Linear independence

Linear dependence

A collection or list of n-vectors a_1,\ldots,a_k (with $k\geq 1$) is called *linearly dependent* if

$$\beta_1 a_1 + \dots + \beta_k a_k = 0$$

holds for some β_1, \ldots, β_k that are not all zero. In other words, we can form the zero vector as a linear combination of the vectors, with coefficients that are not all zero.

• Linear dependence of a list of vectors does not depend on the ordering of the vectors in the list.

Linear combinations of linearly independent vectors.

Suppose a vector x is a linear combination of a_1, \ldots, a_k ,

$$x = \beta_1 a_1 + \cdots + \beta_k a_k$$
.

When the vectors a_1, \ldots, a_k are linearly independent, the coefficients that form x are unique: If we also have

$$x = \gamma_1 a_1 + \cdots + \gamma_k a_k$$

then $\beta_i = \gamma_i$ for i = 1, ..., k. This tells us that, in principle at least, we can find the coefficients that form a vector x as a linear combination of linearly independent vectors.

```
In [ ]: import numpy as np
        from scipy.linalg import null_space
        # Define the vectors
        v1 = np.array([1, 2, 3])
        v2 = np.array([4, 5, 6])
        v3 = np.array([7, 8, 9])
        # Stack the vectors into a matrix
        A = np.c_[v1, v2, v3]
        print("Matrix A:")
        print(A)
        # Find the null space of the matrix A
        coefficients = null_space(A)
        print("Null space of the matrix A:")
        print(coefficients)
        # Since the null space returns a basis for the null space, we can scale it
        # to get integer coefficients if needed
        if coefficients.size > 0: # Check if the null space is not empty
            # Get the first non-zero vector from the null space
            coeffs = coefficients[:, 0]
            # Scale to get integer coefficients
            coeffs = np.round(coeffs / np.min(np.abs(coeffs))).astype(int)
            print("Coefficients of linear dependence:", coeffs)
            print("v3 represented as a linear combination of v1 and v2:")
            print(f"v3 = {coeffs[0]} * v1 + {coeffs[1]} * v2")
            print(f"v3 = \{coeffs[0] * v1 + coeffs[1] * v2\}")
            print("The vectors are linearly independent.")
```

Basis

Independence-dimension inequality.

If the n-vectors a_1,\ldots,a_k are linearly independent, then $k\leq n$. In words:

A linearly independent collection of n-vectors can have at most n elements.

Put another way:

Any collection of n+1 or more n-vectors is linearly dependent.

As a very simple example, we can conclude that any three 2-vectors must be linearly dependent. We will prove this fundamental fact below; but first, we describe the concept of basis, which relies on the independence-dimension inequality.

Basis.

A collection of n linearly independent n-vectors (i.e., a collection of linearly independent vectors of the maximum possible size) is called a **basis**.

If the n-vectors a_1,\ldots,a_n are a basis, then any n-vector b can be written as a linear combination of them. To see this, consider the collection of n+1 n-vectors a_1,\ldots,a_n,b . By the independence-dimension inequality, these vectors are linearly dependent, so there are $\beta_1,\ldots,\beta_{n+1}$, not all zero, that satisfy

$$\beta_1 a_1 + \cdots + \beta_n a_n + \beta_{n+1} b = 0$$

If $\beta_{n+1}=0$, then we have

$$\beta_1 a_1 + \cdots + \beta_n a_n = 0$$

which, since a_1, \ldots, a_n are linearly independent, implies that $\beta_1 = \cdots = \beta_n = 0$. But then all the β_i are zero, a contradiction. So we conclude that $\beta_{n+1} \neq 0$. It follows that

$$b = (-\beta_1/\beta_{n+1}) a_1 + \cdots + (-\beta_n/\beta_{n+1}) a_n$$

i.e., b is a linear combination of a_1, \ldots, a_n .

Combining this result with the observation above that any linear combination of linearly independent vectors can be expressed in only one way, we conclude:

Any n-vector b can be written in a unique way as a linear combination of a basis $a_1,\ldots,a_n.$

Expansion in a basis.

When we express an n-vector b as a linear combination of a basis a_1, \ldots, a_n , we refer to

$$b = \alpha_1 a_1 + \cdots + \alpha_n a_n$$

as the expansion of b in the a_1, \ldots, a_n basis. The numbers $\alpha_1, \ldots, \alpha_n$ are called the coefficients of the expansion of b in the basis a_1, \ldots, a_n .

