COLUMN SORTING IN PYGSVD

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1. Introduction

This document is intended to document and clarify the details of column ordering for the Python pygsvd library. For convenience, an excerpt from the LAPACK documentation is reproduced in the next section.

2. LAPACK DOCUMENTATION EXCERPT

The generalized singular value decomposition of an $m \times m$ matrix A and a $p \times n$ matrix B is given by the pair of factorizations

$$A = U\Sigma_1[0, R]Q^T$$
 and $B = V\Sigma_2[0, R]Q^T$

The matrices in these factorizations have the following properties:

- (1) U is $m \times m$, V is $p \times p$, Q is $n \times n$ and all three matrices are orthogonal. If A and B are complex, these matrices are unitary instead of orthogonal, and Q^T should be replaced by Q^H in the pair of factorizations.
- (2) R is $r \times r$, upper triangular and nonsingular. [0, R] is $r \times n$ (in other words, the 0 is an $r \times (n-r)$ zero matrix. The integer r is the rank of $\begin{pmatrix} A \\ B \end{pmatrix}$, and satisfies $r \leq n$.
- (3) Σ_1 is $m \times r$, Σ_2 is $p \times r$, both are real, nonnegative and diagonal, and $\Sigma_1^T \Sigma_1 + \Sigma_2^T \Sigma_2 = I$. Write $\Sigma_1^T \Sigma_1 = \text{diag } (\alpha_1^2, \dots, \alpha_r^2)$ and $\Sigma_2^T \Sigma_2 = \text{diag } (\beta_1^2, \dots, \beta_r^2)$, where α_i and β_i lie in the interval from 0 to 1. The ratios $\alpha_1/\beta_1, \dots, \alpha_r/\beta_r$ are called the **generalized singular values of the pair** A, B. If $\beta_i = 0$, then the generalized singular value α_i/β_i is infinite

 Σ_1 and Σ_2 have the following detailed structures, depending on whether $m-r \geq 0$ or m-r < 0. In the first case, $m-r \geq 0$, then

$$\Sigma_1 = \begin{pmatrix} k & l \\ k \begin{pmatrix} I & 0 \\ 0 & C \\ m - k - l \end{pmatrix} \quad \text{and} \quad \Sigma_2 = \begin{pmatrix} k & l \\ 0 & S \\ p - l \begin{pmatrix} 0 & S \\ 0 & 0 \end{pmatrix}$$

Here l is the rank of B, k = r - l, C and S are diagonal matrices satisfying $C^2 + S^2 = I$, and S is nonsingular. We may also identify $\alpha_1 = \cdots = \alpha_k = 1$, $\alpha_{k+i} = c_{ii}$ for $i = 1, \ldots, l$, $\beta_1 = \cdots = \beta_k = 0$, and $\beta_{k+i} = s_{ii}$ for $i = 1, \ldots, l$. Thus the first k generalized singular values $\alpha_1/\beta_1, \ldots, \alpha_k/\beta_k$ are infinite, and the remaining l generalized singular values are finite.

In the second case, when m-r<0,

$$\Sigma_{1} = \begin{pmatrix} k & m-k & k+l-m \\ k & 0 & 0 \\ m-k & 0 & C & 0 \end{pmatrix} \quad \text{and} \quad \Sigma_{2} = \begin{pmatrix} k & m-k & k+l-m \\ m-k & 0 & S & 0 \\ 0 & 0 & I \\ p-l & 0 & 0 & 0 \end{pmatrix}$$

Again, l is the rank of B, k = r - l, C and S are diagonal matrices satisfying $C^2 + S^2 = I$, S is nonsingular, and we may identify $\alpha_1 = \cdots = \alpha_k = 1, \alpha_{k+i} = c_{ii}$ for $i = 1, \ldots, m - k, \alpha_{m+1} = \cdots = \alpha_r = 0, \beta_1 = \cdots = \beta_k = 0$, and $\beta_{k+i} = s_{ii}$ for $i = 1, \ldots, m - k$, and $\beta_{m+1} = \cdots = \beta_r = 1$. Thus the first k generalized singular values $\alpha_1/\beta_1, \ldots, \alpha_k/\beta_k$ are infinite, and the remaining l generalized singular values are finite.

3. Column Sorting Performed in Pygsvd

The GSVD implementation in LAPACK returns singular values in arrays of length n. But in all cases, we are only interested in the first r values. These are precisely the α_i and β_i , $i = 1, \ldots r$ values described in the two cases above.

The problem addressed here is that although α_i corresponds to β_i for each i and the columns of U, V and X are ordered correspondingly, no absolute ordering is guaranteed by the LAPACK algorithm. The authors of pygsvd have chosen to provide the α values in descending order and the β values in ascending order, resulting in generalized singular values in descending order, which corresponds to the conventional ordering for the standard SVD algorithm.

As can be seen in the structures of Σ_1 and Σ_2 above, the non-trivial α and β values that require sorting begin at index i = k+1 and continue to either i = k+l in the first case or i = k+m-k=m in the second case. When the sort operation is limited to this minimal set, we can guarantee that no indexing error will occur during sorting of columns of U, V or X.

