# MAXIMUM SUM SUBSEOUENCE 

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## WHAT DOES THE PROBLEM SAY?

- Determine a subsequence of data set that sums to the maximum value with respect to any other subsequence of the data set.
- More formally, if we are given a sequence $X=\left\langle x_{0}, x_{1}, x_{2}, \ldots, x_{n}\right\rangle$ we are required to find a set of indices $u$ and $v, u \leq v$, such that the subsequence $\left\langle x_{u}, x_{u+1}, \ldots, x_{v}\right\rangle$ has the largest possible sum, $x_{u}+x_{u+1}+\ldots+x_{v}$, among all subsequences of $X$.
- This problem is non-trivial only if the given sequence has both positive and negative values


## IS THAT EVEN A THING?

- Protein and DNA sequence analysis
- To locate biologically meaningful segments, e.g. conserved segments , GC-rich regions, non-coding RNA genes, and transmembrane segments.
- A common approach is to assign a value to each residue, and then look for consecutive subsequences with high sum or average.
- Traffic Monitoring
- Example: Say for a bridge, given the numbers of vehicles entering and exiting at various passes, you can determine the busiest routes.


## ARCHITECTURE

- We worked this problem on a Linear Array
- But, we don't have a Linear Array!
- But we always can simulate the same behavior virtually.
- We wrote the MPI code such that the program behaves as if it were running on a Linear Array.
- It comes with a cost though.
- The communication time is more than what is intuitive.


## ALGORITHM

- The algorithm that I am considering to implement is from "Algorithms Sequential \& Parallel: A Unified Approach" by Russ Miller and Laurence Boxer.
- First compute the parallel prefix sums $S=\left\{p_{0}, p_{1}, \ldots, p_{n-1}\right\}$ of $X=\left\{x_{0}, x_{11}, \ldots, x_{n-1}\right\}$, where $p_{i}=x_{0} \otimes \ldots \otimes x_{i}$.
- Next, compute the parallel postfix maximum of $S$ so that for each index $i$, the maximum $\mathrm{pj}, \mathrm{j} \geq \mathrm{i}$, is determined, along with the value j .
- Let $m_{i}$ denote the value of the postfix-max at position $i$, and let $a_{i}$ be the associated index, i.e., $\mathrm{p}_{\mathrm{ai}}=\max \left\{\mathrm{p}_{i,} \mathrm{p}_{\mathrm{i}+1}, \ldots, \mathrm{p}_{\mathrm{n}-1}\right\}$.


## ALGORITHM CONTINUED

- Next, for each $i$, compute $b_{i}=m_{i}-p_{i}+x_{i}$, the maximum prefix value of anything to the right minus the prefix sum plus the current value.
- Finally, the solution corresponds to the maximum of the $b_{i}$ 's, where $u$ is the index of the position where the maximum of the $b_{i}^{\prime}$ 's is found and $v=a_{u}$.


## TIME COMPLEXITY

- This algorithm runs in $\Theta(n)$ time on a Linear Array.
- And the optimal cost of $\Theta(n)$ is achieved with $n^{1 / 2}$ processors.


## RUNNING TIME

| ProcesslValues | 1000 | 10000 | 100000 | 1000000 |
| :--- | :--- | :--- | :--- | :--- |
| $4(4 \times 1)$ | 0.0142 | 0.0832 | 1.1636 | 12.5648 |
| $8(8 \times 1)$ | 0.0271 | 0.1795 | 0.7540 |  |
| $16(8 \times 2)$ | 0.0301 | 0.1571 | 2.8025 | 19.2160 |
| $32(8 \times 4)$ | 0.0306 | 0.5545 | 6.7950 | 27.9722 |

## RUNNING TIME

Process vs Values


## RUNNING TIME

Process vs Value


## CONCLUSION

- Linear Array is not a very efficient architecture to go with.
- With the increase in no of processes the running time increases too because of the communication diameter.
- After a while the communication time completely overshadows the execution time.


## FUTURE GOALS

- For the rest of the semester, we would like to conduct experiments with even larger data.
- In the next phase we plan to work this out on various other architectures as well.
- A comparative analysis would be great!


## ACKNOWLEDGEMENT AND REFERENCES

- "Algorithms Sequential \& Parallel: A Unified Approach" by Russ Miller and Laurence Boxer
- http://wordaligned.org/articles/the-maximum-subsequence-problem
- "Genomic Sequence Analysis: A Case Study in Constrained Heaviest Segments" Kun-Mao Chao


## LET'S TALK!

- Session open for discussion

Thank you

