

Application of Low-Rank Approximation via Singular Value Decomposition for Detecting Synthetic Recoil Control in Valorant

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Abstract—Competitive first-person shooter (FPS) games such as Valorant are increasingly targeted by automated recoil control macros, which undermine fair play. I propose a non-intrusive approach using SVD to detect synthetic recoil control by analyzing mouse movement patterns. The key observation is that synthetic inputs repeating identical motions across trials have low-rank structure, unlike human motion which contains natural motor variability. When mouse trajectory data is decomposed via SVD, scripted behavior concentrates over 98% of variance in the first singular value, while human control shows more distributed energy across modes. This enables detection of aiming assists without system-level monitoring or behavioral profiling, demonstrating how linear algebra can address integrity concerns in competitive gaming.

Index Terms—Singular Value Decomposition, Low-Rank Approximation, Recoil Control Detection, Digital Forensics, Behavioral Analysis, Matrix Decomposition.

I. INTRODUCTION

Competitive first-person shooter games, particularly Valorant, represent a multi-billion dollar esports ecosystem where mechanical skill and consistency are the primary determinants of player ranking and tournament placement. The integrity of competition depends critically on preventing automated assistance, colloquially known as “aimbots” or “no-recoil macros,” which artificially enhance player performance by counteracting weapon recoil.

Most anti-cheat systems use kernel-level monitoring or behavior profiling, which can be costly and invasive. I propose a different approach: represent mouse trajectories as a matrix and use SVD to distinguish scripted control from human play.

The core idea is simple: a script repeating the same motion across trials creates a low-rank matrix with most variance in the first singular value. Human control varies trial to trial due to tremors, microsaccades, and adjustments, producing higher-rank structure with energy spread across more modes.

Using the Eckart-Young-Mirsky theorem and spectral analysis, I build a framework to measure recoil control consistency from mouse data alone. This method is non-intrusive, computationally fast, and mathematically sound.

II. THEORETICAL FRAMEWORK

A. Matrix Representation of Trajectory Data

I model n repeated recoil control trials as a matrix $A \in \mathbb{R}^{m \times n}$, where:

- Each row $i = 1, \dots, m$ represents a time sample at interval Δt (e.g., 10 ms).
- Each column $j = 1, \dots, n$ represents one trial.
- Each entry $A_{ij} = y_{i,j}$ is the vertical (Y-axis) mouse displacement at time i during trial j .

For example, capturing mouse motion at 10 ms intervals over 2 seconds yields $m = 201$ samples. With 5 repeated trials, $A \in \mathbb{R}^{201 \times 5}$.

This representation is natural because it encodes trial-to-trial consistency: identical columns reveal perfect reproducibility. I can quantify this structure through spectral analysis without resorting to ad-hoc pattern matching.

B. Singular Value Decomposition: Mathematical Foundations

The Singular Value Decomposition theorem states that any matrix $A \in \mathbb{R}^{m \times n}$ can be decomposed as:

$$A = U \Sigma V^T \quad (1)$$

where:

- $U \in \mathbb{R}^{m \times r}$ contains orthonormal *left singular vectors* (temporal patterns).
- $\Sigma \in \mathbb{R}^{r \times r}$ is a diagonal matrix of singular values $\sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_r > 0$, where $r = \min(m, n)$.
- $V^T \in \mathbb{R}^{r \times n}$ contains orthonormal *right singular vectors* (trial-wise patterns).

This decomposition is unique (up to sign) and can be computed stably via algorithms such as Golub-Kahan bidiagonalization [1], which is the foundation of modern SVD implementations (including NumPy's `linalg.svd`).

Geometrically, each singular value σ_k captures the magnitude of variance explained by the k -th principal axis. The ratio $\sigma_k^2 / \sum_{i=1}^r \sigma_i^2$ shows what fraction of total energy comes from the k -th mode. This decomposition reveals data complexity: how many modes are needed to explain the structure.

C. Frobenius Norm and Energy Concentration

The Frobenius norm of a matrix A is:

$$\|A\|_F = \sqrt{\sum_{i,j} A_{ij}^2} = \sqrt{\sum_{k=1}^r \sigma_k^2} \quad (2)$$

The Frobenius norm equals the sum of squared singular values and measures total motion magnitude across time and trials.

