

Advanced Linear Algebra AOL Documentation

Nathanael Romaloburju¹, Richie Rizawardana², Robben Wijanathan³, and Weneville⁴

¹2802538132, *Computer Science & Mathematics, Binus University, Jakarta, Indonesia*

²2802403100, *Computer Science & Mathematics, Binus University, Jakarta, Indonesia*

³2802461681, *Computer Science & Mathematics, Binus University, Jakarta, Indonesia*

⁴2802549016, *Computer Science & Mathematics, Binus University, Jakarta, Indonesia*

ABSTRACT

This project presents a simple Python-based Matrix Analyzer program designed to perform key linear algebra operations. The system provides an interactive menu with four main features: checking if a matrix is diagonalizable, performing LU decomposition, computing the dominant eigenvalue using the Power Method, and conducting Singular Value Decomposition (SVD). Each operation is implemented using fundamental numerical methods in Python.

1 MENU USER INTERFACE

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```
44         print("\nExiting Matrix Analyzer Tool. Goodbye!")
45         break
46     else:
47         print("\nInvalid choice. Please enter a number from 1 to 5.")
```

- The function `clear()` clears the terminal screen depending on the operating system (Windows or Linux/macOS).
- The function `mainMenu()` displays an ASCII art banner and lists five main options for matrix operations:
 1. Check Diagonalizability
 2. Perform LU Decomposition
 3. Find Dominant Eigenvalue
 4. Perform Singular Value Decomposition (SVD)
 5. Exit the program
- It then continuously prompts the user for input (1-5) and calls the corresponding function (e.g., `isDiagonalizable()`, `LUdecomposition()`, etc.).
- If the input is invalid, an error message is displayed until a valid choice is entered.

Output

(.)

> <

,

/

1. Check Diagonalizability
2. Perform LU Decomposition
3. Find Dominant Eigenvalue
4. Singular Value Decomposition (SVD)
5. Exit

Enter your choice (1-5): a

Invalid choice. Please enter a number from 1 to 5.

Enter your choice (1-5): 0

Invalid choice. Please enter a number from 1 to 5.

Enter your choice (1-5): 1

2 INPUT MATRIX FUNCTIONS

These two functions handle user input for matrix creation, ensuring valid and properly formatted entries before performing matrix operations.

2.1 $m \times n$ Matrix (For Singular Value Decomposition)

```

1 def input_matrix():
2     while True:
3         try:
4             rows = int(input("Enter number of rows: "))
5             cols = int(input("Enter number of columns: "))
6             break
7         except ValueError:
8             print("Please enter valid integers for rows and columns.\n")
9
10    print("\nEnter the matrix values row by row (separated by spaces):")
11    matrix = []
12

```

```

13     for i in range(rows):
14         while True:
15             row_input = input(f"Row {i + 1}: ").split()
16
17             if len(row_input) != cols:
18                 print(f"Row {i + 1} must have exactly {cols} entries. Try again.")
19                 continue
20
21             try:
22                 row = [float(x) for x in row_input]
23                 matrix.append(row)
24                 break
25             except ValueError:
26                 print(f"Row {i + 1} contains non-numeric values. Try again.")
27
28     return matrix

```

The function `input_matrix()` is designed to input a general rectangular matrix of size $m \times n$

- The user is first prompted to enter the number of rows (m) and columns (n), with input validation to ensure they are integers.
- Then, for each row, the user must input numeric values separated by spaces.
- The program checks that:
 - Each row contains exactly n entries.
 - All entries are valid numeric values (converted to float).
- If invalid input is detected, an error message is displayed and the user is asked to re-enter that row.
- Once all inputs are valid, the function returns the complete matrix as a list of lists.

2.2 $n \times n$ Square Matrix (For Checking Diagonalizability, LU Decomposition, & Finding Dominant Eigenvalue)

