

15

Mathematical Tools for Multivariate Character Analysis

This chapter introduces a variety of tools from matrix algebra and multivariate statistics useful in the analysis of selection on multiple characters. Our primary intent is to introduce the reader to the idea of vectors and matrices as **geometric structures**, viewing matrix operations as transformations converting a vector into a new vector by a change in geometry (rotation and scaling). The **eigenvalues** and their associated **eigenvectors** of a matrix describe the geometry of the transformation associated with that matrix. Using the multivariate normal, we then develop the multivariate breeders' equation and examine properties of Gaussian fitness functions. We conclude with some elementary concepts in vector calculus, focusing on derivatives of vectors and finding local extrema of vector-valued functions. The reader should be aware that this chapter is rather dense in terms of mathematical machinery and focus on getting an overview of the various methods during the first reading, referring back to relevant sections for specific details as applications arise. Readers who feel a little uncomfortable with matrices might wish to review Chapter 7 before proceeding further.

THE GEOMETRY OF VECTORS AND MATRICES

There are numerous excellent texts on matrix algebra, so we will make little effort to prove most of the results given below. For statistical applications, concise introductions can be found in the chapters on matrix methods in Johnson and Wichern (1988) and Morrison (1976), while Dhrymes (1978) and Searle (1982) provide a more extended treatment. Wilf's (1978) short chapter on matrix methods provides a very nifty review of methods useful in applied mathematics. Franklin (1968), Horn and Johnson (1985), and Gantmacher (1960), respectively, give increasingly sophisticated treatments of matrix analysis.

Comparing Vectors: Lengths and Angles

As Figure 15.1 shows, a vector \mathbf{x} can be treated as a geometric object, an arrow leading from the origin to the n dimensional point whose coordinates are given by the elements of \mathbf{x} . By changing coordinate systems, we change the resulting vector,

potentially changing both its direction (**rotating** the vector) and length (**scaling** the vector). This geometric interpretation suggests several ways for comparing vectors, such as the **angle** between two vectors and the **projection** of one vector onto another.

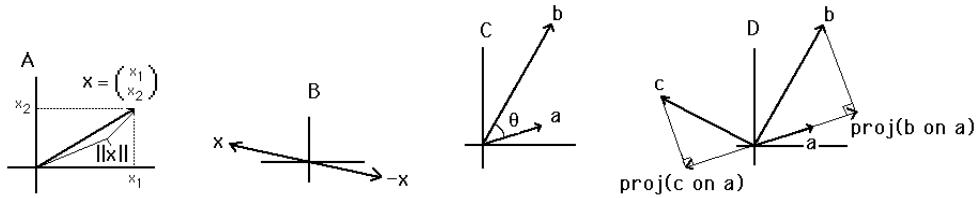


Figure 15.1. Some basic geometric concepts of vectors. While we use examples from two dimensions, these concepts easily extend to n dimensions. **A:** A vector \mathbf{x} can be thought of as an arrow from the origin to a point in space whose coordinates are given by the elements of \mathbf{x} . **B:** Multiplying a vector by -1 results in a *reflection* about the origin. **C:** One measure of the difference in direction between two vectors is the angle θ between them. **D:** $\text{Proj}(\mathbf{b} \text{ on } \mathbf{A})$ is the vector resulting from the projection of \mathbf{b} on \mathbf{A} . Note that the resulting projection vector is either in the same direction as \mathbf{A} or in the direction of the reflection of \mathbf{A} , as seen for $\text{Proj}(\mathbf{c} \text{ on } \mathbf{A})$.

Consider first the length (or **norm**) of a vector. The most common length measure is the Euclidean distance of the vector from the origin, $\|\mathbf{x}\|$, which is defined by

$$\|\mathbf{x}\| = \sqrt{x_1^2 + x_2^2 + \cdots + x_n^2} = \sqrt{\mathbf{x}^T \mathbf{x}} \quad (15.1a)$$

Hence for any scalar a , $\|a\mathbf{x}\| = |a|\|\mathbf{x}\|$. If $a < 0$, the vector $a\mathbf{x}$ is scaled by $|a|$ and reflected about the origin as is shown in Figure 15.1. Similarly, the Euclidean distance between \mathbf{x} and \mathbf{y} is

$$\|\mathbf{x} - \mathbf{y}\|^2 = \sum_{i=1}^n (x_i - y_i)^2 = (\mathbf{x} - \mathbf{y})^T (\mathbf{x} - \mathbf{y}) = (\mathbf{y} - \mathbf{x})^T (\mathbf{y} - \mathbf{x}) \quad (15.1b)$$

Vectors can differ by length, direction, or both. The angle θ between two vectors \mathbf{x} and \mathbf{y} provides a measure of how much they differ in direction (Figure 15.1). If the vectors satisfy $a\mathbf{x} = \mathbf{y}$ where $a > 0$ they point in exactly the same direction, and they are defined to be zero degrees apart. If $a < 0$, they are exactly 180 degrees apart and differ in direction only by a reflection of the axes about the origin. At the other extreme, two vectors can be at right angles to each other ($\theta = 90^\circ$ or 270°), in which case the vectors are said to be **orthogonal**. Orthogonal vectors of unit length are further said to be **orthonormal**. For any two n dimensional vectors, θ

satisfies

$$\cos(\theta) = \frac{\mathbf{x}^T \mathbf{y}}{\|\mathbf{x}\| \|\mathbf{y}\|} = \frac{\mathbf{y}^T \mathbf{x}}{\|\mathbf{x}\| \|\mathbf{y}\|} \tag{15.2}$$

Note that since $\cos(90^\circ) = \cos(270^\circ) = 0$, two vectors are orthogonal if and only if their inner product is zero, $\mathbf{x}^T \mathbf{y} = 0$.

Another way to compare vectors, illustrated in Figure 15.1, is to consider the **projection** of one vector on another. For any two n dimensional vectors, the projection of \mathbf{x} on \mathbf{y} generates a vector defined by

$$\text{Proj}(\mathbf{x} \text{ on } \mathbf{y}) = \frac{\mathbf{x}^T \mathbf{y}}{\mathbf{y}^T \mathbf{y}} \mathbf{y} = \frac{\mathbf{x}^T \mathbf{y}}{\|\mathbf{y}\|^2} \mathbf{y} = \left(\cos(\theta) \frac{\|\mathbf{x}\|}{\|\mathbf{y}\|} \right) \mathbf{y} \tag{15.3 a}$$

If $\|\mathbf{y}\| = 1$, then

$$\text{Proj}(\mathbf{x} \text{ on } \mathbf{y}) = (\mathbf{x}^T \mathbf{y}) \mathbf{y} = (\cos(\theta) \|\mathbf{x}\|) \mathbf{y} \tag{15.3b}$$

Note that since the term involving cosines in Equations 15.3a/b is a scalar, the vector resulting from the projection of \mathbf{x} on \mathbf{y} is in the same direction as \mathbf{y} , unless $90^\circ < \theta < 270^\circ$ in which case $\cos(\theta) < 0$ and the projection vector is in exactly the opposite direction (the reflection of \mathbf{y} about the origin). The length of the projection vector is

$$\|\text{Proj}(\mathbf{x} \text{ on } \mathbf{y})\| = |\cos(\theta)| \|\mathbf{x}\| \tag{15.3c}$$

If two vectors lie in exactly the same direction, the projection of one on the other just recovers the vector ($\text{Proj}(\mathbf{x} \text{ on } \mathbf{y}) = \mathbf{x}$). Conversely, if two vectors are orthogonal, then the projection of one on the other yields a vector of length zero. An important use of projection vectors is that if $\mathbf{y}_1, \mathbf{y}_2, \dots, \mathbf{y}_n$ is any set of mutually orthogonal n dimensional vectors, then any n dimensional vector \mathbf{x} can be represented as the sum of projections of \mathbf{x} onto the members of this set,

$$\mathbf{x} = \sum_{i=1}^n \text{Proj}(\mathbf{x} \text{ on } \mathbf{y}_i) \tag{15.4}$$

Matrices Describe Vector Transformations

When we multiply a vector \mathbf{x} by a matrix \mathbf{A} to create a new vector $\mathbf{y} = \mathbf{A}\mathbf{x}$, \mathbf{A} rotates and scales the original vector \mathbf{x} to give \mathbf{y} . Thus \mathbf{A} describes a transformation of the original coordinate system of \mathbf{x} into a new coordinate system \mathbf{y} (which has a different dimensions than \mathbf{x} unless \mathbf{A} is square).