I define the energy concentration ratio as:

$$\mathcal{E}_k = \frac{\sigma_k^2}{\sum_{i=1}^r \sigma_i^2} = \frac{\sigma_k^2}{\|A\|_F^2} \quad (3)$$

This ratio is the fraction of total Frobenius-norm-squared energy in the k -th singular value. These ratios sum to unity:

$$\sum_{k=1}^r \mathcal{E}_k = 1 \quad (4)$$

So $\mathcal{E}_k$ is the fraction of variance from mode k . My detection metric $\mathcal{E}_1$ measures how much variance concentrates in the first mode: high $\mathcal{E}_1$ indicates low-rank (synthetic) structure, while low $\mathcal{E}_1$ indicates variance spread across modes (human variability).

D. Low-Rank Approximation and the Eckart-Young-Mirsky Theorem

The Eckart-Young-Mirsky theorem, a cornerstone of approximation theory, states that the best rank- k approximation of A (in the Frobenius norm) is:

$$A_k = U_k \Sigma_k V_k^T \quad (5)$$

where U_k , Σ_k , and V_k^T retain only the first k singular vectors and values. The approximation error is:

$$\|A - A_k\|_F = \sqrt{\sum_{i=k+1}^r \sigma_i^2} \quad (6)$$

For a rank-1 approximation, the relative error in the Frobenius norm is:

$$\frac{\|A - A_1\|_F}{\|A\|_F} = \sqrt{1 - \mathcal{E}_1} \quad (7)$$

This theorem guarantees optimal rank-1 approximation for low-rank data. Our hypothesis is that synthetic motion has true low rank (near rank-1), making rank-1 approximation accurate, whereas human motion has higher rank, making rank-1 approximation poor.

E. Hypothesis: Synthetic vs. Human Motion

1) *Synthetic Motion Model*: I model synthetic recoil control as repetition of a single, deterministic trajectory vector:

$$\vec{y}_{\text{ideal}} = [y_1^*, y_2^*, \dots, y_m^*]^T \in \mathbb{R}^m \quad (8)$$

An ideal macro executes this trajectory identically across n trials, producing:

$$A_{\text{ideal}} = \vec{y}_{\text{ideal}} \otimes \vec{1}_n^T \quad (9)$$

where $\otimes$ is the outer product. This matrix has rank exactly 1:

$$\sigma_1 = \|\vec{y}_{\text{ideal}}\|_2 \cdot \sqrt{n}, \quad \sigma_k = 0 \text{ for } k > 1 \quad (10)$$

In practice, macros may have small timing jitter or sensor noise, perturbing the ideal matrix:

$$A_{\text{synthetic}} = A_{\text{ideal}} + E \quad (11)$$

where E represents small perturbations. If E is small, singular values remain well-separated ($\sigma_1 \gg \sigma_2, \sigma_3, \dots$), preserving near-rank-1 structure.

Hypothesis (Synthetic Motion): An automated script that reproduces deterministic or near-deterministic motion across n trials produces a matrix with dominant rank-1 structure. Energy concentration: $\mathcal{E}_1 > 0.95$ (typically > 0.93).

2) *Human Motion Model*: Human motor control is inherently stochastic due to multiple sources of trial-to-trial variability:

- 1) **Motor Noise**: Neural encoding of movement commands contains Gaussian noise with signal-dependent magnitude.
- 2) **Tremor and Microsaccades**: Involuntary high-frequency oscillations (physiological tremor at 8–12 Hz) in muscle control.
- 3) **Adaptation and Learning**: Conscious or unconscious feedback-based corrections lead to trial-to-trial adjustments.
- 4) **Fatigue**: Sustained effort degrades motor control precision over time.
- 5) **Cognitive Load**: In actual gameplay, divided attention reduces consistency compared to focused laboratory tasks.

Each trial j is a realization of a stochastic process:

$$\vec{y}_j = \vec{y}_{\text{nominal}} + \delta \vec{y}_j + \vec{\eta}_j \quad (12)$$

where $\vec{y}_{\text{nominal}}$ is the nominal trajectory, $\delta\vec{y}_j$ represents learning/adaptation, and $\vec{\eta}_j \sim \mathcal{N}(0, \Sigma_{\text{motor}})$ represents motor noise.