```

1 def input_squareMatrix():
2     while True:
3         try:
4             n = int(input("Enter the size of the square matrix (n x n): "))
5             if n <= 0:
6                 print("Matrix size must be a positive integer.\n")
7                 continue
8             break
9         except ValueError:
10            print("Please enter a valid integer for the matrix size.\n")
11
12    print(f"\nEnter the {n}x{n} matrix values row by row (separated by spaces):")
13    matrix = []
14
15    for i in range(n):
16        while True:
17            row_input = input(f"Row {i + 1}: ").split()
18
19            if len(row_input) != n:
20                print(f"Row {i + 1} must have exactly {n} entries. Try again.")
21                continue
22
23            try:
24                row = [float(x) for x in row_input]
25                matrix.append(row)
26                break
27            except ValueError:
28                print(f"Row {i + 1} contains non-numeric values. Try again.")
29
30    return matrix

```

The function `input_squareMatrix()` is used for creating a square matrix of size $n \times n$.

- It starts by asking the user for the matrix size n and checks that the input is a positive integer.
- The user is then prompted to enter each row of the matrix, with input separated by spaces.
- The program validates that:
 - Each row contains exactly n numeric entries.
 - Non-numeric or mismatched inputs trigger an error message and require re-entry.
- After successful input, the matrix is returned as a list of lists to be used in other operations such as diagonalizability checking, LU decomposition, or eigenvalue calculation.

3 CORE MAIN FUNCTIONS

3.1 Diagonalizable Matrix

Theory A square matrix A of size $n \times n$ is said to be **diagonalizable** if it can be expressed as:

$$A = PDP^{-1}$$

where D is a diagonal matrix whose diagonal elements are the eigenvalues of A , and P is a matrix whose columns are the corresponding eigenvectors of A .

- **Diagonalizability:** A matrix is diagonalizable if there exists an invertible matrix P such that $A = PDP^{-1}$. This means the linear transformation represented by A can be simplified to scaling operations along independent directions defined by the eigenvectors.
- **Eigenvalues:** The eigenvalues of a matrix A are scalars λ that satisfy the equation:

$$A\mathbf{v} = \lambda \mathbf{v}$$

where \mathbf{v} is a non-zero vector. Each eigenvalue represents how its corresponding eigenvector is scaled when transformed by A .

- **Eigenvectors:** The eigenvectors are non-zero vectors that do not change direction when multiplied by A ; only their magnitude changes by the factor of their eigenvalue.

A matrix is diagonalizable if and only if it has n linearly independent eigenvectors, allowing it to be represented as a diagonal matrix in the basis of those eigenvectors.

[illegible]

Press Enter to go back to main menu...

Press Enter to go back to main menu...

- **Computing Determinants:** The determinant of A can be easily computed as:

$$\det(A) = \det(L) \times \det(U)$$

Since $\det(L) = 1$, it follows that $\det(A) = \prod_i U_{ii}$, the product of the diagonal elements of U .

- **Finding Inverses:** LU decomposition also aids in finding the inverse of a matrix by solving multiple systems of linear equations for each column of the identity matrix.

- **Numerical Stability:** When row exchanges are required for stability, a **permutation matrix** P is introduced:

$$PA = LU$$

This process, known as **partial pivoting**, minimizes numerical errors in floating-point computations.

[illegible]

Explanation The function `LUdecomposition()` performs an LU decomposition of a user-input square matrix without using external libraries. It breaks the given matrix A into a **lower triangular matrix** L and an **upper triangular matrix** U such that:

$$A = LU$$

- The function first clears the screen and displays an ASCII art title for better presentation.
- It then calls the `input_squareMatrix()` function to receive the matrix input from the user.
- Two empty matrices, `lower` and `upper`, of size $n \times n$ are initialized with zeros.
- The decomposition process is done in two nested loops:
 - **Upper Triangular Calculation:** For each row i , the elements of U are computed using:

$$U_{i,k} = A_{i,k} - \sum_{j=0}^{i-1} L_{i,j} U_{j,k}$$

This ensures that U contains nonzero values only on and above its main diagonal.

- **Lower Triangular Calculation:** For each column i , the diagonal of L is set to 1. The remaining elements below the diagonal are computed as:

$$L_{k,i} = \frac{A_{k,i} - \sum_{j=0}^{i-1} L_{k,j} U_{j,i}}{U_{i,i}}$$

This ensures that L contains ones on its diagonal and nonzero values below it.

- The function also checks if any pivot element $U_{i,i} = 0$, in which case the matrix is **singular** and LU decomposition cannot be performed.
- After the decomposition, both L and U are printed with values rounded to two decimal places for readability.
- Finally, the program waits for the user to press Enter before returning to the main menu.

Output

[The following section contains extremely faint, illegible markings that appear to be bleed-through from another page or are artifacts of the scanning process.]

Enter the size of the square matrix (n x n): 3

Enter the 3x3 matrix values row by row (separated by spaces):

Row 1: 1 2 3

Row	1:	2	3	4	5
Row	2:	3	4	5	

Row	3:	5	6	7
-----	----	---	---	---

Lower Triangular Matrix:

```
['1.00', '0.00', '0.00']
```

```
['3.00', '1.00', '0.00']
```

```
['5.00', '2.00', '1.00']
```

Upper Triangular Matrix:

```
['1.00', '2.00', '3.00']
```

```
['0.00', '-2.00', '-4.00']
```

```
[ '0.00 ', '0.00 ', '0.00 ']
```

Press Enter to go back to main menu...

Output (Singular)

[illegible]

Enter the size of the square matrix (n x n): 3

Enter the 3x3 matrix values row by row (separated by spaces):

Row 1: 2 4 6

Row 2: 3 6 1

Row 3: 7 8 5

Matrix is singular, cannot perform LU decomposition.

Press Enter to go back to main menu...

3.3 Dominant Eigenvalue

Theory In the context of a square matrix $A \in \mathbb{R}^{n \times n}$, the *dominant eigenvalue* is usually understood to be an eigenvalue λ_1 satisfying

$$|\lambda_1| > |\lambda_j| \quad \text{for all } j = 2, 3, \dots, n.$$

That is, the magnitude of λ_1 strictly exceeds that of every other eigenvalue of A . A corresponding eigenvector $\mathbf{v}_1 \neq \mathbf{0}$ satisfies

$$A \mathbf{v}_1 = \lambda_1 \mathbf{v}_1.$$

A common numerical method to find λ_1 (and \mathbf{v}_1) is the *Power Method*. The basic algorithm proceeds as follows:

- Choose an initial non-zero vector \mathbf{b}_0 .
- For $k = 0, 1, 2, \dots$:

$$\mathbf{b}_{k+1} = \frac{A \mathbf{b}_k}{\|A \mathbf{b}_k\|}.$$

- After each iteration one may estimate the eigenvalue via the Rayleigh quotient:

$$\mu_k = \frac{\mathbf{b}_k^T (A \mathbf{b}_k)}{\mathbf{b}_k^T \mathbf{b}_k}.$$

Convergence Criteria and Remarks:

- The method converges (i.e., $\mathbf{b}_k \rightarrow \pm \mathbf{v}_1$ and $\mu_k \rightarrow \lambda_1$) provided that:
 1. A has a unique eigenvalue λ_1 such that $|\lambda_1| > |\lambda_j|$ for all $j > 1$.
 2. The initial vector \mathbf{b}_0 has a non-zero component in the direction of \mathbf{v}_1 ; equivalently, when writing $\mathbf{b}_0 = c_1 \mathbf{v}_1 + c_2 \mathbf{v}_2 + \dots + c_n \mathbf{v}_n$, the coefficient $c_1 \neq 0$.
- The rate of convergence is governed by the ratio

$$\left| \frac{\lambda_2}{\lambda_1} \right|,$$

where λ_2 is the eigenvalue of next largest absolute value (i.e., the “sub-dominant” eigenvalue). If $\left| \frac{\lambda_2}{\lambda_1} \right|$ is close to zero, convergence is very rapid; if it is close to one, convergence is slow.

- If $|\lambda_1| = |\lambda_2|$ (or there’s a dominant eigenspace of dimension ≥ 1), or if \mathbf{b}_0 is orthogonal to \mathbf{v}_1 , the power method may fail to converge to the correct eigen-pair.