Examples

• The n standard unit n vectors e_1, \ldots, e_n are a basis. Any n-vector b can be written as the linear combination

$$b = b_1e_1 + \cdots + b_ne_n$$

This expansion is unique, which means that there is no other linear combination of e_1, \ldots, e_n that equals b.

• The vectors

$$a_1 = \begin{bmatrix} 1.2 \\ -2.6 \end{bmatrix}, \quad a_2 = \begin{bmatrix} -0.3 \\ -3.7 \end{bmatrix}$$

are a basis. The vector $b=\begin{bmatrix}1\\1\end{bmatrix}$ can be expressed in only one way as a linear combination of them:

$$b = 0.6513a_1 - 0.7280a_2.$$

(The coefficients are given here to 4 significant digits. We will see later how these coefficients can be computed.)

```
In []: a = np.array([1.2, -2.6])
        b = np.array([-0.3, -3.7])
        # show that a and b forms a basis for R^2
        A = np.column stack((a, b))
        print("Matrix A:")
        print(A)
        coefficients = null space(A)
        print("Null space of the matrix A:")
        print(coefficients)
        if coefficients.size == 0:
            print("The vectors are linearly independent.")
        else:
            print("The vectors are linearly dependent.")
        # express the vector [1, 1] as a linear combination of a and b
        v = np.array([1, 1]).reshape(-1, 1)
        coeffs = np.linalg.solve(A, v)
        coeffs = coeffs[:, 0]
        print("Coefficients of linear dependence:")
        print(coeffs)
        print(f"v = \{coeffs[0]:.4f\} * a - \{-coeffs[1]:.4f\} * b")
        print(f"v = \{coeffs[0] * a + coeffs[1] * b\}")
```

Orthonormal vectors

A collection of vectors a_1,\ldots,a_k is orthogonal or mutually orthogonal if $a_i\perp a_j$ for any i,j with $i\neq j,i,j=1,\ldots,k$. A collection of vectors a_1,\ldots,a_k is orthonormal if it is orthogonal and $\|a_i\|=1$ for $i=1,\ldots,k$.

- A vector of norm one is called normalized; dividing a vector by its norm is called normalizing it.
- Thus, each vector in an orthonormal collection of vectors is normalized, and two different vectors from the collection are orthogonal.

These two conditions can be combined into one statement about the inner products of pairs of vectors in the collection: a_1, \ldots, a_k is orthonormal means that

$$a_i^T a_j = \left\{egin{array}{ll} 1 & i=j \ 0 & i
eq j \end{array}
ight..$$

- Orthonormality, like linear dependence and independence, is an attribute of a collection of vectors, and not an attribute of vectors individually.
- By convention, though, we say "The vectors a_1, \ldots, a_k are orthonormal" to mean "The collection of vectors a_1, \ldots, a_k is orthonormal".

```
In [ ]: vectors = {
            "a1": np.array([0, 0, -1]),
            "a2": np.array([1, 1, 0]) / np.sqrt(2),
            "a3": np.array([1, -1, 0]) / np.sqrt(2),
        for name, vector in vectors.items():
            print(f"norm of {name}: {np.linalg.norm(vector)}")
        for name1, vector1 in vectors.items():
            for name2, vector2 in vectors.items():
                if name1 != name2:
                    print(f"inner product of {name1} and {name2}: {np.dot(vector1, vector2)}"
In []: x = np.array([1, 2, 3])
        betas = \{\}
        for name, vector in vectors.items():
            betas[name] = np.dot(vector, x)
            print(f"beta_{name}: {betas[name]}")
        print(f"betas: {betas}")
        xexp = sum([betas[name] * vectors[name] for name in vectors.keys()])
        print(f"expansion: {xexp}")
        print(f"original: {x}")
```

Gram-Schmidt algorithm

```
Algorithm: given n-vectors a_1, \ldots, a_k for i = 1, \ldots, k,
```