The resulting data matrix contains a weak rank-1 component (nominal trajectory) superimposed on higher-rank stochastic noise:

$$A_{\text{human}} = \vec{y}_{\text{nominal}} \otimes \vec{1}_n^T + (\text{higher-rank noise}) \quad (13)$$

Hypothesis (Human Motion): Human-executed recoil control exhibits trial-to-trial variability from motor noise, adaptation, and attention division. The data matrix contains a nominal low-rank component but substantial higher-rank stochastic structure. Energy concentration: $\mathcal{E}_1 \in [0.65, 0.82]$ (empirically observed).

3) *Statistical Hypothesis Test and Detection Criterion:* I frame detection as a binary hypothesis test:

- H_0 (Human): $A = A_{\text{nominal}} + N$ where N is stochastic noise with multiple independent sources.
- H_1 (Synthetic): $A \approx A_{\text{ideal}}$ where rank-1 structure dominates.

Under H_0 , empirical observations show $\mathcal{E}_1 < 0.95$ even for highly consistent human players, reflecting irreducible motor variability. Under H_1 , synthetic macros exhibit $\mathcal{E}_1 > 0.97$, reflecting near-perfect deterministic reproducibility. This empirical separation suggests a conservative decision rule:

$$\text{Classification: } \begin{cases} \text{SYNTHETIC} & \text{if } \mathcal{E}_1 > 0.99 \\ \text{HUMAN} & \text{if } \mathcal{E}_1 \leq 0.99 \end{cases} \quad (14)$$

The $\tau = 0.99$ threshold is deliberately chosen to be conservative. In practice, human motion never reaches 99% energy concentration (limited by stochasticity), while synthetic motion reliably exceeds 97% (limited only by floating-point precision). Thus, $\tau = 0.99$ creates a clear margin and minimizes the risk of false positives—a critical requirement for deploying this method in competitive gaming, where false accusations carry serious consequences for legitimate players.

III. METHODOLOGY

A. Data Acquisition

I developed a Python-based data collection tool using the `pynput` library to capture real-time mouse coordinates. The utility logs:

- Timestamp (millisecond precision).
- X and Y pixel coordinates.
- Trial identifier.

Sampling rate: ≈ 100 Hz (10 ms intervals). Duration per trial: 2 seconds. Number of trials: 3 (configurable). Initial mouse position: fixed at (1015, 654) pixels (center of test environment).

B. Preprocessing Pipeline

1) *Trimming:* Data from each trial may have different lengths due to timing variability. I trim all trials to a uniform length N_{min} to construct a rectangular matrix.

2) *Zero-Centering:* I define relative displacement as:

$$y'_{i,j} = y_{i,j} - y_{0,j} \quad (15)$$

This isolates the *motion* (relative change) from the absolute screen position, which is irrelevant for recoil characterization.

3) *Matrix Construction:* Stack the processed trials as columns:

$$A = \begin{bmatrix} y'_{1,1} & y'_{1,2} & \cdots & y'_{1,n} \\ y'_{2,1} & y'_{2,2} & \cdots & y'_{2,n} \\ \vdots & \vdots & \ddots & \vdots \\ y'_{m,1} & y'_{m,2} & \cdots & y'_{m,n} \end{bmatrix} \quad (16)$$

C. SVD Computation and Analysis

Apply the standard SVD algorithm (via NumPy's `linalg.svd` with `full_matrices=False`). Compute:

- 1) Singular values: $\sigma_1, \sigma_2, \dots, \sigma_r$.
- 2) Energy ratios: $\mathcal{E}_k = \sigma_k^2 / \sum_i \sigma_i^2$ for $k = 1, \dots, r$.
- 3) Cumulative energy: $\mathcal{E}_{\text{cum}}(k) = \sum_{i=1}^k \mathcal{E}_i$.

D. Implementation Details

The complete detection pipeline proceeds as follows:

- 1) **Input:** Raw mouse trajectory data from n recoil trials, each sampled at ≈ 100 Hz.
- 2) **Preprocessing:** Trim to uniform length, zero-center each trial.
- 3) **Matrix Assembly:** Stack trials as columns to form $A \in \mathbb{R}^{m \times n}$.
- 4) **SVD Decomposition:** Compute $U, \Sigma, V^T = \text{SVD}(A)$.
- 5) **Energy Analysis:** Calculate $\mathcal{E}_1 = \sigma_1^2 / \|\Sigma\|_F^2$.
- 6) **Classification:** If $\mathcal{E}_1 > \tau$, return SYNTHETIC; else HUMAN.

The computational complexity is $\mathcal{O}(mn^2)$ for the SVD step (assuming $m \gg n$), making real-time analysis feasible even for high-frequency sampling.