```

1 def power_method(A, iter=2000, tolerance=1e-9):
2     n, _ = A.shape
3     x = np.ones(n, dtype=float)
4     x = x / np.linalg.norm(x)
5
6     eigen_start = 0.0
7     x_prevprev = None
8
9     for _ in range(iter):

```

```

10     Ax = A @ x
11     normAx = np.linalg.norm(Ax)
12     if normAx == 0.0:
13         return 0.0, x
14
15     x_new = Ax / normAx
16     eigen_new = float(np.dot(x_new, Ax) / np.dot(x_new, x_new))
17
18     if x_prevprev is not None:
19         if abs(np.dot(x_new, x_prevprev)) > 1.0 - 1e-8:
20             return None, None # signal: no unique dominant eigenvalue (tie/complex)
21
22     if abs(eigen_new - eigen_start) < tolerance:
23         return eigen_new, x_new
24
25     x_prevprev = x
26     x = x_new
27     eigen_start = eigen_new
28
29     return None, None # non-convergent within iter
30
31 def dominantEigenvalue():
32     clear()
33     print(r"""
34
35 / _ \ _ _ _ _ _ ( _ ) _ _ _ _ _ / _ \
36 / / / _ _ \ / ' \ / _ _ \ / _ _ \ / _ \
37 / _ _ \ _ _ / / / / / / / / \ _ _ \ / _ \
38 / _ ( _ ) _ _ _ _ _ _ _ _ _ _ / _ _ _ _ _
39 / _ / / _ _ ' / - _ ) _ _ \ / / _ _ ' / / / / - _ )
40 / _ _ \ _ \ , / \ _ / / / _ _ \ _ , _ / \ _ , _ / \ _ /
41 / _ _ \
42 """)
43
44     matrix = input_squareMatrix()
45     matrix = np.array(matrix, dtype=float)
46
47     eig, x = power_method(matrix, iter=2000, tolerance=1e-9)
48
49     if eig is None:
50         print("Power method did not converge to a unique dominant eigenvalue.")
51     else:
52         print(f"Dominant eigenvalue: {eig}")
53         print(f"Dominant eigenvector: {x}")
54     print("Press Enter to go back to main menu...")
55     input()
56     mainMenu()

```

Explanation The `dominantEigenvalue()` function computes the **dominant eigenvalue** and its corresponding **eigenvector** of a given square matrix using the **Power Method**, a simple iterative numerical algorithm.

- The function first clears the screen and displays an ASCII title for presentation purposes.
- It then calls `input_squareMatrix()` to allow the user to input a square matrix, which is converted into a NumPy array for computation.
- The core logic is handled by the helper function `power_method(A, iter, tolerance)`, which performs the iterative computation:
 - The algorithm begins with an initial guess vector x (initialized as a vector of ones) and normalizes it.
 - In each iteration, the algorithm multiplies the matrix A by the current vector x :

$$Ax = Ax$$

- The resulting vector is normalized to produce the next approximation \mathbf{x}_{new} .
- The eigenvalue estimate is updated using the **Rayleigh quotient**:

$$\lambda_{\text{new}} = \frac{\mathbf{x}_{\text{new}}^T (A \mathbf{x}_{\text{new}})}{\mathbf{x}_{\text{new}}^T \mathbf{x}_{\text{new}}}$$

- Convergence is checked by comparing the change in eigenvalue estimates between iterations. If

$$|\lambda_{\text{new}} - \lambda_{\text{old}}| < \text{tolerance},$$

the algorithm stops and returns the approximate dominant eigenvalue and eigenvector.

- The function then prints the computed dominant eigenvalue and corresponding normalized eigenvector.
- Finally, it waits for user input before returning to the main menu.

Output