Orthonormal Matrices

Matrix transformations consist of two basic operations, rotations (changes in the direction of a vector) and scalings (changes in its length). We can partition a matrix transformation into these two basic operations by using **orthonormal** matrices.

Writing a square matrix \mathbf{U} as $\mathbf{U} = (\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_n)$ where each \mathbf{u}_i is an n dimensional column vector, \mathbf{U} is orthonormal if

$$\mathbf{u}_i^T \mathbf{u}_j = \begin{cases} 1 & \text{if } i = j \\ 0 & \text{if } i \neq j \end{cases}$$

In other words, each column of \mathbf{U} is independent from every other column and has unit length. Matrices with this property are also referred to as **unitary**, or **orthogonal** and satisfy

$$\mathbf{U}^T \mathbf{U} = \mathbf{U} \mathbf{U}^T = \mathbf{I} \quad (15.5a)$$

Hence,

$$\mathbf{U}^T = \mathbf{U}^{-1} \quad (15.5b)$$

The coordinate transformation induced by an orthonormal matrix has a very simple geometric interpretation in that it is a **rigid rotation** of the original coordinate system — all axes of the original coordinate are simply rotated by the same angle to create the new coordinate system. To see this, note first that orthonormal matrices preserve all innerproducts. Taking $\mathbf{y}_1 = \mathbf{U}\mathbf{x}_1$ and $\mathbf{y}_2 = \mathbf{U}\mathbf{x}_2$,

$$\mathbf{y}_1^T \mathbf{y}_2 = \mathbf{x}_1^T (\mathbf{U}^T \mathbf{U}) \mathbf{x}_2 = \mathbf{x}_1^T \mathbf{x}_2$$

A special case of this is that orthonormal matrices do not change the length of vectors, as $\|\mathbf{y}_1\| = \mathbf{y}_1^T \mathbf{y}_1 = \mathbf{x}_1^T \mathbf{x}_1 = \|\mathbf{x}_1\|$. If θ is the angle between vectors \mathbf{x}_1 and \mathbf{x}_2 , then following transformation by an orthonormal matrix,

$$\cos(\theta | \mathbf{y}_1, \mathbf{y}_2) = \frac{\mathbf{y}_1^T \mathbf{y}_2}{\sqrt{\|\mathbf{y}_1\|} \sqrt{\|\mathbf{y}_2\|}} = \frac{\mathbf{x}_1^T \mathbf{x}_2}{\sqrt{\|\mathbf{x}_1\|} \sqrt{\|\mathbf{x}_2\|}} = \cos(\theta | \mathbf{x}_1, \mathbf{x}_2)$$

and the angle between any two vectors remains unchanged following their transformation by the same orthonormal matrix.

Eigenvalues and Eigenvectors

The **eigenvalues** and their associated **eigenvectors** of a square matrix describe the geometry of the transformation induced by that matrix. Eigenvalues describe how the original coordinate axes are scaled in the new coordinate system while eigenvectors describe how the original axes are rotated.

Suppose that the vector \mathbf{y} satisfies the matrix equation

$$\mathbf{A}\mathbf{y} = \lambda\mathbf{y} \quad (15.6)$$

for some scalar value λ . Geometrically, this means that the new vector resulting from transformation of \mathbf{y} by \mathbf{A} points in the same direction (or is exactly reflected about the origin if $\lambda < 0$) as \mathbf{y} . For such vectors, the only action of the matrix transformation is to scale them by some amount λ . Hence, it is natural that the

system of such vectors along with their corresponding scalar multipliers completely describes the geometry of the transformation associated with \mathbf{A} . Vectors satisfying Equation 15.6 are referred to as **eigenvectors** and their associated scaling factors are **eigenvalues**. If \mathbf{y} is an eigenvector, then $a\mathbf{y}$ is also an eigenvector as $\mathbf{A}(a\mathbf{y}) = a(\mathbf{A}\mathbf{y}) = \lambda(a\mathbf{y})$. Note, however, that the associated eigenvalue remains unchanged. Hence, we typically scale eigenvectors to be of unit length to give **unit** or **normalized** eigenvectors. In particular, if \mathbf{u}_i is the eigenvector associated with the i th eigenvalue, then the associated normalized eigenvector is $\mathbf{e}_i = \mathbf{u}_i / \|\mathbf{u}_i\|$.

The eigenvalues of square matrix \mathbf{A} of dimension n are solution of Equation 15.6, which is usually expressed as the **characteristic equation** $|\mathbf{A} - \lambda\mathbf{I}| = 0$. This can be also be expressed using the **Laplace expansion** as

$$|\mathbf{A} - \lambda\mathbf{I}| = (-\lambda)^n + S_1(-\lambda)^{n-1} + \dots + S_{n-1}(-\lambda) + S_n = 0 \quad (15.7)$$

where S_i is the sum of all **principal minors** (minors including diagonal elements of the original matrix) of order i . Minors were defined in Chapter 7. Finding the eigenvalues thus requires solving a polynomial equation of order n . In practice, for $n > 2$ this is usually done numerically, and most statistical and numerical analysis packages offer routines to accomplish this task.

Two of these principal minors are easily obtained and provide some information on the nature of the eigenvalues. The only principal minor having the same order of the matrix is the full matrix itself, so that $S_n = |\mathbf{A}|$, the determinant of \mathbf{A} . S_1 is also related to an important matrix quantity, the **trace**. This is denoted by $\text{tr}(\mathbf{A})$ and is the sum of the diagonal elements of the matrix,

$$\text{tr}(\mathbf{A}) = \sum_{i=1}^n a_{ii}$$

Observe that $S_1 = \text{tr}(\mathbf{A})$ as the only principal minors of order one are the diagonal elements themselves, the sum of which equals the trace. The trace and determinant can be expressed as functions of the eigenvalues,

$$\text{tr}(\mathbf{A}) = \sum_{i=1}^n \lambda_i \quad \text{and} \quad |\mathbf{A}| = \prod_{i=1}^n \lambda_i \quad (15.8)$$

Hence \mathbf{A} is singular ($|\mathbf{A}| = 0$) if and only if at least one eigenvalue is zero.

Let \mathbf{e}_i be the (unit-length) eigenvector associated with eigenvalue λ_i . If the eigenvectors of \mathbf{A} can be chosen to be mutually orthogonal, e.g., $\mathbf{e}_i^T \mathbf{e}_j = 0$ for $i \neq j$, we can express \mathbf{A} as

$$\mathbf{A} = \lambda_1 \mathbf{e}_1 \mathbf{e}_1^T + \lambda_2 \mathbf{e}_2 \mathbf{e}_2^T + \dots + \lambda_n \mathbf{e}_n \mathbf{e}_n^T \quad ((15.9b))$$

This decomposition is called the **spectral decomposition** of \mathbf{A} . Hence,

$$\begin{aligned} \mathbf{A}\mathbf{x} &= \lambda_1 \mathbf{e}_1 \mathbf{e}_1^T \mathbf{x} + \lambda_2 \mathbf{e}_2 \mathbf{e}_2^T \mathbf{x} + \dots + \lambda_n \mathbf{e}_n \mathbf{e}_n^T \mathbf{x} \\ &= \lambda_1 \text{Proj}(\mathbf{x} \text{ on } \mathbf{e}_1) + \lambda_2 \text{Proj}(\mathbf{x} \text{ on } \mathbf{e}_2) + \dots + \lambda_n \text{Proj}(\mathbf{x} \text{ on } \mathbf{e}_n) \end{aligned} \quad (15.9b)$$

The last equality follows since $\mathbf{e}_i(\mathbf{e}_i^T \mathbf{x}) = (\mathbf{e}_i^T \mathbf{x})\mathbf{e}_i$ as $\mathbf{e}_i^T \mathbf{x}$ is a scalar.

Properties of Symmetric Matrices

Many of the matrices we will encounter are **symmetric**, satisfying $\mathbf{A} = \mathbf{A}^T$. Examples include variance-covariance matrices and the γ matrix of quadratic coefficients in the Pearson-Lande-Arnold fitness regression. Here we give some of the more useful properties of symmetric matrices. Proofs can be found in Dhrymes (1978), Horn and Johnson (1985), and Wilf (1978).