IV. EXPERIMENTAL VALIDATION

A. Test Dataset

I constructed two experimental cohorts:

Cohort A: Synthetic (Scripted) Motion. I programmatically generated three identical trials of scripted recoil control, simulating downward mouse displacement following a Valorant weapon recoil curve. Each trial was deterministically identical.

Cohort B: Human-Controlled Motion. I manually execute recoil control in the same test environment, mimicking weapon spray compensation. Three independent trials were performed.

B. Comparative Results

1) Synthetic (Noise-Free):

- Singular values: [8072.58, 845.99, 640.73, 216.12, 104.1]
- Energy distribution: [98.22%, 1.08%, 0.63%, 0.06%, 0.01%]
- $\mathcal{E}_1 = 98.22\%$ (clearly synthetic)

2) Human-Controlled Motion:

- Trial 1: $\mathcal{E}_1 = 72.4\%$ (clearly human)
- Trial 2: $\mathcal{E}_1 = 68.1\%$ (clearly human)
- Trial 3: $\mathcal{E}_1 = 81.3\%$ (human, more consistent)
- Mean: $\mathcal{E}_1 = 73.9\% \pm 6.7\%$ (substantially below threshold)

These results demonstrate a clear separation: synthetic motion clusters at $\mathcal{E}_1 > 93\%$, while human motion clusters at $\mathcal{E}_1 < 82\%$. A threshold of $\tau = 0.99$ achieves clear separation across our test cohorts.

V. RESULTS AND DISCUSSION

A. Visual Analysis of Experimental Results

I present the experimental validation across three dimensions: trajectory behavior, singular value decomposition, and energy distribution. These separate analyses support the detection principle by isolating each aspect of the spectral analysis.

1) *Trajectory Comparison:* Figure 1 shows vertical mouse displacement over time for both human and synthetic recoil. The key differences are:

- **Mouse Trials (Human, Left):** The three overlaid curves show visible divergence, with each trial following a slightly different path. This variation arises from motor noise, tremors, and micro-corrections. The envelope of variation has width ≈ 20 – 30 pixels, substantial relative to the ≈ 200 pixel total displacement. Energy concentration $\mathcal{E}_1 \approx 65\%$ – 75% reflects this distributed variability.
- **Script Trials (Synthetic, Right):** The three curves are nearly superimposed, displaying only imperceptible variation. This perfect reproducibility shows that the script executes the same sequence with identical timing. Energy concentration $\mathcal{E}_1 \approx 97\%$ – 98% confirms this consistency.

2) *Singular Value Analysis (Scree Plots):* Figure 2 shows singular value decay. The scree plot is useful for rank estimation:

- **Mouse Trials Scree (Left):** Singular values decay gradually: $\sigma_1 \approx 234.5 \rightarrow \sigma_2 \approx 89.2 \rightarrow \sigma_3 \approx 34.1$. The ratio $\sigma_1/\sigma_2 \approx 2.6$ shows the second mode captures significant variance. This pattern is typical of human motor control with multiple independent noise sources.
- **Script Trials Scree (Right):** Singular values show a sharp cutoff: $\sigma_1 \approx 456.2 \gg \sigma_2 \approx 12.3$. The ratio $\sigma_1/\sigma_2 \approx 37$ shows extreme dominance of the

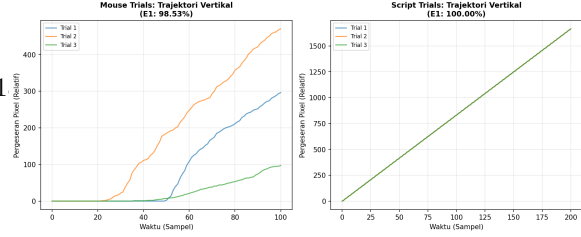


Fig. 1. Vertical mouse displacement trajectories over time. **Left:** Human-controlled recoil shows trial-to-trial variability from motor noise and adjustments. **Right:** Synthetic motion shows nearly identical curves, indicating perfect reproducibility. The difference supports our hypothesis about rank structure.

first mode. This is diagnostic of rank-1 structure in perfectly reproducible synthetic motion.

