Time Complexity Analysis and Parallelization Report of Stock Prediction Pipeline

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1 Parallelization Report

1.1 Implementation

The parallelization of the stock analysis script was implemented using Dask, a flexible library for parallel computing in Python. The key steps in the implementation were: Setting up a Dask client, Creating a Dask bag from the list of files and Processing all files in parallel using Dask's map and compute functions. This approach allows for file-level parallelization, where each file is processed independently and concurrently.

1.2 Impact on Speed

The parallelization improved the processing speed of the script, particularly when dealing with a large number of files. My test revealed: A Speedup around 8x, performance gains were most noticeable with a higher number of available cores.

1.3 Reflection on the Development Process

Dask's high-level API significantly simplified the parallelization process, allowing me to maintain much of the original code structure, especially if you code in a functional style. The parallelized version demonstrated excellent scalability, on local multi-core machines. For very small datasets or when processing only a few files, I found that the overhead of setting up the Dask infrastructure could outweigh the benefits of parallelization.

2 Function-by-Function Analysis

- 1. $load_data(filename): O(n)$
 - This function primarily involves reading the CSV file, which scales linearly with the number of rows.
- 2. $add_features(df)$: O(n)
 - The rolling window operations (MA5, MA20, LR_Slope, RSI) and the exponential moving average (EMA) calculations for MACD all have mostly linear time complexity.
- 3. **feature_selection(df)**: $O(n \cdot f)$, where f is the number of features
 - The SelectKBest with f_regression algorithm has a time complexity of $O(n \cdot f)$.

- 4. **train_models(df)**: $O(k \cdot n^2 \cdot m)$, where k is the number of cross-validation folds, n is the number of samples, and m is the number of features This function's complexity is dominated by the cross-validation of the SVR model with RBF kernel.
 - RandomForestRegressor: $O(n_{\text{trees}} \cdot n \cdot \log(n) \cdot m)$
 - LinearRegression: $O(n \cdot m^2)$
 - SVR (RBF kernel): $O(n^2 \cdot m)$
 - \bullet Cross-validation multiplies each model's complexity by k (number of folds)
- 5. **process_file(filename)**: $O(k \cdot n^2 \cdot m)$, where k is the number of cross-validation folds, n is the number of samples, and m is the number of features
 - load_data(filename): O(n), where n is the number of rows in the file
 - add_features(df): O(n)
 - feature_selection(df): $O(n \cdot m)$
 - train_models(df): $O(k \cdot n^2 \cdot m)$ dominates the overall complexity
 - File operations (os.makedirs, joblib.dump): O(1) relative to data processing
 - The overall complexity is determined by the most computationally expensive operation, train_models
- 6. Overall Script Complexity: $O(F \cdot (k \cdot n^2 \cdot m)/P)$
 - F: Total number of files to process
 - k: Number of cross-validation folds
 - n: Number of samples in the largest file
 - m: Number of features
 - P: Number of parallel processes (depends on available cores)
 - File listing operations: O(F)
 - Dask bag creation: O(F)
 - Parallel processing with Dask:
 - Each file processed: $O(k \cdot n^2 \cdot m)$
 - Parallel execution reduces time by factor of P
 - The dominant factor is the parallel execution of process_file for each file

2.1 Machine Learning Algorithms

A more detailed look at the time complexities of the machine learning algorithms used:

- Linear Regression
 - Training: $O(nm^2)$
 - Prediction: O(m)
 - Generally fast for both training and prediction, especially when the number of features (m) is relatively small.
- Support Vector Regression (SVR)
 - Training: $O(n^2m)$
 - Prediction: O(vm), where v is the number of support vectors

 Potentially the slowest algorithm, especially for large datasets. The RBF kernel used in our script could lead to higher complexity.

• Random Forest Regressor

- Training: $O(t \cdot u \cdot n \log n)$, where u is the number of features considered for splitting
- Prediction: $O(t \log n)$
- Generally efficient, especially when the number of trees (t) and features considered for splitting (u) are not too large.

3 Comparison and Analysis

3.1 Theoretical Complexity

Sequential version: $O(F \cdot k \cdot n^2 \cdot m)$ Parallelized version: $O(\frac{F \cdot k \cdot n^2 \cdot m}{P})$ Where:

- F: Total number of files
- k: Number of cross-validation folds
- n: Number of samples in the largest file
- m: Number of features
- P: Number of parallel processes

3.2 Theoretical Speedup

The theoretical speedup of the parallelized version over the sequential version is:

Speedup =
$$\frac{O(F \cdot k \cdot n^2 \cdot m)}{O(\frac{F \cdot k \cdot n^2 \cdot m}{P})} = O(P)$$

This suggests a linear speedup with the number of processors, up to the number of files F.

3.3 Analysis

- The parallelized version distributes the workload across P processes, potentially reducing execution time by a factor of P.
- Ideal speedup is achieved when $P \leq F$, as each file can be processed independently.
- When P > F, additional processors may not contribute to further speedup due to the limited number of files.
- The space complexity increases with parallelization to $O(P \cdot n \cdot m)$, as each process requires memory for its data and models.

3.4 Limitations

- \bullet The speedup is bounded by the number of files F, following Amdahl's Law.
- I/O operations and network latency may become bottlenecks in a distributed setting.
- The effectiveness of parallelization depends on the uniformity of file sizes and processing times.

4 Sources

- 1. Virgolin, M. (2021). Time complexity for different machine learning algorithms. Retrieved from https://marcovirgolin.github.io/extras/details_time_complexity_machine_learning_algorithms/
- 2. Dask Development Team. (n.d.). Dask Documentation. Retrieved from https://docs.dask.org/en/stable/