1. If \mathbf{A} is symmetric, then if \mathbf{A}^{-1} exists, it is also symmetric.
2. The eigenvalues and eigenvectors of a symmetric matrix are all real.
3. For any n -dimensional symmetric matrix, a corresponding set of orthonormal eigenvectors can be constructed, i.e., we can obtain a set of eigenvalues \mathbf{e}_i for $1 \leq i \leq n$ that satisfies

$$\mathbf{e}_i^T \mathbf{e}_j = \begin{cases} 1 & \text{if } i = j \\ 0 & \text{if } i \neq j \end{cases}$$

In particular, this guarantees that a spectral decomposition of \mathbf{A} exists. This can be restated as:

4. A symmetric matrix \mathbf{A} can be **diagonalized** as

$$\mathbf{A} = \mathbf{U}\mathbf{A}\mathbf{U}^T \quad (15.10a)$$

where \mathbf{A} is a diagonal matrix, and \mathbf{U} is an orthonormal matrix ($\mathbf{U}^{-1} = \mathbf{U}^T$). If λ_i and \mathbf{e}_i are the i th eigenvalue and its associated unit eigenvector of \mathbf{A} , then

$$\mathbf{A} = \text{diag}(\lambda_1, \lambda_2, \dots, \lambda_n) = \begin{pmatrix} \lambda_1 & 0 & \dots & 0 \\ 0 & \lambda_2 & \dots & 0 \\ \vdots & & \ddots & \vdots \\ 0 & \dots & \dots & \lambda_n \end{pmatrix} \quad (15.10b)$$

and

$$\mathbf{U} = (\mathbf{e}_1, \mathbf{e}_2, \dots, \mathbf{e}_n) \quad (15.10c)$$

Geometrically, \mathbf{U} describes a rigid rotation of the original coordinate system while \mathbf{A} is the amount by which unit lengths in the original coordinate system are scaled in the transformed system. Using Equation 15.10a, it is easy to show that

$$\mathbf{A}^{-1} = \mathbf{U}\mathbf{A}^{-1}\mathbf{U}^T \quad (15.11a)$$

$$\mathbf{A}^{1/2} = \mathbf{U}\mathbf{A}^{1/2}\mathbf{U}^T \quad (15.11b)$$

where the **square root matrix** $\mathbf{A}^{1/2}$ (which is also symmetric) satisfies

$$\mathbf{A}^{1/2} \mathbf{A}^{1/2} = \mathbf{A}$$

Since \mathbf{A} is diagonal, the i th diagonal elements of \mathbf{A}^{-1} and $\mathbf{A}^{1/2}$ are λ_i^{-1} and $\lambda_i^{1/2}$ respectively, implying that if λ_i is an eigenvalue of \mathbf{A} , then λ_i^{-1} and $\sqrt{\lambda_i}$ are eigenvalues of \mathbf{A}^{-1} and $\mathbf{A}^{1/2}$. Note that Equations 15.11a/b imply that \mathbf{A} , \mathbf{A}^{-1} and $\mathbf{A}^{1/2}$ all have the same eigenvectors. Finally, using Equation 15.10a we see that premultiplying \mathbf{A} by \mathbf{U}^T and then postmultiplying by \mathbf{U} gives a diagonal matrix whose elements are the eigenvalues of \mathbf{A} ,

$$\begin{aligned} \mathbf{U}^T \mathbf{A} \mathbf{U} &= \mathbf{U}^T (\mathbf{U} \mathbf{A} \mathbf{U}^T) \mathbf{U} = (\mathbf{U}^T \mathbf{U}) \mathbf{A} (\mathbf{U}^T \mathbf{U}) \\ &= \mathbf{A} \end{aligned} \tag{15.12}$$

5. The **Rayleigh-Ritz** theorem gives useful bounds on quadratic products associated with the symmetric matrix \mathbf{A} : if the eigenvalues of \mathbf{A} are ordered as $\lambda_{max} = \lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_n = \lambda_{min}$, then for any vector of constants \mathbf{c} ,

$$\lambda_1 \|\mathbf{c}\| \geq \mathbf{c}^T \mathbf{A} \mathbf{c} \geq \lambda_n \|\mathbf{c}\| \tag{15.13a}$$

Alternatively, if \mathbf{c} is of unit length

$$\max_{\|\mathbf{c}\|=1} \mathbf{c}^T \mathbf{A} \mathbf{c} = \lambda_1 \tag{15.13b}$$

$$\min_{\|\mathbf{c}\|=1} \mathbf{c}^T \mathbf{A} \mathbf{c} = \lambda_n \tag{15.13c}$$

Where the maximum and minimum occur when $\mathbf{c} = \mathbf{e}_1$ and $\mathbf{c} = \mathbf{e}_n$, the eigenvectors associated with λ_1 and λ_n . This is an especially useful result for bounding variances. Consider a univariate random variable $y = \mathbf{c}^T \mathbf{x}$ formed by a linear combination of the elements of a random vector \mathbf{x} . From Equation 7.20, $\sigma^2(y) = \mathbf{c}^T \boldsymbol{\Sigma}_{\mathbf{x}} \mathbf{c}$ and applying Equation 15.13a,

$$\lambda_1 \|\mathbf{c}\|^2 \geq \sigma^2(y) \geq \lambda_n \|\mathbf{c}\|^2 \tag{15.14}$$

where λ_1 is the largest (**leading** or **dominant**) and λ_n the smallest eigenvalue of $\boldsymbol{\Sigma}_{\mathbf{x}}$.

Correlations can be Removed by a Matrix Transformation

A particularly powerful use of diagonalization is that it allows us to extract a set of n uncorrelated variables when the variance-covariance matrix $\boldsymbol{\Sigma}_{\mathbf{x}}$ is nonsingular and of dimension n . Consider the transformation

$$\mathbf{y} = \mathbf{U}^T \mathbf{x} \tag{15.15a}$$

where $\mathbf{U} = (\mathbf{e}_1, \mathbf{e}_2, \dots, \mathbf{e}_n)$ contains the normalized eigenvectors of $\Sigma_{\mathbf{x}}$. Since \mathbf{U} is an orthonormal matrix, this transformation is a rigid rotation of the axes of the original (x_1, \dots, x_n) coordinate system to a new system given by (y_1, \dots, y_n) . Applying Equations 7.22b and 15.12, the variance-covariance matrix for \mathbf{y} is

$$\Sigma_{\mathbf{y}} = \mathbf{U}^T \Sigma_{\mathbf{x}} \mathbf{U} = \Lambda \quad (15.15b)$$

where Λ is a diagonal matrix whose elements are the eigenvalues of $\Sigma_{\mathbf{x}}$, so that

$$\sigma(y_i, y_j) = \begin{cases} \lambda_i & \text{if } i = j \\ 0 & \text{if } i \neq j \end{cases}$$

The rigid rotation introduced by \mathbf{U} thus creates a set of n uncorrelated variables, the i th of which is

$$y_i = \mathbf{e}_i^T \mathbf{x} \quad (15.15c)$$

Since \mathbf{e}_i are defined to be of unit length, from Equation 15.3b we have $y_i = \mathbf{e}_i^T \mathbf{x} = \text{Proj}(\mathbf{x} \text{ on } \mathbf{e}_i)$, so that this new variable is the projection of \mathbf{x} onto the i th eigenvector of $\Sigma_{\mathbf{x}}$, implying that the axes of new coordinate system are given by the orthogonal set of eigenvectors of $\Sigma_{\mathbf{x}}$.