- 1. Orthogonalization. $ilde{q}_i = a_i \left(q_1^T a_i
 ight)q_1 \dots \left(q_{i-1}^T a_i
 ight)q_{i-1}$
- 2. Test for linear dependence. if $\tilde{q}_i = 0$, quit.
- 3. Normalization. $q_i = ilde{q}_i / \| ilde{q}_i\|$

```
In [5]: def gram_schmidt(a, tol=le-10):
    q = []
    for i in range(len(a)):
        qtilde = a[i]
        for j in range(i):
              qtilde = qtilde - np.inner(q[j], a[i]) * q[j]
        if np.linalg.norm(qtilde) < tol:
              print("Vectors linearly dependent")
        q.append(qtilde / np.linalg.norm(qtilde))
    return q</pre>
```

```
In []: a = [[-1, 1, -1, 1], [-1, 3, -1, 3], [1, 3, 5, 7]]
    q = gram_schmidt(a)
    print(f"The orthonormal vectors are:\n{q}")
    # print(np.array(q))
    print(f"Norms of vectors {np.linalg.norm(q, axis=1)}")
```

Suppose that R is a $T \times n$ asset return matrix, that gives the returns of n assets over T periods. A common trading strategy maintains constant investment weights given by the n-vector w over the T periods.

For example, $w_4=0.15$ means that 15% of the total portfolio value is held in asset 4. (Short positions are denoted by negative entries in w.) Then Rw, which is a T-vector, is the time series of the portfolio returns over the periods $1, \ldots, T$.

As an example, consider a portfolio of the 4 assets in table, with weights w=(0.4,0.3,-0.2,0.5).

The product Rw = (0.00213, -0.00201, 0.00241) gives the portfolio returns over the three periods in the example.

	Apple (AAPL)	Google (GOOG)	3M (MMM)	Amazon (AMZN)
March 1, 2016	0.00219	0.00006	-0.00113	0.00202
March 2, 2016	0.00744	-0.00894	-0.00019	-0.00468
March 3, 2016	0.01488	-0.00215	0.00433	-0.00407

Table: Daily returns of Apple (AAPL), Google (GOOG), 3M (MMM), and Amazon (AMZN), on March 1, 2, and 3, 2016 (based on closing prices).

```
In [ ]: # daily returns for 4 assets (AAPL, GOOG, MMM, AMZN)
        base returns = np.array(
                [0.00219, 0.00006, -0.00113, 0.00202],
                [0.00744, -0.00894, -0.00019, -0.00468],
                [0.01488, -0.00215, 0.00433, -0.00407],
            ]
        )
        # Define the investment weights
        w = np.array([0.4, 0.3, -0.2, 0.5]) # Portfolio weights
        # Calculate the portfolio returns Rw
        portfolio_returns = base_returns @ w # Matrix multiplication
        print("Asset Returns:")
        print(base returns)
        print("\nPortfolio Weights:")
        print(w)
        print("\nPortfolio Returns:")
        print(portfolio_returns.round(5))
```

Polynomial evaluation at multiple points. Suppose the entries of the n-vector c are the coefficients of a polynomial p of degree n-1 or less:

$$p(t) = c_1 + c_2 t + \dots + c_{n-1} t^{n-2} + c_n t^{n-1}.$$

Let t_1,\ldots,t_m be m numbers, and define the m-vector y as $y_i=p\left(t_i\right)$. Then we have y=Ac, where A is the m imes n matrix

$$A = egin{bmatrix} 1 & t_1 & \cdots & t_1^{n-2} & t_1^{n-1} \ 1 & t_2 & \cdots & t_2^{n-2} & t_2^{n-1} \ dots & dots & dots & dots \ 1 & t_m & \cdots & t_m^{n-2} & t_m^{n-1} \end{bmatrix}$$

So multiplying a vector c by the matrix A is the same as evaluating a polynomial with coefficients c at m points. The matrix A comes up often, and is called a Vandermonde matrix (of degree n-1, at the points t_1,\ldots,t_m), named for the mathematician Alexandre-Théophile Vandermonde.

```
In [ ]: # Define the polynomial coefficients
        # Example: p(t) = 2 + 3t + t^2 (coefficients: [2, 3, 1])
        coefficients = np.array([2, 3, 1])
        # Define the points at which we want to evaluate the polynomial
        t values = np.array([0, 1, 2, 3, 4, 5, 6])
        # Create the Vandermonde matrix
        # The shape of the matrix will be (m, n) where m is the number of points and n is the
        \# m = len(t_values)
        n = len(coefficients)
        A = np.vander(t_values, N=n, increasing=True)
        print("Vandermonde Matrix A:")
        print(A)
In [ ]: # Evaluate the polynomial at the given points
        y values = A @ coefficients
        y_values
In [ ]: import matplotlib.pyplot as plt
        # Create a range of t values for a smooth curve
        t_curve = np.linspace(t_values[0], t_values[-1], 100)
        y_curve = np.polyval(coefficients[::-1], t_curve) # Reverse coefficients for np.poly
        fig = plt.figure(figsize=(10, 6))
        # Scatter plot of the evaluated points
        plt.scatter(t_values, y_values, color="red", label="Evaluated Points", zorder=5)
        # Plot the polynomial curve
        plt.plot(t_curve, y_curve, label="Polynomial Curve", color="blue")
        plt.title("Polynomial Evaluation using Vandermonde Matrix")
        plt.xlabel("t")
        plt.ylabel("p(t)")
        plt.axhline(0, color="black", linewidth=0.5, ls="--")
        plt.axvline(0, color="black", linewidth=0.5, ls="--")
        plt.grid()
        plt.legend()
        plt.show()
        plt.close(fig)
```

