a) left half
                (minLeft, maxLeft, leftSum) = findMaxSubArr(A, left, mid);
                // b) right half
                (minRight, maxRight, rightSum) = findMaxSubArr(A, mid+1, right);
                // c) subarray that crosses the midpoint
                (crossLeft, crossRight, crossSum) = findMaxSubArrCrossingMid(A, left, mid, right);
                // leftSum is the mss
                if (leftSum >= rightSum and leftSum >= crossSum)
                         return (minLeft, maxLeft, leftSum);
                // rightSum is the mss
                else if (rightSum >= leftSum and rightSum >= crossSum)
                         return (minRight, maxRight, rightSum);
                // crossSum is the mss
                else
                         return (crossLeft, crossRight, crossSum);
        }
```

The above algorithm also uses the divide and conquer concept using the observation that (1) the max subarray can only be contained in the first half, (2) only contained in the second half, or (3) composed of a suffix of the first half and a prefix of the second half. (1) and (2) are solved by recursion. Therefore, we are repeatedly dividing the length of our array in half every time we go down a level in our tree until we finally reach the base case (an array containing a single element), giving us a depth of O(lg n). The non-recursive work for (3) is done in linear O(n) time because we are simply finding the max suffix of the first half and the max prefix of the second half and combining them together.

(O(n) non-recursive work) * (O(lg n) tree depth) = O(n lg n) run time

Algorithm 4: Linear-time

```
linearMSS(A[0...n-1], n) {
       maxSoFar = 0;
       maxEndingHere = 0;
       startIndex, endIndex, s = 0
       for(i to n) {
              maxEndingHere += A[i];
              if(maxSoFar < maxEndingHere) {</pre>
                      maxSoFar = maxEndingHere;
                      startIndex = s;
                      endIndex = i;
              if(maxEndingHere < 0) {</pre>
                      maxEndingHere = 0;
                      s = i + 1;
              }
       subArrSize = (endIndex - startIndex) + 1; // calculate sub array size
       maxSubArr[subArrSize];
                                            // make new array of appropriate size
       for (i to subArrSize) {
                                            // fill new sub array
              maxSubArr[i] = A[startIndex];
              increment startIndex;
       }
       return maxSoFar, maxSubArr;
                                            // return max subarray sum and subarray
}
```

The above algorithm uses dynamic programming by using a bottom-up approach to find our max sum subarray. This means we start from our smallest problem (1 element) and work our way up to the biggest problem (the entire array). The max subarray we are looking for either uses the last element in our subset or it doesn't. The work of finding our max subarray and max suffix takes constant O(1) time. There are O(n) elements to compute in our array, therefore our running time is O(n).

(O(n)) elements to compute) * (O(1)) compute max array and max suffix) = O(n) run time

Testing

Given that we are to assume all elements in the input arrays will be integers with at least one positive element in each array, we used the following guidelines for test sets against our program:

Test Input	Example
Array of size 1, positive	[5]
Array of all negative values with one positive value	[-1-10-9-2-82]
Array of all positive values	[7361210002]
Array of even size, mixed pos/neg values	[-100 0 -12 20 -50 30 5 -7 9 100]
Array of odd size, mixed pos/neg values	[3 8 -25 150 -10 0 1 2 -40]
Array containing all pos values in the first half and all negative values in the second half	[11111-1-1-1-1]
Array containing all neg values in the first half and all pos values in the second half	[-100 -100 -100 -100 100 100 100]

We also tested our algorithms against the MSS_Problems.txt and MSS_TestProblems.txt files. While MSS_Problems.txt produced the expected results, MSS_TestProblems.txt contained a bug where it would sometimes duplicate the last number of our subset array. After some investigation, we determined that a few lines in MSS_TestProblems.txt contained a trailing space at the end, causing the duplicate number to show up in our results. The MSS_Problems.txt file did not contain these trailing spaces. After removing the trailing spaces from the MSS_TestProblems.txt file our code produced the correct results. Since we are using MSS_Problems as the file we are submitting with our code, we used it's format as a "standard" and therefore did not leave trailing spaces at the end of the lines of input when testing our code.