Canonical Axes of Quadratic Forms

The transformation $\mathbf{y} = \mathbf{U}^T \mathbf{x}$ given by Equation 15.15a applies to any symmetric matrix, and is referred to as the **canonical transformation** associated with that matrix. The canonical transformation simplifies the interpretation of the quadratic form $\mathbf{x}^T \mathbf{A} \mathbf{x}$ as rotation of the original axes to align them with the eigenvalues of \mathbf{A} removes all cross-product terms on this new coordinate system. Applying Equations 15.15a and 15.12 transforms the quadratic form to one in which the square matrix is diagonal,

$$\begin{aligned} \mathbf{x}^T \mathbf{A} \mathbf{x} &= (\mathbf{U} \mathbf{y})^T \mathbf{A} \mathbf{U} \mathbf{y} = \mathbf{y}^T (\mathbf{U}^T \mathbf{A} \mathbf{U}) \mathbf{y} \\ &= \mathbf{y}^T \Lambda \mathbf{y} \\ &= \sum_{i=1}^n \lambda_i y_i^2 \end{aligned} \quad (15.16)$$

where λ_i and \mathbf{e}_i are the eigenvalues and associated (normalized) eigenvectors of \mathbf{A} and $y_i = \mathbf{e}_i^T \mathbf{x}$. The new axes defined by \mathbf{e}_i are the **canonical** (or **principal**) axes. Since the $y_i^2 \geq 0$, Equation 15.16 immediately shows the connection between the signs of the eigenvalues of a matrix and whether that matrix is positive definite, negative definite, or indefinite. **If all eigenvalues are positive (all $\lambda_i > 0$), then the quadratic form is always positive (unless all the y_i are zero) and hence \mathbf{A} is positive definite.** If all eigenvalues are negative (all $\lambda_i < 0$), then \mathbf{A} is negative definite as the quadratic form is always negative. If at least one eigenvalue is zero, then \mathbf{A} is at most semidefinite, while if \mathbf{A} has both positive and negative eigenvalues it is **indefinite.**

Equations of the form

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \sum_{i=1}^n \sum_{j=1}^n a_{ij} x_i x_j = c^2 \tag{15.17a}$$

arise fairly frequently. For example, they describe surfaces of constant variance (tracing out the surface created by vectors \mathbf{b} such that $\mathbf{b}^T \mathbf{x}$ has constant variance c^2 , see Figure 15.5) or surfaces of constant fitness in quadratic fitness regressions (those vectors of phenotypic values \mathbf{z} such that $w(\mathbf{z}) = a + (\mathbf{z} - \boldsymbol{\mu})^T \boldsymbol{\gamma} (\mathbf{z} - \boldsymbol{\mu})$ is constant). Solutions to Equation 15.17a describe **quadratic surfaces** — for two dimensions these are the familiar conic sections (ellipses, parabolas, or hyperbolas). Equation 15.16 greatly simplifies the interpretation of these surfaces by removing all cross product terms, giving

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \sum_{i=1}^n \lambda_i y_i^2 = c^2 \tag{15.17b}$$

Since $(y_i)^2$ and $(-y_i)^2$ have the same value, the canonical axes of \mathbf{A} are also the axes of symmetry for the quadratic surface generated by quadratic forms involving \mathbf{A} . **When all the eigenvalues of \mathbf{A} are positive** (as occurs with non-singular variance-covariance and other positive definite matrices), **Equation 15.17b describes an ellipsoid whose axes of symmetry are given by the eigenvectors of \mathbf{A}** . The distance from the origin to the surface along the axis given by \mathbf{e}_i is $\lambda_i y_i^2 = c^2$ or $y_i = c \lambda_i^{-1/2}$, as can be seen by setting all the y_k equal to zero except for y_i , giving $\mathbf{x}^T \mathbf{A} \mathbf{x} = \lambda_i y_i^2 = c^2$. Figure 15.5 shows an example of a two-dimensional constant-variance surface: if we plot the entire set of vectors \mathbf{b} such that the variable $y = \mathbf{b}^T \mathbf{x}$ has variance $c^2 = \mathbf{b}^T \boldsymbol{\Sigma}_X \mathbf{b}$, the tips of these vectors sweep out the ellipse.

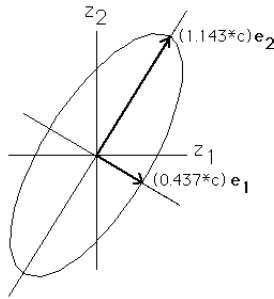


Figure 15.5. The general shape of surfaces of constant variance for the additive genetic variance-covariance matrix \mathbf{G} given in Example 1. Defining a new composite character $y = az_1 + bz_2$, the rotated ellipse represents the set of (a, b)

values that give y the same additive genetic variance c^2 . The major axis of the ellipse is along \mathbf{e}_2 , the eigenvector associated with the smallest eigenvalue of \mathbf{G} , where $\lambda_2 \simeq 0.765$, giving $\lambda_2^{-1/2} \simeq 1.143$. The minor axis of the ellipse is along \mathbf{e}_1 , the eigenvector associated with the largest eigenvalue of \mathbf{G} , where $\lambda_1 \simeq 5.241$, giving $\lambda_1^{-1/2} \simeq 0.437$.

Principal Components of the Variance-Covariance Matrix

We are very interested in how the variance of a random vector can be decomposed into independent components. For example, even though we may be measuring n variables, only one or two of these may account for the majority of the variation. If this is the case we may wish to exclude those variables contributing very little variation from further analysis. More generally, if the random variables are correlated, then certain **linear combinations** of the elements of \mathbf{x} may account for most of the variance. The procedure of **principal component analysis** extracts these combinations by decomposing the variance of \mathbf{x} into a series of orthogonal vectors, the first of which explains the most variation possible for any single vector, the second the next possible amount, and so on until the entire variance of \mathbf{x} has been accounted for.

Consider Figure 15.5. Since the set of points comprising the ellipse represents those linear combinations of the random variables of \mathbf{z} giving **equal** variance, we see that the closer a point on this curve is to the origin, the more variance there is in that direction. The points closest to the origin are those that lie along the axis defined by \mathbf{e}_1 , while those furthest away lie along the axis defined by \mathbf{e}_2 . Here \mathbf{e}_1 and \mathbf{e}_2 are the principal components of \mathbf{G} , with the first principal component accounting for most of the variation of \mathbf{G} . In particular, the ratio of additive variances for the characters $y_1 = \mathbf{e}_1^T \mathbf{z}$ and $y_2 = \mathbf{e}_2^T \mathbf{z}$ is $\sigma^2(y_1)/\sigma^2(y_2) = \sigma^2(\mathbf{e}_1^T \mathbf{z})/\sigma^2(\mathbf{e}_2^T \mathbf{z}) = \mathbf{e}_1^T \mathbf{G} \mathbf{e}_1 / \mathbf{e}_2^T \mathbf{G} \mathbf{e}_2 = \lambda_1/\lambda_2 \simeq 5.241/0.765 \simeq 6.85$, so that a character in the direction of \mathbf{e}_1 has almost seven times as much additive variance as a character lying in the direction of \mathbf{e}_2 .

In general, suppose we have an n -dimensional variance-covariance matrix $\Sigma_{\mathbf{x}}$. Ordering the eigenvalues of $\Sigma_{\mathbf{x}}$ as $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_n$ then from Equation 15.13b the maximum variance for any linear combination $y = \mathbf{c}_1^T \mathbf{x}$ (subject to the constraint that $\|\mathbf{c}_1\| = 1$) is

$$\max \sigma^2(y) = \max_{\|\mathbf{c}_1\|=1} \sigma^2(\mathbf{c}_1^T \mathbf{x}) = \mathbf{c}_1^T \Sigma_{\mathbf{x}} \mathbf{c}_1 = \lambda_1$$

which occurs when $\mathbf{c}_1 = \mathbf{e}_1$. This vector is the **first principal component** (often abbreviated as PC1). Excluding PC1, consider how the remaining variance can be explained. The vector \mathbf{c}_2 orthogonal to PC1 (e.g., $\mathbf{c}_2^T \mathbf{c}_1 = 0$) that maximizes the remaining variance, e.g., maximizes $\sigma^2(\mathbf{c}_2^T \mathbf{x})$, can be shown to be \mathbf{e}_2 and that the amount of the remaining variation it explains is λ_2 (e.g., Morrison 1976, Johnson

and Wichern 1988). Proceeding in this fashion, we see that the i th PC is given by \mathbf{e}_i and that the amount of variation it accounts for is

$$\lambda_i / \sum_{k=1}^n \lambda_k = \frac{\lambda_i}{\text{tr}(\boldsymbol{\Sigma}_{\mathbf{X}})} \tag{15.18}$$

Example 5. Again consider the additive genetic variance-covariance matrix \mathbf{G} as given in Example 1. Since $\lambda_1 \simeq 5.241$, $\lambda_2 \simeq 0.765$ and $\text{tr}(\mathbf{G}) = 4 + 2 = 6$, the first PC explains $5.241/6 \simeq 0.8735$ or 87 percent of the variance in \mathbf{G} . Note, however, that although the first PC accounts for the majority of variation, the amount of variation explained by PC1 for any *particular* variable $y = \mathbf{b}^T \mathbf{x}$ depends on the projection of \mathbf{b} onto PC1. For example, if $\mathbf{b} = \mathbf{e}_2$, then the projection of \mathbf{b} onto PC1 has length zero and hence PC1 accounts for no variation of y .