We next develop a strategy for re-ordering columns to accomplish the desired sorting. We note that if X is defined by

$$X = Q \left(\begin{array}{cc} I & 0 \\ 0 & R^{-1} \end{array} \right)$$

That is, if X is the inverse transpose of the default X returned by pygsvd, then

$$U^TAX = U^TU\Sigma_1 \left(\begin{array}{cc} 0 & R \end{array} \right) Q^TQ \left(\begin{array}{cc} I & 0 \\ 0 & R^{-1} \end{array} \right) = \Sigma_1 \left(\begin{array}{cc} 0 & R \end{array} \right) \left(\begin{array}{cc} I & 0 \\ 0 & R^{-1} \end{array} \right) = \Sigma_1 \left(\begin{array}{cc} 0 & I \end{array} \right)$$

Similarly

$$V^T B X = \Sigma_2 \left(\begin{array}{cc} 0 & I \end{array} \right)$$

Thus we see that if r < n, we have an additional n-r columns of zeros in these products preceding the structures of Σ_1 and Σ_2 . In the first case, $m-r \ge 0$, we get

$$U^{T}AX = \begin{pmatrix} n-r & k & l \\ k & 0 & I & 0 \\ l & 0 & 0 & C \\ m-k-l & 0 & 0 & 0 \end{pmatrix} \quad \text{and} \quad V^{T}BX = \begin{pmatrix} n-r & k & l \\ l & 0 & 0 & S \\ p-l & 0 & 0 & 0 \end{pmatrix}$$

In the second case, when m-r<0,

$$U^{T}AX = \begin{pmatrix} n-r & k & m-k & k+l-m \\ k & 0 & I & 0 & 0 \\ m-k & 0 & C & 0 \end{pmatrix} \quad \text{and} \quad$$

$$V^{T}BX = k + l - m \begin{pmatrix} 0 & 0 & S & 0 \\ k + l - m & 0 & 0 & I \\ p - l & 0 & 0 & 0 & 0 \end{pmatrix}$$

To see clearly how columns must be reordered, it is useful to write U^TAX as

$$\begin{pmatrix} u_1^T \\ u_1^T \\ \vdots \\ u_m^T \end{pmatrix} A \begin{pmatrix} x_1 & x_2 & \cdots & x_n \end{pmatrix} = \begin{pmatrix} u_1^T A x_1 & u_1^T A x_2 & \cdots & u_1^T A x_n \\ u_2^T A x_1 & u_2^T A x_2 & \cdots & u_2^T A x_n \\ \vdots & \vdots & \vdots & \vdots \\ u_m^T A x_1 & u_m^T A x_2 & \cdots & u_m^T A x_n \end{pmatrix}$$

and $V^T B X$ as

$$\begin{pmatrix} v_1^T \\ v_1^T \\ \vdots \\ v_m^T \end{pmatrix} B \begin{pmatrix} x_1 & x_2 & \cdots & x_n \end{pmatrix} = \begin{pmatrix} v_1^T B x_1 & v_1^T B x_2 & \cdots & v_1^T B x_n \\ v_2^T B x_1 & v_2^T B x_2 & \cdots & v_2^T B x_n \\ \vdots & \vdots & \vdots & \vdots \\ v_p^T B x_1 & v_p^T B x_2 & \cdots & v_p^T B x_n \end{pmatrix}$$

Pre-multiplication of a square matrix A by a compatible permutation matrix P^T reorders the rows of A and post-multiplication of A by P reorders the corresponding columns. So in particular, if D is a diagonal matrix, P^TDP will re-order the diagonal elements of D. In the case, for example where U, A and X are $n \times n$, and r = l = n we could let $\tilde{U} = UP$, $\tilde{X} = XP$ and get

$$\tilde{U}^T A \tilde{X} = (UP)^T A X P = P^T U^T A X P = P^T \Sigma_1 P$$

So we see that the desired ordering of the diagonal entries can be achieved by an appropriate ordering of the columns of U and A. Now consider the two representations of U^TAX above. In the case m-r>0. We wish to permute columns of U and columns of X in such a way that the diagonal entries of C are in descending order. The top left element of C corresponds to $u_{k+1}^TAx_{n-r+k+1}$. The desired entries of C can be sorted performing the desired sort on the correct ranges of columns of U and X. For U, we see that we must skip the first k columns and sort the next l columns. For X, we must skip the first n-r+k columns and sort the next l columns. Since X always has n columns, this is equivalent to sorting its last l columns. To sort S, we must sort the first l values of V and the last l columns of X, which agrees with the sort already performed on X. These operations correspond to the Python code:

```
if m - r >= 0:
    ix = np.argsort(C[k:r])[::-1] # sort l values
    X[:, -l:] = X[:, -l:][:, ix]
    if compute_uv[0]:
        U[:, k:k+1] = U[:, k:k+1][:, ix]
    if compute_uv[1]:
        V[:, :1] = V[:, :1][:, ix]
```

We also note that the given structures of U^TAX and V^TBX do not place the singular values on the diagonal in cases when the rank r of the stacked AB matrix is less than n and/or in cases when k = r - l > 0, the difference between the rank of AB and the rank of B. In most cases, this can be corrected by rotating the columns of X to the left by n - r and rotating the columns of V down by V. This is impossible only in the case when V0 and V1 and V2 and V3 are the number of rows of V4 is less than the rank of V5. These operations correspond to the Python code:

```
if n-r > 0:
    X = np.roll(X, r-n, axis=1)
if k > 0 and p >= r:
    V = np.roll(V, k, axis=1)
```