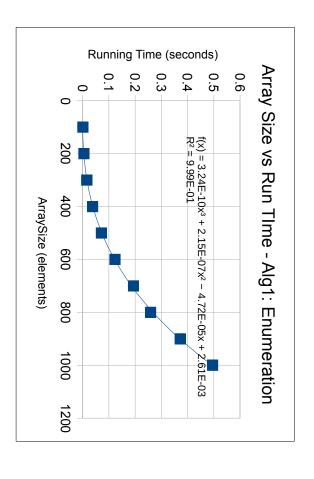
Conclusion

For the problem where we are given an array of small integers and are to compute the max sum and corresponding subarray, we designed, implemented, and analyzed four algorithms to solve this problem. From this analysis, we have come to the conclusion that algorithm 4, using dynamic programming and a bottom-up approach was the most efficient. This means that algorithm design does indeed matter- especially when dealing with large data sets. To illustrate the difference we just have to look at our experimental analysis to see that algorithm 1 can only handle an input of 5,490 in 60 seconds compared to algorithm 4 which handles 1.41E+10 in the same amount of time! In addition, since algorithm 1 follows a cubic curve, it's efficiency will taper as our data set grows larger, but algorithm 4 follows a linear curve, so it's efficiency will remain linear to the size of the input.

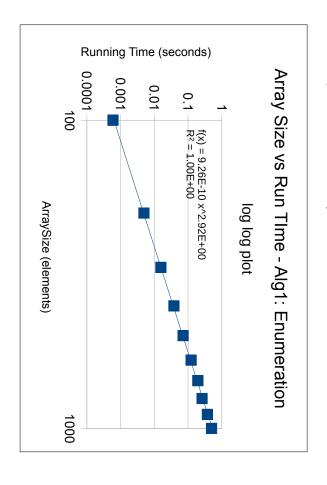
	Theoretical RT	Curve fit to data	R^2	Largest input in 60 secs
Algorithm 1	O(n³)	Cubic	.999	5,490
Algorithm 2	O(n ²)	Quadratic	.995	229,000
Algorithm 3	O(nlgn)	nlgn/linear	.980	1.35E+09
Algorithm 4	O(n)	linear	.995	1.41E+10

0.4951969	1000
0.3719375	900
0.2591869	800
0.1936407	700
0.1233996	600
0.0715286	500
0.0381729	400
0.0155972	300
0.0049087	200
0.0005969	100
Avg run time (s)	n
ation	Alg1 : Enumeration

be solved in a given time, we set the regression equal to the given times:	To predict the largest input (array size) that could
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which is predicted theoretically. The regression curves show a polynomial (cubic) relationship,

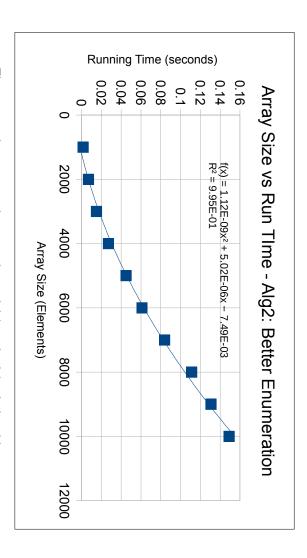


0.1490488	10000
0.1307949	9000
0.1112906	8000
0.0839139	7000
0.0610665	6000
0.0450681	5000
0.0271719	4000
0.0151424	3000
0.0070292	2000
0.001633	1000