Example 6. Jolicoeur and Mosimann (1960) measured three carapace characters in 24 males of the painted turtle *Chrysemys picta marginata*. Letting z_1 be carapace length, z_2 maximum carapace width, and z_3 carapace height, the resulting sample variance-covariance matrix ($\mathbf{S}_{\mathbf{Z}}$, the sample estimate of $\boldsymbol{\Sigma}_{\mathbf{Z}}$) for these data is

$$\mathbf{S}_{\mathbf{Z}} = \begin{pmatrix} 13.77 & 79.15 & 37.13. \\ 79.15 & 50.04 & 21.65 \\ 37.13. & 21.65 & 13.26 \end{pmatrix}$$

Hence, $\text{tr}(\mathbf{S}_{\mathbf{Z}}) = 13.77 + 50.04 + 13.26 = 200.07$. The eigenvalues for this matrix are found to be

$$\lambda_1 = 195.281, \quad \lambda_2 = 3.687, \quad \lambda_3 = 1.103$$

and the associated normalized eigenvectors are

$$\mathbf{e}_1 = \begin{pmatrix} 0.840 \\ 0.492 \\ 0.229 \end{pmatrix}, \quad \mathbf{e}_2 = \begin{pmatrix} 0.488 \\ -0.870 \\ 0.079 \end{pmatrix}, \quad \mathbf{e}_3 = \begin{pmatrix} 0.213. \\ 0.043 \\ -0.971 \end{pmatrix}$$

PC1 accounts for 97.6% of the variation (as $195.281/200.07 = 0.976$), while PC2 and PC3 account for 1.84% and 0.55%, respectively. Jolicoeur and Mosimann interpret PC1 as measuring overall size as the new variable is

$$y_1 = \mathbf{e}_1^T \mathbf{z} = 0.840z_1 + 0.492z_2 + 0.229z_3$$

which corresponds to a simultaneous change in all three variables, as is expected to occur as individuals change their overall size. Likewise PC2 and PC3 are

$$y_2 = \mathbf{e}_2^T \mathbf{z} = 0.488z_1 - 0.870z_2 + 0.079z_3$$

$$y_3 = \mathbf{e}_3^T \mathbf{z} = 0.213z_1 + 0.043z_2 - 0.971z_3$$

which Jolicoeur and Mosimann interpret as measures of shape. Since the coefficient on z_3 is small relative to the others in PC2, they interpret PC2 as measuring the tradeoff between length (z_1) and width (z_2). After removing the variation in size, 1.84% of the remaining variation can be accounted for by differences in the shape measured by length versus width. Likewise, since the coefficient in z_2 is very small in PC3, it measures shape differences due to length (z_1) versus height (z_3).

This example points out some of the advantages, and possible pitfalls, of using principal components analysis to reduce the data. Essentially all (over 97 percent) of the variance in the three measured characters is accounted for by variation in overall size, with the remaining variation accounted for by differences in shape. While the temptation is strong to simply consider overall size and ignore all shape information, it might be the case that selection is largely ignoring variation in size and instead is focusing on (size-independent) shape differences. In this case, an analysis ignoring shape (such as would occur if only the new character generated by PC1 is considered) would be very misleading. A further complication with principal component analysis is that it can often be very difficult to give biological interpretations to the new characters resulting from the rotation of the coordinate system. This example serves as a brief introduction to the important field of **morphometrics**, which is concerned with how to quantify and compare the size and shape of organisms. The reader is referred to Pimentel (1979), Reyment et al. (1984), and especially Bookstein et al. (1985), Rohlf and Bookstein (1990), and Reyment (1991) for detailed treatments.

THE MULTIVARIATE NORMAL DISTRIBUTION

Recall the density of the multivariate normal distribution,

$$\phi(\mathbf{x}) = (2\pi)^{-n/2} |\boldsymbol{\Sigma}_{\mathbf{x}}|^{-1/2} \exp \left[-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}_{\mathbf{x}}^{-1} (\mathbf{x} - \boldsymbol{\mu}) \right] \quad (15.19a)$$

Thus surfaces of equal probability for MVN distributed vectors satisfy

$$(\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}_{\mathbf{x}}^{-1} (\mathbf{x} - \boldsymbol{\mu}) = c^2 \quad (15.19b)$$

From the discussion following Equation 15.17b, these surfaces are n -dimensional ellipsoids centered at $\boldsymbol{\mu}$ whose axes of symmetry are given by the principal components (the eigenvectors) of $\boldsymbol{\Sigma}_{\mathbf{x}}$. The length of the ellipsoid along the i th axis is $c\sqrt{\lambda_i}$ where λ_i is the eigenvalue associated with the eigenvector \mathbf{e}_i (Figure 15.6).

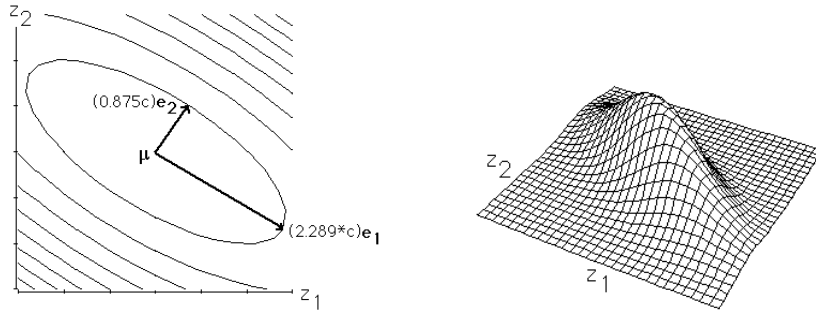


Figure 15.6. Left: Surfaces of equal probability assuming that the additive genetic values associated with the characters z_1 and z_2 in Example 1 are $\text{MVN}(\boldsymbol{\mu}, \mathbf{G})$. These surfaces are ellipses centered at $\boldsymbol{\mu}$, with the major axis of the ellipse along \mathbf{e}_1 and the minor axis along \mathbf{e}_2 . Right: A plot of the associated probability density. Slicing along either the major or minor axis gives a normal curve. Since the variance in the major axis is greater, the curve is much broader along this axis. The covariance between the breeding values of z_1 and z_2 rotates the distribution so that the principal axes do not coincide with the (z_1, z_2) axes.

Applying the canonical transformation (Equation 15.15a), we can change coordinate systems by a rigid rotation to remove any correlations between the variables in \mathbf{x} . Taking $\mathbf{y} = \mathbf{U}^T(\mathbf{x} - \boldsymbol{\mu})$,

$$\mathbf{y} \sim \text{MVN}(\mathbf{0}, \boldsymbol{\Lambda}) \tag{15.20a}$$

where $\boldsymbol{\Lambda}$ and \mathbf{U} are the matrices defined by Equations 15.10b/c for the diagonalization of $\boldsymbol{\Sigma}_{\mathbf{x}}$. In particular,

$$y_i = \mathbf{e}_i^T(\mathbf{x} - \boldsymbol{\mu}) \quad \text{where} \quad y_i \sim \text{N}(0, \lambda_i) \tag{15.20b}$$

Note from Equation 15.19 that since the y_i are uncorrelated, they are independent as the joint probability density is the product of n individual univariate normal densities. We can further transform the original vector by taking

$$y_i = \frac{\mathbf{e}_i^T(\mathbf{x} - \boldsymbol{\mu})}{\sqrt{\lambda_i}} \quad \text{giving} \quad y_i \sim \text{N}(0, 1) \tag{15.20c}$$

Thus, the transformation

$$\mathbf{y} = \boldsymbol{\Lambda}^{-1/2}\mathbf{U}^T(\mathbf{x} - \boldsymbol{\mu}) \tag{15.20d}$$

implies that $\mathbf{y} \sim \text{MVN}(\mathbf{0}, \mathbf{I})$, the elements of \mathbf{y} being n independent unit normal random variables.