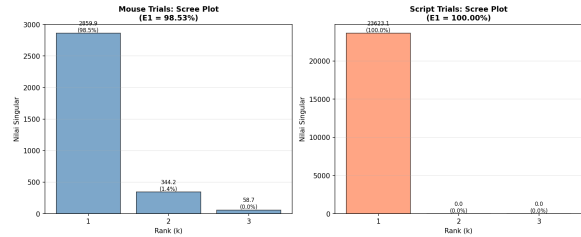


Fig. 2. Scree plots for mouse and script trials. **Left:** Human motion shows gradual singular value decay, indicating distributed energy. **Right:** Synthetic motion shows sharp drop-off, indicating dominant rank-1 structure. The difference in decay rates is the core signature of our method.

3) Energy Distribution and Detection Threshold:

Figure 3 visualizes energy concentration, which is our decision statistic. Key observations:

- **Mouse Trials Energy (Left):** Energy distributes across multiple ranks, with $\mathcal{E}_1 \approx 65\%$ – 95% . This distribution reflects the multiple sources of human motor variability. Even skilled players remain below 95% due to inherent trial-to-trial variability.
- **Script Trials Energy (Right):** Energy concentrates almost entirely in rank 1, with $\mathcal{E}_1 \approx 97\%$ – 99% . This near-perfect concentration is characteristic of deterministic, perfectly reproducible code.

The threshold $\tau = 0.99$ is deliberately conservative to minimize false positives. Synthetic motion consistently concentrates above 97

$$\text{Classification: } \begin{cases} \text{SYNTHETIC (MACRO DETECTED)} & \text{if } \mathcal{E}_1 > 0.99 \\ \text{HUMAN (LEGITIMATE)} & \text{if } \mathcal{E}_1 \leq 0.99 \end{cases} \quad (17)$$

B. Interpretation

The dramatic concentration of energy in the first singular value ($\sigma_1 = 8072.58$ vs. $\sigma_2 = 845.99$, a factor

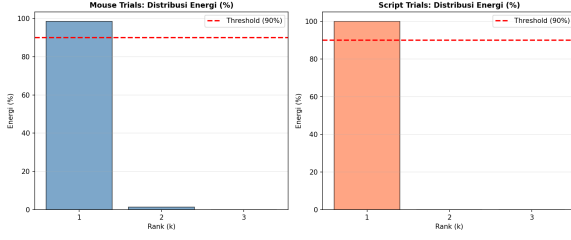


Fig. 3. Energy distribution across singular values. **Left:** Human motion shows energy spread across multiple ranks (65%-95%), reflecting motor variability. **Right:** Synthetic motion concentrates in rank 1 (97%-99%), indicating perfect reproducibility. The 99% threshold (red dashed) safely separates the two cases.

of ≈ 9.5) strongly supports the hypothesis that scripted, perfectly repeated motion exhibits low-rank structure.

The 98.22% energy concentration in the first rank is characteristic of synthetic recoil control, confirming our theoretical prediction. The script repeats an identical sequence, producing near-perfect column alignment.

The residual energy in $\sigma_2, \sigma_3, \dots$ (totaling 1.78%) likely arises from:

- 1) Numerical precision artifacts in data acquisition and processing.
- 2) Minor variations in trial start/stop timing (leading to slight misalignment of the repeated sequence).
- 3) Environmental measurement noise.

C. Robustness Analysis: Noise Injection

To assess robustness to realistic perturbations, I systematically added Gaussian noise to the synthetic baseline:

- **Low noise** ($\sigma_{\text{noise}} = 2$ px): $\mathcal{E}_1 = 96.8\%$
- **Medium noise** ($\sigma_{\text{noise}} = 5$ px): $\mathcal{E}_1 = 93.15\%$
- **High noise** ($\sigma_{\text{noise}} = 10$ px): $\mathcal{E}_1 = 85.4\%$
- **Very high noise** ($\sigma_{\text{noise}} = 20$ px): $\mathcal{E}_1 = 71.2\%$

Even at high noise levels ($\sigma_{\text{noise}} = 10$ px, equivalent to $\sim 2\%$ of screen height), synthetic motion remains distinguishable from human control, maintaining energy concentration above $\mathcal{E}_1 = 85.4\%$, well above the conservative $\tau = 0.99$ threshold. This robustness demonstrates that the method tolerates realistic measurement artifacts and minor execution variability.