Avg run time (s)	n
numeration	Alg2: Better Enumeration

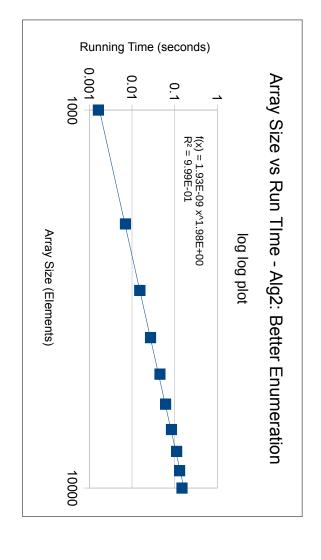
10000	9000	8000	7000	6000	5000	
0.1490488	0.1307949	0.1112906	0.0839139	0.0610665	0.0450681	

To predict the largest input (array size) that could be solved in a given time, we set the regression equation equal to the given times:

229000	60
92000	10
65000	5
Array Size (n)	Time (s)
quation equal to the given times:	quation equal
e solved in a given time, we set the regre	e solved in a

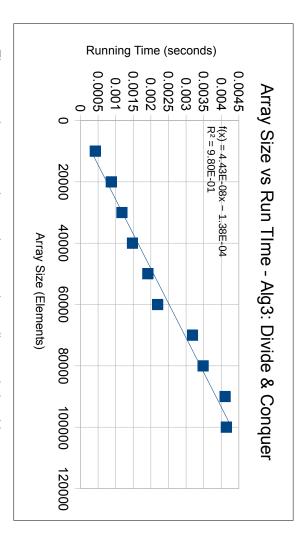


which is predicted theoretically. The regression curves show a polynomial (quadratic) relationship,

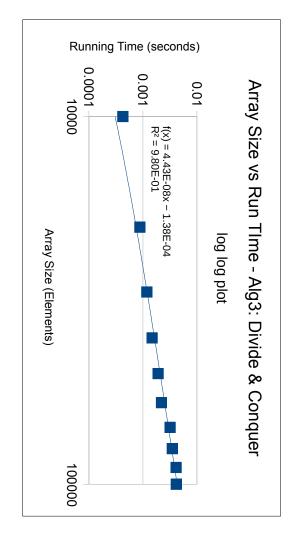


0.0041421	100000
0.0041069	90000
0.003483	80000
0.0031834	70000
0.0021938	60000
0.0019137	50000
0.0014758	40000
0.0011774	30000
0.0008745	20000
0.0004214	10000
Avg run time (s)	n
nd Conquer	Alg3: Divide and Conquer

1 35E+09	60
2.26E+08	10
1.13E+08	ഗ
Array Size (n)	Time (s)
equation equal to the given times:	equation equal



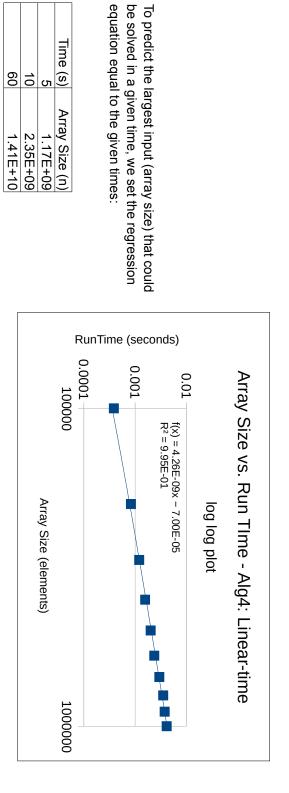
The regression curves show a close match to a linear relationship. Theoretical predictions indicate a nlgn relationship. However, the variation due to lgn is not significant compared to n, so this graph appears linear.



0.0041141	1000000
0.0037737	900000
0.0035241	800000
0.0029723	700000
0.0023643	600000
0.0020156	500000
0.0015683	400000
0.0012008	300000
0.0008302	200000
0.0003848	100000
Avg run time (s)	n
ne	Alg4: Linear-time

0 0 +	RunTime 0.00 1	(seconds)	Array 0.005
200000 400000 600000 800000 1000000 1200000 Array Size (elements)		f(x) = 4.26E-09x - 7.00E-05 R ² = 9.95E-01	Array Size vs. Run Tlme - Alg4: Linear-time

predicted theoretically. The regression curves show a linear relationship, which is



Time (s) 60 5 Array Size (n) 1.17E+09 2.35E+09 1.41E+10

equation equal to the given times:

