

# Scalability of Hybrid Sparse Matrix Dense Vector (SpMV) Multiplication

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**Abstract**—TBD

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## I. INTRODUCTION

## II. RELATED WORK

Due to the computation impact that SpMV operations have on a many scientific applications there has been an effort to analyze its performance and scalability characteristics. Bylina, Bylina, Stpiczunski, and Szalkowski [1] introduced and evaluated the performance of both multicore and multinodal implementations of SpMV on various chip architectures. A modified version of the SpMV algorithm found in the SPARSKIT Fortran library [2] for the multicore implementation. Using matrices from the University of Florida Sparse Matrix Collection (UFSMC), they found that for their multicore algorithm, similar performance was experienced across all matrices tested when the number of threads remained low. Alternatively as architectures allow for increased thread count, higher performance can be obtained, and it was noted that the use of OpenMP allowed for performance comparable to that of the optimized Intel MKL version of SpMV [cite Intel MKL ?](#).

Bylina et al's multinodal implementation distributed equal sized sub matrices of a given benchmark matrix to each MPI process, where in each process would then work on the non-zeros contained within that submatrix via a multithreaded version of Intel's MKL SpMV routines. For the distribution method chosen the density in addition to distribution of non-zeros within the matrix had the greatest impact on the scalability of their multinodal algorithm.

[-discuss memory bandwidth as it impacts SpMV](#)

## III. IMPLEMENTATION

### A. Work Distribution and MPI

The application that was written to emulate the behavior of that used in the study performed by Bylina et al was written using the C++ programming language. We chose to forgo the use of proprietary libraries such as BLACS and MKL deciding instead to write explicit MPI and OpenMP directives to control distributed and shared memory behavior across the cluster environment. Special care was paid to insure that the communication pattern matched that of the

2d cluster methods in the BLACS library which were used in the prior work. It was felt that by not including these packages greater control over communication and memory access parameters could be achieved, even though the result may not be as highly optimized for particular applications, architectures, or compilers.

Benchmark matrices in the Matrix Market Format were chosen from the University of Florida Sparse Matrix Collection. The characteristics of the matrices chosen is discussed in greater detail in section 4a "Benchmarks". Matrices are read from file by the master MPI process and converted to Compressed Sparse Row (CSR) or Compressed Sparse Column (CSC) format, depending on the individual matrix being input. Both CSR and CSC formats provide a reduction on memory requirements thereby increasing performance while reducing data transfer in multinode environments since only data about non-zero elements in the input matrix are kept [this needs a cite](#). While the matrix is being read in, the distribution of work amongst the MPI processes is being determined. The distribution pattern used by Bylina et al and which we have emulated, we call the *sub matrix method*, splits the input matrix  $A$  into  $p^2$  sub matrices in which each piece has nearly identical dimensions based on the number of processes  $p$  and the row or column count of  $A$ . Therein the size of each sub matrix from  $A_p$  will be  $A_{rows}/\sqrt{p} \times A_{cols}/\sqrt{p}$ . We chose to require that  $p$  be a non-negative square value.

The `csrSpMV` class was created to contain submatrix information in CSR/CSC format so that work allocation across MPI processes could be performed prior to MPI communication amongst those processes taking place, and stores information about each non-zero in three vectors. As each new non-zero is read the sub matrix it is to be assigned to is easily determined from its row and column, and is subsequently added to the `csrSpMV` object representing that submatrix.

Given that there are the same number of processes as submatrices,  $p$ , we can view the processes as being laid out in a matrix  $P$  in which process  $P_{ij}$  will receive data corresponding to submatrix  $A_{ij}$ . The first row of the process matrix containing processes with global MPI ranks 0 to  $p - 1$  are termed *column masters*, while the first column of processes with global ranks such that  $rank \% \sqrt{p}$  are *row*

*masters*. The MPI master process sends every column master the data contained within the csrSpMV object containing all data to be distributed amongst its column. Column masters will then send information about non-zeros to each process in its column. Each process receives its work allotment, if any, and proceeds to computation.

It is important to note that given the *sub matrix* distribution method considerable work imbalances including process with no work can occur between processes. As discussed in Bylina et al such imbalances can lead to entire computing nodes sitting idle as they have no data to process, thereby potentially decreasing overall performance.

#### B. OpenMP SpMV

#### C. Reduction and Validation

Upon completion of computation all nodes within a row perform an MPI reduction where each process's results are summed and stored within the row master's *result* vector. This is possible as each process within the same row of the process matrix are working on the same rows from  $A$ , but only on those non-zero elements contained within their assigned submatrix from  $A$ . After having performed the reduction a gather is performed on the process column containing the global master process (also a row master), and all other row masters. At this point the global master process now has all results and can proceed with secondary computation if necessary.

During development it was necessary to insure that the hybrid portion of the program was computing the correct SpMV result for a given matrix  $A$  and a dense vector. In order to verify accuracy of the hybrid version, a sequential version of the SpMV algorithm was performed on the master process only, prior to the hybrid portion of the program being performed. The results from each method were then compared and any differences indicated an error in computation. This was performed with a series of matrices, increasing in size, until no discrepancies were found amongst the test matrices. With the validity of the hybrid algorithms communication and computation tested and verified, the sequential master only computation was removed so that benchmarking tests could be performed.

#### D. Work Distribution

### IV. EVALUATION

- cluster architecture, network connectivity, number of nodes (max used), etc.
- how timings were taken and GFlops calculated based on these time measurements.
- note clock precision (nanoseconds)
- number of tests run per test permutation
- best, worst, and averages were taken for all times recorded and GFlops calculated
- then tables and pretty graphs to talk about

#### A. Benchmarks

why Parabolic\_Fem, bmw3\_2, torso1, and nd24k were chosen. -sparsity/density -size -symmetry  
 - how the GFlops will be calculated for symmetric and non-symmetric matrices (they are the same, however the Bylina paper did some weird shit with their numbers!)

#### B. Impact: Process Count

#### C. Impact: Thread Count

asdfsadf

#### D. Impact: Sparsity

### V. FUTURE WORK

### VI. CONCLUSIONS

### ACKNOWLEDGMENT

### REFERENCES

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- [2] Y. Saad, "Sparskit: A basic tool kit for sparse matrix computations," *Proceedings of the 2014 Federated Conference on Computer Science and Information Systems*, vol. 2, May 1990.