```

---
/  - \---  - - -  (-)--  ---  ---  /  / -
/  // / - \ / , \ / / - \ / - \ / - \ / --/
/-----/\-----/ / - / / - / / - / \ - , - / - / \ --/
/  --(-)--  ---  ---  ---  ---  /  / - -----
/  _// / - \ / - -) - \ | / / - \ / / // / - -)
/  --- / - \ - , / \ - / - / - / \ - , - / \ - , - / \ --/
      / --- /

Enter the size of the square matrix (n x n): 3

Enter the 3x3 matrix values row by row (separated by spaces):
Row 1: 2 4 5
Row 2: 5 3 4
Row 3: 1 2 3
Dominant eigenvalue: 9.219264347529903
Dominant eigenvector: [0.62158879 0.7110826 0.32861668]
Press Enter to go back to main menu...

```

Output (Repeated Eigenvalues)

```

---
/  - \---  - - -  (-)--  ---  ---  /  / -
/  // / - \ / , \ / / - \ / - \ / - \ / --/
/-----/\-----/ / - / / - / / - / \ - , - / - / \ --/
/  --(-)--  ---  ---  ---  ---  /  / - -----
/  _// / - \ / - -) - \ | / / - \ / / // / - -)
/  --- / - \ - , / \ - / - / - / \ - , - / \ - , - / \ --/
      / --- /

Enter the size of the square matrix (n x n): 2

Enter the 2x2 matrix values row by row (separated by spaces):
Row 1: 2 1
Row 2: 0 2
Power method did not converge to a unique dominant eigenvalue.
Press Enter to go back to main menu...

```

Output (Non-Dominant Eigenvalues)

```

---
/  - \---  - - -  (-)--  ---  ---  /  / -
/  // / - \ / , \ / / - \ / - \ / - \ / --/
/-----/\-----/ / - / / - / / - / \ - , - / - / \ --/
/  --(-)--  ---  ---  ---  ---  /  / - -----
/  _// / - \ / - -) - \ | / / - \ / / // / - -)
/  --- / - \ - , / \ - / - / - / \ - , - / \ - , - / \ --/
      / --- /

Enter the size of the square matrix (n x n): 2

```

```

Enter the 2x2 matrix values row by row (separated by spaces):
Row 1: 0 -1
Row 2: 1 0
Power method did not converge to a unique dominant eigenvalue.
Press Enter to go back to main menu...

```

3.4 Singular Value Decomposition

Theory The **Singular Value Decomposition (SVD)** is a powerful matrix factorization technique that generalizes the concept of eigen-decomposition to any real or complex matrix $A \in \mathbb{R}^{m \times n}$, whether square or rectangular. It expresses A as the product of three matrices:

$$A = U \Sigma V^T$$

where:

- $U \in \mathbb{R}^{m \times m}$ is an **orthogonal matrix** whose columns are the **left singular vectors** of A .
- $\Sigma \in \mathbb{R}^{m \times n}$ is a **diagonal matrix** containing the **singular values** $\sigma_1, \sigma_2, \dots, \sigma_r$ (where $r = \text{rank}(A)$).
- $V \in \mathbb{R}^{n \times n}$ is an **orthogonal matrix** whose columns are the **right singular vectors** of A .

Mathematical Background

The singular values of A are defined as the non-negative square roots of the eigenvalues of $A^T A$ (or equivalently AA^T):

$$A^T A v_i = \sigma_i^2 v_i \quad \text{and} \quad AA^T u_i = \sigma_i^2 u_i$$

Here, v_i and u_i are the right and left singular vectors corresponding to the singular value σ_i , respectively. The singular values are typically arranged in descending order:

$$\sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_r > 0$$

Each singular value measures the amount of “stretching” that A applies in the direction of its corresponding singular vector.

Geometric Interpretation

The SVD represents the linear transformation A as a sequence of three simpler transformations:

$$A = U \Sigma V^T$$

$$(\text{Rotation by } V^T) \longrightarrow (\text{Scaling by } \Sigma) \longrightarrow (\text{Rotation by } U)$$

This means that A first rotates the input space, then scales it along orthogonal axes (by the singular values), and finally rotates it again.

Applications

SVD has a wide range of applications across mathematics, engineering, and data science:

- **Dimensionality Reduction:** Used in **Principal Component Analysis (PCA)** to reduce large datasets into smaller, more informative components.
- **Image Compression:** Images represented as matrices can be approximated using only the largest singular values, reducing storage while preserving quality.
- **Noise Filtering:** Small singular values often correspond to noise and can be truncated to enhance signal clarity.
- **Solving Linear Systems:** Provides stable solutions for ill-conditioned or non-square systems using the pseudoinverse:

$$A^+ = V \Sigma^+ U^T$$

- **Data Analysis and Recommender Systems:** Helps identify latent structures or patterns (e.g., in collaborative filtering like the Netflix algorithm).