TESTING FOR MULTIVARIATE NORMALITY

While multivariate normality is often assumed, it is rarely tested. In Chapter 10 we briefly discussed two approaches for testing univariate normality, one graphical and the other based on if the observed skewness and/or kurtosis exceeds that expected for a Gaussian. Both of these can be extended to testing for multivariate normality. Additional methods are reviewed by Gnanadesikan (1977) and Seber (1984).

Graphical Tests: Chi-square Plots

A fairly simple graphical test can be developed by extending the notion of the normal probability plot used to check univariate normality (Chapter 10). Recall that in this case the observations are ranked and then plotted against their expected values under normality. Departures from linearity signify departures from normality.

We can apply this same approach to check for multivariate normality. From Equation 15.20d, if $\mathbf{z} \sim \text{MVN}(\boldsymbol{\mu}, \boldsymbol{\Sigma}_{\mathbf{z}})$, then each element of the vector

$$\mathbf{y} = \boldsymbol{\Lambda}^{-1/2} \mathbf{U}^T (\mathbf{z} - \boldsymbol{\mu})$$

is an independent unit normal ($\mathbf{y} \sim \text{MVN}(\mathbf{0}, \mathbf{I})$). Solving for \mathbf{z} gives

$$(\mathbf{z} - \boldsymbol{\mu}) = \mathbf{U} \boldsymbol{\Lambda}^{1/2} \mathbf{y}$$

Using this and recalling Equation 15.11a,

$$\begin{aligned} (\mathbf{z} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}_{\mathbf{z}}^{-1} (\mathbf{z} - \boldsymbol{\mu}) &= (\mathbf{U} \boldsymbol{\Lambda}^{1/2} \mathbf{y})^T (\mathbf{U} \boldsymbol{\Lambda}^{-1} \mathbf{U}^T) (\mathbf{U} \boldsymbol{\Lambda}^{1/2} \mathbf{y}) \\ &= \mathbf{y}^T \boldsymbol{\Lambda}^{1/2} (\mathbf{U}^T \mathbf{U}) \boldsymbol{\Lambda}^{-1} (\mathbf{U}^T \mathbf{U}) \boldsymbol{\Lambda}^{1/2} \mathbf{y} \\ &= \mathbf{y}^T \mathbf{y} = \sum_{i=1}^n y_i^2 \end{aligned} \quad (15.21)$$

Thus if $\mathbf{z} \sim \text{MVN}$, the quadratic form given by Equation 15.21 is the sum of n independent squared unit normal random variables. By definition, this sum is a χ^2 random variable with n degrees of freedom (e.g., Morrison 1976), suggesting that one test for multivariate normality is to compare the goodness of fit of the scaled distances

$$d_i^2 = (\mathbf{z}_i - \bar{\mathbf{z}})^T \mathbf{S}_{\mathbf{z}}^{-1} (\mathbf{z}_i - \bar{\mathbf{z}}) \quad (15.22)$$

to a χ_n^2 . Here \mathbf{z}_i is the vector of observations from the i th individual, $\bar{\mathbf{z}}$ the vector of sample means, and $\mathbf{S}_{\mathbf{z}}^{-1}$ the inverse of the sample variance-covariance matrix. (We use the term distance because $\Sigma_{\mathbf{y}} = \mathbf{I}$, giving the variance of any linear combination $\mathbf{c}^T \mathbf{y}$ as $\mathbf{c}^T \Sigma_{\mathbf{y}} \mathbf{c} = \mathbf{c}^T \mathbf{I} \mathbf{c} = \|\mathbf{c}\|^2$. Thus, regardless of orientation, any two \mathbf{y} vectors having the same length also have the same variance, which equals their squared Euclidean distance.) We can order these distances as

$$d_{(1)}^2 \leq d_{(2)}^2 \leq \dots \leq d_{(m)}^2$$

where m is the number of individuals sampled. Note that $d_{(i)}^2$ is the i th smallest distance, whereas d_i^2 is the distance associated with the vector of observations for the i th individual. Let $\chi_n^2(\alpha)$ correspond to the value of a chi-square random variable x with n degrees of freedom that satisfies $\text{Prob}(x \leq \chi_n^2(\alpha)) = \alpha$. Under multivariate normality, we expect the points

$$\left(d_{(i)}^2, \chi_n^2 \left(\frac{i - 1/2}{m} \right) \right)$$

to fall along a line, as the i th ordered distance has i/m observations less than or equal to it (the factor of $1/2$ is added as a correction for continuity). As with normal probability plots, departures from multivariate normality are indicated by departures from linearity. Complete tables of the χ^2 may be difficult to locate, in which case the appropriate $\chi_n^2(\alpha)$ values can be numerically obtained using the incomplete gamma function (see Press et al. 1986 for details).

Example 7. Consider again the data of Jolicoeur and Mosimann (1960) on carapace characters in 24 male turtles. Are the characters z_1 (carapace length) and z_2 (maximum carapace width) jointly bivariate normally distributed? Here $n = 2$ and $m = 24$ and

$$\bar{\mathbf{z}} = \begin{pmatrix} 113.13 \\ 88.29 \end{pmatrix}, \quad \mathbf{S}_{\mathbf{z}} = \begin{pmatrix} 13.77 & 79.15 \\ 79.15 & 50.04 \end{pmatrix}, \quad \mathbf{S}_{\mathbf{z}}^{-1} = \begin{pmatrix} 0.0737 & -0.1165 \\ -0.1165 & 0.2043 \end{pmatrix}$$

where $\mathbf{S}_{\mathbf{z}}$ is the sample covariance matrix. A partial list of the 24 vectors of observations are

$$\mathbf{z}_1 = \begin{pmatrix} 93 \\ 74 \end{pmatrix}, \quad \dots, \quad \mathbf{z}_{11} = \begin{pmatrix} 113 \\ 88 \end{pmatrix}, \quad \dots, \quad \mathbf{z}_{24} = \begin{pmatrix} 135 \\ 106 \end{pmatrix}$$

Applying Equation 15.22, these observations translate into the distances

$$d_1^2 = 4.45, \quad \dots, \quad d_{11}^2 = 0.002, \quad \dots, \quad d_{24}^2 = 9.277$$

After rank ordering, these correspond to $d_{(23)}^2$, $d_{(1)}^2$, and $d_{(24)}^2$, respectively. For $d_{(23)}^2$, the matching value when distances are chi-squared distributed is

$$\chi_2^2 \left(\frac{23 - 1/2}{24} \right) = \chi_2^2(0.913.)$$

From chi-square tables, we find $\text{Prob}(\chi_2^2 \leq 5.561) = 0.913$. so that the data point generated from \mathbf{z}_1 is (4.45, 5.561). Likewise, the chi-square values for $d_{(1)}^2$ and $d_{(24)}^2$ are 0.043 and 7.727, respectively. Proceeding similarly for the other values, we obtain the curve plotted in Figure 15.7. This curve departs somewhat from linearity. Further, under the assumption of multivariate normality, the line is expected to pass through the origin, while the best linear fit of these data departs from the origin. Transforming the data by taking logs gives a slightly better fit to a MVN (Figure 15.7).

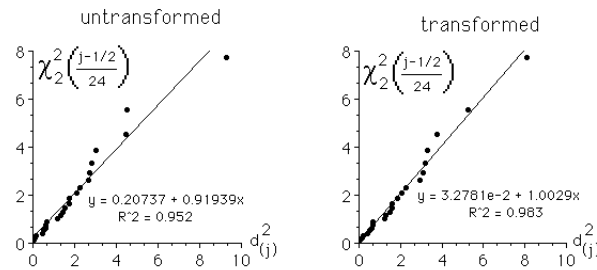


Figure 15.7. Plots of ranked distance data ($d_{(j)}^2$ being the j th smallest distance) versus the expected corresponding χ^2 value for the data in Example 7. **Left:** The untransformed data does not appear to depart significantly from linearity. **Right:** Log-transforming the data gives a slightly better linear fit ($r^2 = 0.983$ versus $r^2 = 0.952$ for the untransformed data), with the best-fitting line passing through the origin as expected if the distance data follow a χ^2 distribution.

