## Linear Algebra for Deep Learning

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# Symmetric, Orthogonal, and Unitary Matrices

If for a square matrix, A, we have

$$A^{\top} = A$$

then A is said to be a **symmetric matrix**.

If the following is true,

$$AA^{\top} = A^{\top}A = I$$

then A is an **orthogonal matrix**, and  $A^{-1} = A^{\top}$ .

And as a result,  $det(\mathbf{A}) = \pm 1$ .

If we count complex matrices, then if

$$U^*U = UU^* = I$$

we say that U is a **unitary matrix** with  $U^*$  being the conjucate transpose of U.

### **Eigenvalues and Eigenvectors**

Suppose the eigenvalue and eigenvector of matrix A is v and  $\lambda$ , the property

$$Av = \lambda v$$

holds.

The code in Python np.linalg.eig(a) returns two values. np.linalg.eig(a)[0] are the eigen values, and np.linalg.eig(a)[1] are the eigen vectors for the corresponding eigenvalues.

## Vector Norms and Distance Matrix

For an n-dimensional vector, x, we define the p-norm of the vector to be

$$\left\|x\right\|_p = \left(\sum_i \left|x_i\right|^p\right)^{\frac{1}{p}}$$

where p is a real number.

The L2-norm,

$$\left|\left|x\right|\right|_{2} = \sqrt{x_{0}^{2} + x_{1}^{2} + \ldots + x_{n-1}^{2}} = \sqrt{\boldsymbol{x}^{\top}\boldsymbol{x}}$$

and the L1-norm

$$\left|\left|x\right|\right|_1 = \sum_i |x_i|$$

are the most frequently used norms in deep learning.

A funny thing is that the  $L_{\infty}$ -norm finds the maximum absolute value for all the components of x.

Switching from norm to distance makes a change to the equations,

$$\mathrm{L}_p(oldsymbol{x},oldsymbol{y}) = \left(\sum_i \left|x_i - y_i
ight|^p
ight)^{rac{1}{p}}$$

this is called the *Euclidean distance* between two vectors. The L1-distance is often called the *Manhattan distance* ( $\mathbf{L}_1 = \sum_i \left| x_i - y_i \right|$ ).

#### **Covariance Matrix**

The covariance matrix  $\Sigma$  discribes how two columns of data vary together, that is to say

$$\Sigma_{ij} = \frac{1}{n-1} \sum_{k=0}^{n-1} (x_{ki} - \overline{x}_i) \left(x_{kj} - \overline{x}_j\right)$$

Numpy function for calculating the covariance matrix is np.cov(X, rowvar = False). We also

need to set rowvar = False because our data for individual group are stored in the columns.

#### **Mahalanobis Distance**

With the mean value vector  $\mu$  and the covariance matrix  $\Sigma$ , with these values, we can define a distance metric called the *Mahalanobis distance*.

$$D_M = \sqrt{\left(\boldsymbol{x} - \boldsymbol{\mu}\right)^{\top} \boldsymbol{\Sigma}^{-1} (\boldsymbol{x} - \boldsymbol{\mu})}$$

We can use this *Mahalanobis distance* as a simple classifier, called the nearest centroid classifier.  $D_M$  is more accurate than the L1-distance in classifiers.

### **Kullback-Leiber Divergence**

The *Kullback-Leiber divergence* (*KL-divergence*), or *relative entropy*, is a measure of the similarity between two probability distributions: the lower the value, the more similar the distributions.

If P and Q are discrete probability distributions, the KL-divergence is

$$D_{\mathrm{KL}}(P \parallel Q) = \sum_{x} P(x) \mathrm{log}_{2} \bigg( \frac{P(x)}{Q(x)} \bigg)$$

The KL-divergence isn't a distance metrix in mathematical sense because the symmetry property doesn't hold,  $D_{\mathrm{KL}}(P\|Q) \neq D_{\mathrm{KL}}(Q\|P)$ .

## **Principle Component Analysis**

*Principle Component Analysis (PCA)* is the technique to learn the directions of the scatter in the dataset, starting with the direction aligned along the greatest scatter. The PCA algorithm generally involves the following steps:

- 1. Find the mean center of the data.
- 2. Calculate the covariance matrix,  $\Sigma$ , of the mean-centered data.
- 3. Calculate the eigenvalues and the eigenvectors of  $\Sigma$ .
- 4. Sort the eigenvalues by decreasing absolute value.
- 5. Discard the weaker eigenvalues and eigenvectors.
- 6. Generate the new transformed values from the existing dataset,  $oldsymbol{x}' = oldsymbol{W} oldsymbol{x}.$

PCA are often used in machine learning to reduce the model size, thus usually enhancing the results of the deep learning.

## Singlar Value Decomposition and Pseudoinverse

Singular value decomposition (SVD) is a power technique to transform any matrix into the product of three matrices. For example, for an input matrix A, with real elements and shape  $m \times n$ , where m might not be equal to n. Then the SVD for A is

$$A = U\Sigma V^{\top}$$

A have been decomposed into three matrices. A  $m \times m$  orthogonal matrix U, a  $m \times n$  "diagonal" matrix  $\Sigma$  and a  $n \times n$  orthogonal matrix V.

The "sigular" comes from the fact that the diagonal elements of the "diagonal" matrix  $\Sigma$ , are singular values, the square roots of the positive eigenvalues of the matrix  $A^{T}A$ .

In SciPy, the svd function returns three values,  $U, \Sigma$  and  $V^{\top}$ .

We can use the SVD for the PCA, or calculating the pseudoinverse of a matrix.