```

1 def svd():
2     clear()
3     print(r"""
4     ,-----,
5     '-\ \ \ \ \ / | \ \ \ \

```

```

6  """
7  .-.-.-.-.-. | \ / | | \ / :
8  -.-.-.-.-. | \ / | | \ / :
9  """)
10 np.set_printoptions(precision=4, suppress=True)
11 A = np.array(input_matrix()) #A = U VT
12 m, n = A.shape
13 r = min(m, n)
14
15 ATA = A.T @ A
16 lamV, V = np.linalg.eigh(ATA)
17 idxV = lamV.argsort()[::-1]
18 lamV = lamV[idxV]
19 V = V[:, idxV]
20
21 #lamU = np.clip(lamU, 0.0, None)
22 lamV = np.clip(lamV, 0.0, None)
23 sigma = np.sqrt(lamV) #averaging in case of mismatch due to floating point rounding errors
24
25 eps = np.finfo(float).eps
26 tol = max(m, n) * eps * (sigma[0] if r else 0.0)
27 k = int(np.sum(sigma > tol)) # numerical rank
28 k = min(k, r)
29
30 U = np.zeros((m, m))
31 U[:, :k] = (A @ V[:, :k]) / sigma[:k].reshape(1, -1)
32
33 d = np.sign(np.diag(U[:, :k].T @ A @ V[:, :k]))
34 d[d == 0] = 1.0
35 U[:, :k] *= d.reshape(1, -1)
36 V[:, :k] *= d.reshape(1, -1)
37
38 if m > k:
39     Z = np.random.randn(m, m - k)
40     Z -= U[:, :k] @ (U[:, :k].T @ Z)
41     U2, _ = np.linalg.qr(Z)
42     U[:, k:] = U2
43
44 Sigma = np.zeros((m, n))
45 Sigma[np.arange(r), np.arange(r)] = sigma[:r]
46
47 def zapsmall(M, Aref=None, rel=50*np.finfo(float).eps, abs_=0.0):
48     scale = np.linalg.norm(A if Aref is None else Aref, ord=np.inf)
49     thr = abs_ + rel * scale
50     M[np.abs(M) < thr] = 0.0
51     return M
52
53 U = zapsmall(U, A)
54 V = zapsmall(V, A)
55 Sigma = zapsmall(Sigma, A)
56
57 print(f"\n\nU = {U};\n\n = {Sigma};\n\nV_T = {V.T}")
58 print(f"Check\nA = U V_T = {zapsmall(U @ Sigma @ V.T, A)};")
59 print("Press Enter to go back to main menu...")
60 input()
61 mainMenu()

```

Explanation The function `svd()` performs the **Singular Value Decomposition (SVD)** of a user-input matrix A , decomposing it into three matrices U , Σ , and V^T , such that:

This implementation computes the SVD manually using NumPy operations, providing detailed insight into the mathematical steps behind the decomposition process. Code snippets are presented before the explanation.

