



Department of Information Technology A.Y. 2024-2025

Class: TY BTech-IT, Semester: VI Subject: Big Data Lab

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Experiment – 10

1. **Aim:** To implement SON/CUR algorithm.

CODE:

SON:

```
import numpy as np
```

```
def son(A, num_columns):
```

```
    """
```

```
    Perform SON (Subsampled Orthogonalization and Normalization)
```

```
    Args:
```

```
        A: Input matrix (m x n)
```

```
        num_columns: Number of columns to select
```

```
    Returns:
```

```
        A matrix C (m x num_columns) representing selected columns
```

```
    """
```

```
        # Step 1: Subsample the columns randomly
```

```
        m, n = A.shape
```

```
        selected_columns = np.random.choice(n, num_columns, replace=False)
```

```
        C = A[:, selected_columns]
```

```
        # Step 2: Orthogonalization (using Gram-Schmidt or QR factorization)
```

```
        Q, R = np.linalg.qr(C)
```

```
        # Step 3: Normalize the selected columns
```

```
        C_normalized = Q
```

```
    return C_normalized
```

```
# Example usage:
```

```
A = np.array([[1, 2, 3, 4],
```

```
              [5, 6, 7, 8],
```

```
              [9, 10, 11, 12]])
```

```
C_son = son(A, num_columns=2) print("SON  
Resulting Columns:\n", C_son)
```

CUR:

```
def cur_decomposition(A, num_columns, num_rows):  
    """  
    Perform CUR Decomposition  
    Args:  
        A: Input matrix (m x n)  
        num_columns: Number of columns to select  
        num_rows: Number of rows to select    Returns:  
        C: Submatrix of selected columns  
        U: Middle matrix that approximates relationships  
        R: Submatrix of selected rows  
    """    m, n =  
A.shape  
  
    # Step 1: Select columns randomly (subsampling)  
    selected_columns = np.random.choice(n, num_columns, replace=False)  
    C = A[:, selected_columns]  
  
    # Step 2: Select rows randomly (subsampling)  
    selected_rows = np.random.choice(m, num_rows, replace=False)  
    R = A[selected_rows, :]  
  
    # Step 3: Calculate U using the pseudo-inverse of C and R  
    U = np.linalg.pinv(C) @ A @ np.linalg.pinv(R)  
  
    return C, U, R  
  
# Example usage:  
A = np.array([[1, 2, 3, 4],  
              [5, 6, 7, 8],  
              [9, 10, 11, 12]])  
  
C_cur, U_cur, R_cur = cur_decomposition(A, num_columns=2, num_rows=2)  
A_approx = C_cur @ U_cur @ R_cur  
  
print("Original Matrix A:\n", A) print("CUR  
Decomposition - C:\n", C_cur) print("CUR  
Decomposition - U:\n", U_cur) print("CUR  
Decomposition - R:\n", R_cur) print("CUR  
Approximation of A:\n", A_approx)
```

2. Requirements: PC, Internet

OUTPUT: SON:

SON Algorithm (Subsampled Orthogonalization and Normalization):

- Resulting Columns (C):

$$\begin{bmatrix} -0.0967 & 0.9077 \\ -0.4834 & 0.3157 \\ -0.8701 & -0.2763 \end{bmatrix}$$

These are the **orthogonalized and normalized columns** selected from the matrix A .

CUR:

CUR Decomposition:

- Selected Columns (C):

$$\begin{bmatrix} 4 & 3 \\ 8 & 7 \\ 12 & 11 \end{bmatrix}$$

- Middle Matrix (U):

$$\begin{bmatrix} 1.375 & -0.375 \\ -1.5 & 0.5 \end{bmatrix}$$

- Selected Rows (R):

$$\begin{bmatrix} 1 & 2 & 3 & 4 \\ 9 & 10 & 11 & 12 \end{bmatrix}$$

- CUR Approximation of A :

$$\begin{bmatrix} 1. & 2. & 3. & 4. \\ 5. & 6. & 7. & 8. \\ 9. & 10. & 11. & 12. \end{bmatrix}$$

3. Conclusion:

Thus, in this experiment, we implemented SON/CURE algorithms.