Mardina's Test: Multivariate Skewness and Kurtosis

As was the case for univariate normality, we can test for multivariate normality by examining the skewness and kurtosis of the sample. Mardina (1970) proposed multivariate extensions of skewness and kurtosis and suggested a large sample test based on the asymptotic distribution of these statistics. Let \mathbf{z}_i be the i -th vector of observations, $\bar{\mathbf{z}}$ the vector of sample means, and $\mathbf{S}_{\mathbf{z}}$ sample covariance matrix.

If there are m vectors of observations (with each vector measuring n characters), then the multivariate skewness is estimated by

$$b_{1,n} = \frac{1}{m^2} \sum_{i=1}^m \sum_{j=1}^m ((\mathbf{z}_i - \bar{\mathbf{z}})^T \mathbf{S}_{\mathbf{z}}^{-1} (\mathbf{z}_j - \bar{\mathbf{z}}))^3 \quad (15.23a)$$

while the multivariate kurtosis is estimated by

$$b_{2,n} = \frac{1}{m} \sum_{i=1}^m ((\mathbf{z}_i - \bar{\mathbf{z}})^T \mathbf{S}_{\mathbf{z}}^{-1} (\mathbf{z}_i - \bar{\mathbf{z}}))^2 \quad (15.23b)$$

If $\mathbf{z} \sim \text{MVN}$, then $b_{1,n}$ and $b_{2,n}$ have expected values 0 and $n(n+2)$. For large m , Mardina (1970) showed that the (scaled) multivariate skewness is asymptotically distributed as a chi-square random variable with f degrees of freedom, viz.,

$$\frac{m}{6} b_{1,n} \sim \chi_f^2, \quad \text{where } f = \frac{n(n+1)(n+2)}{6} \quad (15.24a)$$

Likewise for large m , the multivariate kurtosis (following appropriate scaling) is distributed as a unit-normal, viz.,

$$\frac{b_{2,n} - n(n+2)}{\sqrt{8n(n+2)/m}} \sim N(0, 1) \quad (15.24b)$$

If either Equation 15.24a or 15.24b is significant, then multivariate normality is rejected.

Example 8. Again, let us examine the data of Jolicoeur and Mosimann (1960). Does the data considered in Example 7 display significant skewness or kurtosis? Here $n = 2$ and $m = 24$. Applying Equations 15.24a/b gives $b_{1,2} = 0.6792$, $b_{2,2} = 7.6043$. Considering skewness first,

$$\frac{m}{6} b_{1,2} = \frac{24}{6} 0.6792 = 2.717$$

is approximately chi-square distributed with $f = 2(2+1)(2+2)/6 = 4$ degrees of freedom. Since $\text{Prob}(\chi_4^2 \geq 2.717) \simeq 0.606$, this is not significant. Turning to kurtosis, Equation 15.24b gives

$$\frac{b_{2,n} - n(n+2)}{\sqrt{8n(n+2)/m}} = \frac{7.6043 - 8}{1.633} \simeq -0.2423$$

which is also not significant as $\text{Prob}(|N(0, 1)| \geq 0.2423) \simeq 0.81$. Transforming the data by taking logs gives $b_{1,2} = 0.2767$ and $b_{2,2} = 7.1501$, improving the departure from skewness but increasing the departure from kurtosis. Applying Equations 15.24a/b gives 1.068 and -0.5204 , again these are not significant. Reyment (1971) gives a number of other biological examples using Mardina's test.

Derivatives of Vectors and Vector-valued Functions

The **gradient** (or **gradient vector**) of a scalar function of a vector variable is obtained by taking partial derivatives of the function with respect to each variable. In matrix notation, the gradient operator is

$$\nabla_{\mathbf{x}}[f] = \frac{\partial f}{\partial \mathbf{x}} = \begin{pmatrix} \frac{\partial f}{\partial x_1} \\ \frac{\partial f}{\partial x_2} \\ \vdots \\ \frac{\partial f}{\partial x_n} \end{pmatrix}$$

The gradient at point \mathbf{x}_o corresponds to a vector indicating the direction of steepest ascent of the function at that point (the multivariate slope of f at the point \mathbf{x}_o). For example $f(\mathbf{x}) = \mathbf{x}^T \mathbf{x}$ has gradient vector $2\mathbf{x}$. At the point \mathbf{x}_o , $\mathbf{x}^T \mathbf{x}$ locally increases most rapidly if we change \mathbf{x} in the same the direction as the vector going from point \mathbf{x}_o to point $\mathbf{x}_o + 2\delta \mathbf{x}_o$, where δ is a small positive value.

For a vector \mathbf{a} and matrix \mathbf{A} of constants, it can easily be shown (e.g., Morrison 1976, Graham 1981, Searle 1982) that

$$\nabla_{\mathbf{x}} [\mathbf{a}^T \mathbf{x}] = \nabla_{\mathbf{x}} [\mathbf{x}^T \mathbf{a}] = \mathbf{A} \quad (15.39a)$$

$$\nabla_{\mathbf{x}} [\mathbf{A} \mathbf{x}] = \mathbf{A}^T \quad (15.39b)$$

Turning to quadratic forms, if \mathbf{A} is symmetric, then

$$\nabla_{\mathbf{x}} [\mathbf{x}^T \mathbf{A} \mathbf{x}] = 2 \cdot \mathbf{A} \mathbf{x} \quad (15.39c)$$

$$\nabla_{\mathbf{x}} [(\mathbf{x} - \mathbf{A})^T \mathbf{A} (\mathbf{x} - \mathbf{A})] = 2 \cdot \mathbf{A} (\mathbf{x} - \mathbf{A}) \quad (15.39d)$$

$$\nabla_{\mathbf{x}} [(\mathbf{A} - \mathbf{x})^T \mathbf{A} (\mathbf{A} - \mathbf{x})] = -2 \cdot \mathbf{A} (\mathbf{A} - \mathbf{x}) \quad (15.39e)$$

Taking $\mathbf{A} = \mathbf{I}$, Equation 15.39c implies

$$\nabla_{\mathbf{x}} [\mathbf{x}^T \mathbf{x}] = \nabla_{\mathbf{x}} [\mathbf{x}^T \mathbf{I} \mathbf{x}] = 2 \cdot \mathbf{I} \mathbf{x} = 2 \cdot \mathbf{x} \quad (15.39f)$$

Two final useful identities follow from the chain rule of differentiation,

$$\nabla_{\mathbf{x}} [\exp[f(\mathbf{x})]] = \exp[f(\mathbf{x})] \cdot \nabla_{\mathbf{x}} [f(\mathbf{x})] \quad (15.39g)$$

$$\nabla_{\mathbf{x}} [\ln[f(\mathbf{x})]] = \frac{1}{f(\mathbf{x})} \cdot \nabla_{\mathbf{x}} [f(\mathbf{x})] \quad (15.39h)$$

Example 10. Writing the MVN distribution as

$$\varphi(\mathbf{x}) = a \exp\left(-\frac{1}{2} \cdot (\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} (\mathbf{x} - \boldsymbol{\mu})\right)$$

where $a = \pi^{-n/2} |\mathbf{V}_{\mathbf{x}}|^{-1/2}$, then from Equation 15.39g,

$$\nabla_{\mathbf{x}} [\varphi(\mathbf{x})] = \varphi(\mathbf{x}) \cdot \nabla_{\mathbf{x}} \left[\left(-\frac{1}{2}\right) \cdot (\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} (\mathbf{x} - \boldsymbol{\mu}) \right]$$

Applying Equation 15.39d gives

$$\nabla_{\mathbf{x}} [\varphi(\mathbf{x})] = -\varphi(\mathbf{x}) \cdot \mathbf{V}_{\mathbf{x}}^{-1} (\mathbf{x} - \boldsymbol{\mu}) \quad (15.40a)$$

Note here that $\varphi(\mathbf{x})$ is a scalar and hence its order of multiplication does not matter, while the order of the other variables (being matrices) is critical. Similarly, we can consider the MVN as a function of the mean vector $\boldsymbol{\mu}$, in which case Equation 15.39e implies

$$\nabla_{\boldsymbol{\mu}} [\varphi(\mathbf{x}, \boldsymbol{\mu})] = \varphi(\mathbf{x}, \boldsymbol{\mu}) \cdot \mathbf{V}_{\mathbf{x}}^{-1} (\mathbf{x} - \boldsymbol{\mu}) \quad (15.40b)$$