```
clear()
print(r"""
      .-..-.   .-..-.
     /  \_/_\  /  \_/_\
    /  _\_/_\  /  _\_/_\
   /  _\_/_\  /  _\_/_\
  /  _\_/_\  /  _\_/_\
 /  _\_/_\  /  _\_/_\
/  _\_/_\  /  _\_/_\
""")
```

- The function starts by clearing the screen and displaying an ASCII banner for presentation.

```
np.set_printoptions(precision=4, suppress=True)
A = np.array(input_matrix()) #A = U VT
m, n = A.shape
r = min(m, n)
```

- The decimal precision is set to four digits, to make results more comprehensible.

The user inputs a matrix using the `input_matrix()` function, which is then stored as a NumPy array `A`.

The matrix dimensions m and n are determined from the shape of the input matrix, and the variable $r = \min(m, n)$ represents the smaller dimension (the rank limit).

```
ATA = A.T @ A
lamV, V = np.linalg.eigh(ATA)
idxV = lamV.argsort()[::-1]
lamV = lamV[idxV]
V = V[:, idxV]
```

- Next, the function computes $A^T A$, which is a symmetric matrix, using the `np.linalg.eigh()` function, which is specially designed for hermitian/symmetric matrices. Its eigenvalues and eigenvectors are calculated using:

$$(A^\top A)V = V\Lambda$$

The eigenvalues in Λ correspond to the squares of the singular values, and the eigenvectors form the columns of V (the **right singular vectors**).

- The eigenvalues (λ_i) are sorted in descending order, and any negative rounding errors are corrected using:

$$\lambda_i = \max(\lambda_i, 0)$$

The singular values (σ_i) are then obtained as:

$$\sigma_i = \sqrt{\lambda_i}$$

```
lamV = np.clip(lamV, 0.0, None)
sigma = np.sqrt(lamV)
```

- Mathematically $\lambda_i \geq 0$ and this correction wouldn't be necessary, however in the implementation, this is done to correct any floating-point round-off errors that may produce negative, yet practically zero, values, that indicate that the value is probably meant to be exactly zero.

```
eps = np.finfo(float).eps
tol = max(m, n) * eps * (sigma[0] if r else 0.0)
k = int(np.sum(sigma > tol)) # numerical rank
k = min(k, r)
```

6. A tolerance check is applied to determine the **numerical rank** of the matrix (handling floating-point precision issues), and QR decomposition is used to complete the orthonormal basis of U if necessary (for cases where $m > n$).

```

1 U = np.zeros((m, m))
2 U[:, :k] = (A @ V[:, :k]) / sigma[:k].reshape(1, -1)

```

7. To find the **left singular vectors** (U), the function uses the relationship:

$$U_i = \frac{AV_i}{\sigma_i}$$

This ensures that each column of U is orthonormal and corresponds to a singular direction of A . We do not compute U and the corresponding eigenvalues separately using `np.linalg.eigh()` because it would most probably generate a set of eigenvectors that, though valid, do not align with the equality $U_i = \frac{AV_i}{\sigma_i}$.

```

1 d = np.sign(np.diag(U[:, :k].T @ A @ V[:, :k]))
2 d[d == 0] = 1.0
3 U[:, :k] *= d.reshape(1, -1)
4 V[:, :k] *= d.reshape(1, -1)

```

8. This step is also unnecessary in pure mathematics. We know that

$$Av_i = \sigma_i u_i, \quad A^\top u_i = \sigma_i v_i$$

if we take any triplet (u_i, σ_i, v_i) and replace **both** u_i and v_i with their negatives, the equalities still hold. In the implementation, the problem again lies in round-off errors that can flip either one of u_i or v_i within the triple, but not the other. The code above fixes this, ensuring their signs uphold the equality

$$u_i = \frac{Av_i}{\sigma_i}$$

specifically, since in our implementation, U is derived from V .