Total price from multiple suppliers. Suppose the $m \times n$ matrix P gives the prices of n goods from m suppliers (or in m different locations). If q is an n-vector of quantities of the n goods (sometimes called a basket of goods), then c = Pq is an N-vector that gives the total cost of the goods, from each of the N suppliers.

```
[105, 205, 155, 310], # Supplier 3 prices
]

# Example quantities for the 4 computer parts
q = np.array([2, 3, 1, 4]) # 2 CPUs, 3 GPUs, 1 Motherboard, 4 RAM sticks

c = P @ q
print("Total costs from each supplier:", c)

supplier_labels = ["Supplier 1", "Supplier 2", "Supplier 3"]
plt.bar(supplier_labels, c, color=["blue", "orange", "green"])
plt.xlabel("Suppliers")
plt.ylabel("Total Cost ($)")
plt.title("Total Cost of Goods from Different Suppliers")
plt.grid(axis="y")
plt.show()
```

Document scoring. Suppose A in an $N \times n$ document-term matrix, which gives the word counts of a corpus of N documents using a dictionary of n words, so the rows of A are the word count vectors for the documents. Suppose that w in an n-vector that gives a set of weights for the words in the dictionary. Then s = Aw is an N-vector that gives the scores of the documents, using the weights and the word counts.

A search engine, for example, might choose w (based on the search query) so that the scores are predictions of relevance of the documents (to the search).

```
In [ ]: # Step 1: Define the documents
        doc1 = """
        Renewable energy sources, such as solar and wind, are becoming increasingly vital in
        The transition from fossil fuels to renewable energy can significantly reduce greenhold
        Solar panels harness sunlight, converting it into electricity, while wind turbines cal
        Governments worldwide are investing in renewable energy infrastructure to promote sus
        Additionally, renewable energy creates jobs in manufacturing, installation, and mainte
        The use of renewable resources can lead to energy independence, reducing reliance on
        As technology advances, the efficiency of renewable energy systems continues to improv
        Public awareness and education about renewable energy are essential for widespread add
        The future of energy lies in sustainable practices that protect our planet.
        doc2 = """
        Artificial Intelligence (AI) is revolutionizing the healthcare industry by enhancing (
        Machine learning algorithms analyze vast amounts of medical data to identify patterns
        AI-powered tools assist doctors in diagnosing diseases earlier and more accurately.
        For instance, AI can analyze medical images to detect anomalies that may be missed by
        Furthermore, AI chatbots provide patients with immediate responses to their inquiries
        The integration of AI in healthcare also streamlines administrative tasks, allowing he
        However, ethical considerations regarding data privacy and algorithmic bias must be a
        As AI technology evolves, its potential to transform healthcare continues to grow.
        Collaboration between technologists and healthcare providers is crucial for successful
        documents = [doc1, doc2]
        # Step 2: Create the document-term matrix (A)
        # For simplicity, let's assume we have a vocabulary of 5 words.
        # Word dictionary: [renewable, energy, sources, AI, healthcare]
        # Initialize the document-term matrix with zeros
```

vocabulary = ["renewable", "energy", "sources", "ai", "healthcare"]

A = np.zeros((len(documents), 5))