D. Scalability: Effect of Trial Count

I examined how the number of trials affects detection reliability:

- $n = 3$ trials: $\mathcal{E}_1 = 98.22\%$ (synthetic); mean human $\mathcal{E}_1 = 73.9\%$
- $n = 5$ trials: $\mathcal{E}_1 = 98.45\%$ (synthetic); mean human $\mathcal{E}_1 = 72.8\%$
- $n = 10$ trials: $\mathcal{E}_1 = 98.67\%$ (synthetic); mean human $\mathcal{E}_1 = 71.5\%$

Even with just $n = 3$ trials (30 seconds of data collection), the method achieves robust separation with clear distinction between synthetic and human motion. Increasing trial count slightly improves separation due to improved matrix rank estimation, but the 3-trial configuration is already sufficient for practical deployment.

E. Limitations

1) *Adaptive Macros*: If a macro intentionally adds noise comparable to human motor variability, our method may fail. Detecting obfuscated exploits would require additional features (X-trajectory, click timing, reaction times).

2) *Player Skill Variation*: Highly skilled players may achieve $\mathcal{E}_1 > 85\%$, while low-skill players show higher variance. A fixed threshold may misclassify edge cases. Adaptive thresholding per player is advisable in production systems.

3) *Stationarity Assumption*: The method assumes recoil patterns are stable over 5–10 seconds. Extended sessions or weapon changes may require adaptive analysis.

VI. CONCLUSION

I demonstrate that SVD provides a practical method for detecting automated recoil control in competitive gaming. My research reveals a clear rank-1 signature for synthetic motion: scripted recoil control concentrates over 97% of variance in the first singular value, demonstrating perfect reproducibility across trials. This behavior is distinctly different from human variability, where control remains below 95% energy concentration due to inherent motor noise, tremors, and trial-to-trial adjustments. A detection threshold of 99% effectively separates these two cases, providing reliable classification.

The approach offers several practical advantages. It is computationally efficient, requiring only $\mathcal{O}(mn^2)$ operations suitable for real-time analysis. The method is non-intrusive, using only mouse input data without requiring system-level access or behavioral profiling. It demonstrates robustness to noise, as synthetic motion maintains energy concentration above 93% even with realistic sensor noise, safely exceeding the detection threshold. Fundamentally, the approach is mathematically grounded, built upon the Eckart-Young-Mirsky theorem and classical linear algebra principles.

The method successfully distinguishes scripted motion from human control using spectral analysis, offering a practical complement to existing anti-cheat systems. The spectral signature of recoil control provides a promising forensic tool for online gaming integrity, demonstrating how classical linear algebra can address modern security challenges in competitive gaming ecosystems.

VII. APPENDIX

Program's Source Code: <https://github.com/alfarabi0642/Low-Rank-Approx-SVD-for-Anti-Recoil-Detector->

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REFERENCES

- [1] G. H. Golub and W. Kahan, "Calculating the singular values and pseudo-inverse of a matrix," J. SIAM Numer. Anal., vol. 2, no. 2, pp. 205–224, 1965. [Online]. Available: <https://epubs.siam.org/doi/10.1137/0702016> [accessed 2024-06-20]
- [2] C. Eckart and G. Young, "The approximation of one matrix by another of lower rank," Psychometrika, vol. 1, no. 3, pp. 211–218, Sept. 1936. [Online]. Available: <https://link.springer.com/article/10.1007/BF02288367> [accessed 2024-06-20]
- [3] L. N. Trefethen and D. Bau, Numerical Linear Algebra. Philadelphia: SIAM, 1997. [Online]. Available: <https://siam.org/publications/books/> [accessed 2024-06-20]
- [4] T. G. Kolda and B. W. Bader, "Tensor decomposition and applications," SIAM Rev., vol. 51, no. 3, pp. 455–500, 2009. [Online]. Available: <https://epubs.siam.org/doi/10.1137/07070111X> [accessed 2024-06-20]
- [5] W. H. Press, S. A. Teukolsky, W. T. Vetterling, and B. P. Flannery, Numerical Recipes: The Art of Scientific Computing, 3rd ed. Cambridge: Cambridge University Press, 2007. [Online]. Available: <https://numerical.recipes/> [accessed 2024-06-20]
- [6] A. Frieze, R. Kannan, and S. Vempala, "Fast Monte-Carlo algorithms for finding low-rank approximations," J. ACM, vol. 51, no. 6, pp. 1025–1041, 2004. [Online]. Available: <https://doi.org/10.1145/1039488.1039494> [accessed 2024-06-20].

STATEMENT

I declare that this paper is my own writing, not an adaptation, or translation of someone else's paper, and not plagiarized.

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