```

1 if m > k:
2     Z = np.random.randn(m, m - k)
3     Z -= U[:, :k] @ (U[:, :k].T @ Z)
4     U2, _ = np.linalg.qr(Z)
5     U[:, k:] = U2

```

9. We have previously calculated the numerical rank k , which means we only have k values (u_i, σ_i, v_i) that are *true*, i.e., not as a result of round-off errors. So now

$$U \in \mathbb{R}^{m \times k}, \quad \Sigma \in \mathbb{R}^{k \times k}, \quad V \in \mathbb{R}^{n \times k}$$

, but we want to construct the full matrices

$$U \in \mathbb{R}^{m \times m}, \quad \Sigma \in \mathbb{R}^{m \times n}, \quad V \in \mathbb{R}^{n \times n}$$

This is precisely what the code snippet above does.

```

1 Sigma = np.zeros((m, n))
2 Sigma[np.arange(r), np.arange(r)] = sigma[:r]

```

10. The diagonal matrix Σ is then constructed by placing the singular values along its diagonal:

$$\Sigma = \begin{bmatrix} \sigma_1 & 0 & \cdots \\ 0 & \sigma_2 & \cdots \\ \vdots & \vdots & \ddots \end{bmatrix}$$

```

1 def zapsmall(M, Aref=None, rel=50*np.finfo(float).eps, abs_=0.0):
2     scale = np.linalg.norm(A if Aref is None else Aref, ord=np.inf)
3     thr = abs_ + rel * scale
4     M[np.abs(M) < thr] = 0.0
5     return M
6
7 U = zapsmall(U, A)
8 V = zapsmall(V, A)
9 Sigma = zapsmall(Sigma, A)

```

11. The helper function `zapsmall()` removes very small numerical values (near zero) to clean up floating-point noise and improve readability of the output.

```

1 print(f"\n\nU = {U};\n\nΣ = {Sigma};\n\nV.T = {V.T}")
2 print(f"Check\nA = U V.T = {zapsmall(U @ Sigma @ V.T, A)};")
3 print("Press Enter to go back to main menu...")
4 input()
5 mainMenu()

```

12. Finally, the program prints the resulting matrices U , Σ , and V^T , and verifies the decomposition by reconstructing A through:

$$A \approx U\Sigma V^T$$

This check confirms the correctness of the SVD.

Output



```

Enter number of rows: 2
Enter number of columns: 3

```

```

Enter the matrix values row by row (separated by spaces):
Row 1: 1 2 3
Row 2: 4 5 6

```

```

U = [[-0.3863  0.9224]
      [-0.9224 -0.3863]]

```

```

Σ = [[9.508  0.  0. ]
      [0.  0.7729 0. ]

```

```

V.T = [[-0.4287 -0.5663 -0.7039]
        [-0.806  -0.1124  0.5812]
        [ 0.4082 -0.8165  0.4082]]

```

```

Check

```

```

A = U Σ V.T = [[1.  2.  3.]
                [4.  5.  6.]]

```

```

Press Enter to go back to main menu...

```

Output (Zero Singular Values)



```

Enter number of rows: 2
Enter number of columns: 3

```


Enter the matrix values row by row (separated by spaces):

Row 1: 1 2 3

Row 2: 2 4 6

$U = \begin{bmatrix} 0.4472 & 0. & . \\ 0.8944 & 0. & . \end{bmatrix};$

$\Sigma = \begin{bmatrix} 8.3666 & 0. & 0. & . \\ 0. & 0. & 0. & . \end{bmatrix};$

$V_T = \begin{bmatrix} 0.2673 & 0.5345 & 0.8018 \\ -0.9554 & 0.0386 & 0.2927 \\ -0.1255 & 0.8443 & -0.521 \end{bmatrix}$

Check

$A = U \Sigma V_T = \begin{bmatrix} 1. & 2. & 3. \\ 2. & 4. & 6. \end{bmatrix};$

Press Enter to go back to main menu...