Example 13. Consider obtaining the least-squares solution for the general linear model, $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mathbf{e}$, where we wish to find the value of $\boldsymbol{\beta}$ that minimizes the residual error given \mathbf{y} and \mathbf{X} . In matrix form,

$$\begin{aligned} \sum_{i=1}^n e_i^2 &= \mathbf{e}^T \mathbf{e} = (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^T (\mathbf{y} - \mathbf{X}\boldsymbol{\beta}) \\ &= \mathbf{y}^T \mathbf{y} - \boldsymbol{\beta}^T \mathbf{X}^T \mathbf{y} - \mathbf{y}^T \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\beta}^T \mathbf{X}^T \mathbf{X}\boldsymbol{\beta} \\ &= \mathbf{y}^T \mathbf{y} - 2\boldsymbol{\beta}^T \mathbf{X}^T \mathbf{y} + \boldsymbol{\beta}^T \mathbf{X}^T \mathbf{X}\boldsymbol{\beta} \end{aligned}$$

where the last step follows from Equation 7.18. To find the vector $\boldsymbol{\beta}$ that minimizes $\mathbf{e}^T \mathbf{e}$, taking the derivative with respect to $\boldsymbol{\beta}$ and using Equations 15.39a/c gives

$$\frac{\partial \mathbf{e}^T \mathbf{e}}{\partial \boldsymbol{\beta}} = -2\mathbf{X}^T \mathbf{y} + 2\mathbf{X}^T \mathbf{X}\boldsymbol{\beta}$$

Setting this equal to zero gives $\mathbf{X}^T \mathbf{X} \boldsymbol{\beta} = \mathbf{X}^T \mathbf{y}$ giving

$$\boldsymbol{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

The hessian matrix, local maxima/minima, and multidimensional Taylor series.

In univariate calculus, local extrema of a function occur when the slope (first derivative) is zero. The multivariate extension of this is that the gradient vector is zero, so that the slope of the function with respect to all variables is zero. A point \mathbf{x}_e where this occurs is called a **stationary** or **equilibrium** point, and corresponds to either a local maximum, minimum, saddle point or inflection point. As with the calculus of single variables, determining which of these is true depends on the second derivative. With n variables, the appropriate generalization is the **hessian** matrix

$$\mathbf{H}_{\mathbf{x}}[f] = \nabla_{\mathbf{x}} \left[\left(\nabla_{\mathbf{x}} [f] \right)^T \right] = \frac{\partial^2 f}{\partial \mathbf{x} \partial \mathbf{x}^T} = \begin{pmatrix} \frac{\partial^2 f}{\partial x_1^2} & \cdots & \frac{\partial^2 f}{\partial x_1 \partial x_n} \\ \vdots & \ddots & \vdots \\ \frac{\partial^2 f}{\partial x_1 \partial x_n} & \cdots & \frac{\partial^2 f}{\partial x_n^2} \end{pmatrix} \quad (15.42)$$

This matrix is symmetric, as mixed partials are equal under suitable continuity conditions, and measures the local curvature of the function.

Example 14. Compute $\mathbf{H}_{\mathbf{x}}[\varphi(\mathbf{x})]$, the hessian matrix for the multivariate normal distribution. Recalling from Equation 15.40a that $\nabla_{\mathbf{x}}[\varphi(\mathbf{x})] = -\varphi(\mathbf{x}) \cdot \mathbf{V}_{\mathbf{x}}^{-1}(\mathbf{x} - \boldsymbol{\mu})$, we have

$$\begin{aligned} \mathbf{H}_{\mathbf{x}}[\varphi(\mathbf{x})] &= \nabla_{\mathbf{x}} \left[\left(\nabla_{\mathbf{x}}[\varphi(\mathbf{x})] \right)^T \right] \\ &= -\nabla_{\mathbf{x}} \left[\varphi(\mathbf{x}) \cdot (\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} \right] \\ &= -\nabla_{\mathbf{x}}[\varphi(\mathbf{x})] \cdot (\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} - \varphi(\mathbf{x}) \cdot \nabla_{\mathbf{x}} \left[(\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} \right] \\ &= \varphi(\mathbf{x}) \cdot \left(\mathbf{V}_{\mathbf{x}}^{-1}(\mathbf{x} - \boldsymbol{\mu})(\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} - \mathbf{V}_{\mathbf{x}}^{-1} \right) \quad (15.43a) \end{aligned}$$

Likewise,

$$\mathbf{H}_{\boldsymbol{\mu}}[\varphi(\mathbf{x}, \boldsymbol{\mu})] = \varphi(\mathbf{x}, \boldsymbol{\mu}) \cdot \left(\mathbf{V}_{\mathbf{x}}^{-1}(\mathbf{x} - \boldsymbol{\mu})(\mathbf{x} - \boldsymbol{\mu})^T \mathbf{V}_{\mathbf{x}}^{-1} - \mathbf{V}_{\mathbf{x}}^{-1} \right) \quad (15.43b)$$

To see how the hessian matrix determines the nature of equilibrium points, a slight digression on the multidimensional Taylor series is needed. Consider the Taylor series of a function of n variables $f(x_1, \dots, x_n)$ expanded about the point \mathbf{y} ,

$$f(\mathbf{x}) \simeq f(\mathbf{y}) + \sum_{i=1}^n (x_i - y_i) \frac{\partial f}{\partial x_i} + \frac{1}{2} \sum_{i=1}^n \sum_{j=1}^n (x_i - y_i)(x_j - y_j) \frac{\partial^2 f}{\partial x_i \partial x_j} + \dots$$

where all partials are evaluated at \mathbf{y} . In matrix form, the second-order Taylor expansion of $f(\mathbf{x})$ about \mathbf{x}_o is

$$f(\mathbf{x}) \simeq f(\mathbf{x}_o) + \nabla^T (\mathbf{x} - \mathbf{x}_o) + \frac{1}{2} (\mathbf{x} - \mathbf{x}_o)^T \mathbf{H} (\mathbf{x} - \mathbf{x}_o) \quad (15.44)$$

where ∇ and \mathbf{H} are the gradient and hessian with respect to \mathbf{x} evaluated at \mathbf{x}_o , e.g.,

$$\nabla \equiv \nabla_{\mathbf{x}}[f] \Big|_{\mathbf{x}=\mathbf{x}_o} \quad \text{and} \quad \mathbf{H} \equiv \mathbf{H}_{\mathbf{x}}[f] \Big|_{\mathbf{x}=\mathbf{x}_o}$$

At an equilibrium point $\hat{\mathbf{x}}$, all first partials are zero, so that $(\nabla_{\mathbf{x}}[f])^T$ at this point is a vector of length zero. Whether this point is a maximum or minimum is then determined by the quadratic product involving the hessian evaluated at $\hat{\mathbf{x}}$. Considering vector \mathbf{d} of a small change from the equilibrium point,

$$f(\hat{\mathbf{x}} + \mathbf{d}) - f(\hat{\mathbf{x}}) \simeq \frac{1}{2} \cdot \mathbf{d}^T \mathbf{H} \mathbf{d} \quad (15.45a)$$

Applying Equation 15.16, the canonical transformation of \mathbf{H} , simplifies the quadratic form to give

$$f(\hat{\mathbf{x}} + \mathbf{d}) - f(\hat{\mathbf{x}}) \simeq \frac{1}{2} \sum_{i=1}^n \lambda_i y_i^2 \quad (15.45b)$$

where $y_i = \mathbf{e}_i^T \mathbf{d}$, \mathbf{e}_i and λ_i being the eigenvectors and eigenvalues of the hessian evaluated at $\hat{\mathbf{x}}$. Thus, if \mathbf{H} is positive-definite (all eigenvalues of \mathbf{H} are positive), f increases in all directions around $\hat{\mathbf{x}}$ and hence $\hat{\mathbf{x}}$ is a local minimum of f . If \mathbf{H} is negative-definite (all eigenvalues are negative), f decreases in all directions around $\hat{\mathbf{x}}$ and $\hat{\mathbf{x}}$ is a local maximum of f . If the eigenvalues differ in sign (\mathbf{H} is indefinite), $\hat{\mathbf{x}}$ corresponds to a saddle point (to see this, suppose $\lambda_1 > 0$ and $\lambda_2 < 0$; any change along the vector \mathbf{e}_1 results in an increases in f , while any change along \mathbf{e}_2 results in a decrease in f).