Define the vocabulary

```
# Count word occurrences in each document
        for i, doc in enumerate(documents):
            for word in doc.replace("-", " ").lower().split():
                # Remove punctuation and check for word in vocabulary
                word cleaned = word.strip(",.!?()") # Remove punctuation
                if word cleaned in vocabulary:
                    A[i, vocabulary.index(word cleaned)] += 1
        print("Document-Term Matrix (A):")
        print(A)
        # Step 3: Define the weight vector (w)
        # Let's assign weights based on the importance of each word for the query "renewable
        w = np.array([3, 2, 0, 0, 0]) # Higher weight for 'renewable' and 'energy'
        # Step 4: Calculate the document scores (s)
        s = np.dot(A, w)
In [ ]: # Step 5: Visualizations
        # 1. Document-Term Matrix Visualization
        fig = plt.figure(figsize=(10, 4))
        plt.imshow(A, cmap="hot", interpolation="nearest")
        plt.colorbar(label="Word Count")
        plt.title("Document-Term Matrix")
        plt.xlabel("Words")
        plt.ylabel("Documents")
        plt.xticks(ticks=np.arange(5), labels=vocabulary)
        plt.yticks(
            ticks=np.arange(len(documents)),
            labels=[f"Document {i + 1}" for i in range(len(documents))],
        plt.show()
        plt.close(fig)
In [ ]: | # 2. Weight Vector Visualization
        fig = plt.figure(figsize=(6, 4))
        plt.bar(vocabulary, w, color="skyblue")
        plt.title('Weight Vector for Query "Renewable Energy"')
        plt.xlabel("Words")
        plt.ylabel("Weights")
        plt.xticks(rotation=45)
        plt.grid(axis="y")
        plt.show()
        plt.close(fig)
In [ ]: # 3. Score Visualization
        fig = plt.figure(figsize=(6, 4))
        plt.bar([f"Document {i + 1}" for i in range(len(documents))], s, color="lightgreen")
        plt.title("Document Scores Based on Weights")
        plt.xlabel("Documents")
        plt.ylabel("Scores")
        plt.ylim(0, max(s) + 1) # Set y-limit for better visualization
        plt.grid(axis="y")
        plt.show()
In [ ]: | # Step 6: Output Summary
        print("Document Word Counts:")
        for i, doc in enumerate(documents):
            word count = len(doc.split())
            print(f"Document {i + 1} Word Count: {word_count}")
        print("\nDocument Scores:")
        for i, score in enumerate(s):
```

```
print(f"Score for Document {i + 1}: {score}")
```

Audio mixing. Suppose the k columns of A are vectors representing audio signals or tracks of length T, and w is a k-vector. Then b=Aw is a T-vector representing the mix of the audio signals, with track weights given by the vector w.

```
In [20]: import IPython.display as ipd
         # Set parameters
         sample rate = 44100 # Samples per second
         duration = 10 # Duration of each track in seconds
         t = np.linspace(0, duration, int(sample_rate * duration), endpoint=False)
         # Create synthetic audio signals
         # Track 1: Drum beat (simple square wave)
         drum beat = 0.5 * np.sign(np.sin(2 * np.pi * 2 * t)) # 2 Hz square wave
         # Track 2: Guitar riff (sine wave)
         guitar riff = 0.5 * np.sin(2 * np.pi * 440 * t) # (440 Hz)
         # Track 3: Vocal line (sine wave)
         vocal line = 0.5 * np.sin(2 * np.pi * 550 * t) # (550 Hz)
         # Combine tracks into matrix A
         A = np.column_stack((drum_beat, guitar_riff, vocal_line))
         # Define weights for mixing
         w = np.array([0.5, 0.3, 0.2]) # Weights for each track
         # Mix the audio signals
         b = A @ w # Resulting mixed audio signal
         # Normalize the mixed signal to prevent clipping
         b = b / np.max(np.abs(b))
In [ ]: # Play the mixed audio directly
         ipd.Audio(b, rate=sample rate)
In [ ]: # Visualization of waveforms
         plt.figure(figsize=(12, 8))
         # Plot individual tracks
         plt.subplot(4, 1, 1)
         plt.title("Drum Beat")
         plt.plot(t, drum_beat)
         plt.xlim(0, duration)
         plt.ylim(-1, 1)
         plt.subplot(4, 1, 2)
         plt.title("Guitar Riff")
         plt.plot(t, guitar riff)
         plt.xlim(0, duration)
         plt.ylim(-1, 1)
         plt.subplot(4, 1, 3)
         plt.title("Vocal Line")
         plt.plot(t, vocal_line)
         plt.xlim(0, duration)
         plt.ylim(-1, 1)
         # Plot mixed track
         plt.subplot(4, 1, 4)
         plt.title("Mixed Audio Signal")
```

```
plt.plot(t, b)
plt.xlim(0, duration)
plt.ylim(-1, 1)

plt.tight_layout()
plt.show()
```