

# Mathematical Derivation of the Two-Step Sigma Matrix Calculation

Clinical Trial Simulation Documentation

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## **Abstract**

This document provides a comprehensive mathematical derivation of the efficient two-step method for computing covariance matrices from correlation matrices in multivariate normal data generation. Specifically, we derive the identity  $\Sigma = \mathbf{DRD}$ , where  $\Sigma$  is the covariance matrix,  $\mathbf{R}$  is the correlation matrix, and  $\mathbf{D}$  is a diagonal matrix of standard deviations. This decomposition is fundamental to clinical trial simulation, enabling numerical stability and computational efficiency.

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# 1 Introduction

In multivariate normal data generation for clinical trials, we need to construct covariance matrices that are:

1. **Positive definite** (PD) - ensuring mvrnorm() and mvn sampling work correctly
2. **Numerically stable** - avoiding ill-conditioning and eigenvalue issues
3. **Interpretable** - separating correlation structure from magnitude (standard deviations)

The most efficient approach achieves all three by decomposing the covariance matrix into:

$$\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$$

where the correlation matrix  $\mathbf{R}$  is validated for PD independently from scaling by standard deviations in  $\mathbf{D}$ .

This document derives this identity from first principles and explains its computational advantages.

## 2 Foundational Concepts

### 2.1 Covariance and Correlation: Univariate Case

**Definition 1** (Covariance). For two random variables  $X$  and  $Y$  with means  $\mu_X$  and  $\mu_Y$ :

$$\text{Cov}(X, Y) = E[(X - \mu_X)(Y - \mu_Y)]$$

**Definition 2** (Standard Deviation). The standard deviation of a random variable  $X$  is:

$$\sigma_X = \sqrt{\text{Var}(X)} = \sqrt{\text{Cov}(X, X)}$$

**Definition 3** (Correlation Coefficient). The Pearson correlation coefficient between  $X$  and  $Y$  is:

$$\text{Corr}(X, Y) = \frac{\text{Cov}(X, Y)}{\sigma_X \sigma_Y}$$

**Theorem 1** (Covariance-Correlation Relationship). For any two random variables  $X$  and  $Y$ :

$$\text{Cov}(X, Y) = \text{Corr}(X, Y) \cdot \sigma_X \cdot \sigma_Y$$

*Proof.* By definition of correlation (Definition 3):

$$\text{Corr}(X, Y) = \frac{\text{Cov}(X, Y)}{\sigma_X \sigma_Y}$$

Rearranging:

$$\text{Cov}(X, Y) = \text{Corr}(X, Y) \cdot \sigma_X \cdot \sigma_Y \quad \square$$

□

## 3 Multivariate Extension

### 3.1 Covariance Matrix

**Definition 4** (Covariance Matrix). Let  $\mathbf{Z} = [Z_1, Z_2, \dots, Z_n]^\top$  be a random vector with mean  $\boldsymbol{\mu}$ . The covariance matrix is:

$$\Sigma = \text{Cov}(\mathbf{Z}) = E[(\mathbf{Z} - \boldsymbol{\mu})(\mathbf{Z} - \boldsymbol{\mu})^\top]$$

with elements:

$$\Sigma_{ij} = \text{Cov}(Z_i, Z_j) = E[(Z_i - \mu_i)(Z_j - \mu_j)]$$

In particular,  $\Sigma_{ii} = \text{Var}(Z_i) = \sigma_i^2$ .

### 3.2 Correlation Matrix

**Definition 5** (Correlation Matrix). The correlation matrix  $\mathbf{R}$  has elements:

$$R_{ij} = \frac{\Sigma_{ij}}{\sigma_i \sigma_j} = \text{Corr}(Z_i, Z_j)$$

By definition,  $R_{ii} = 1$  (perfect self-correlation) and  $|R_{ij}| \leq 1$  for  $i \neq j$ .

### 3.3 Positive Definiteness

**Definition 6** (Positive Definite Matrix). A symmetric matrix  $\mathbf{A} \in \mathbb{R}^{n \times n}$  is positive definite (PD) if:

$$\mathbf{x}^\top \mathbf{A} \mathbf{x} > 0 \quad \forall \mathbf{x} \in \mathbb{R}^n, \mathbf{x} \neq \mathbf{0}$$

Equivalently, all eigenvalues are strictly positive.

**Remark 1** (Valid Correlation Matrices). A valid correlation matrix  $\mathbf{R}$  must be:

1. Symmetric:  $R_{ij} = R_{ji}$
2. Unit diagonal:  $R_{ii} = 1$
3. Bounded:  $|R_{ij}| \leq 1$
4. Positive semi-definite (PSD)

Note: Conditions 1-3 are necessary but NOT sufficient for PSD.

## 4 The Sigma Matrix Decomposition: $\Sigma = \mathbf{D} \cdot \mathbf{R} \cdot \mathbf{D}$

### 4.1 Defining the Diagonal Standard Deviation Matrix

**Definition 7** (Diagonal Standard Deviation Matrix). Define  $\mathbf{D}$  as the diagonal matrix of standard deviations:

$$\mathbf{D} = \begin{pmatrix} \sigma_1 & 0 & \cdots & 0 \\ 0 & \sigma_2 & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & \sigma_n \end{pmatrix}$$

where  $\sigma_i = \sqrt{\Sigma_{ii}}$  is the standard deviation of  $Z_i$ .

## 4.2 Main Derivation

**Theorem 2** (Sigma Decomposition). The covariance matrix  $\Sigma$  can be decomposed as:

$$\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$$

where  $\mathbf{D}$  is the diagonal standard deviation matrix and  $\mathbf{R}$  is the correlation matrix.

*Proof.* We show this by computing each element of  $\mathbf{D}\mathbf{R}\mathbf{D}$ .

### Step 1: Compute $\mathbf{D}\mathbf{R}$

The  $(i, j)$ -element of  $\mathbf{D}\mathbf{R}$  is:

$$[\mathbf{D}\mathbf{R}]_{ij} = \sum_{k=1}^n D_{ik} R_{kj} = D_{ii} R_{ij} = \sigma_i R_{ij}$$

because  $\mathbf{D}$  is diagonal (so  $D_{ik} = 0$  for  $k \neq i$ ).

### Step 2: Compute $(\mathbf{D}\mathbf{R})\mathbf{D}$

The  $(i, j)$ -element of  $(\mathbf{D}\mathbf{R})\mathbf{D}$  is:

$$[(\mathbf{D}\mathbf{R})\mathbf{D}]_{ij} = \sum_{k=1}^n [\mathbf{D}\mathbf{R}]_{ik} D_{kj} = (\sigma_i R_{ij}) D_{jj} = \sigma_i R_{ij} \sigma_j$$

### Step 3: Express in terms of covariance

From Definition 2 and Theorem 1:

$$R_{ij} = \frac{\Sigma_{ij}}{\sigma_i \sigma_j}$$

Therefore:

$$[(\mathbf{D}\mathbf{R})\mathbf{D}]_{ij} = \sigma_i \cdot \frac{\Sigma_{ij}}{\sigma_i \sigma_j} \cdot \sigma_j = \Sigma_{ij}$$

Since this holds for all elements  $(i, j)$ :

$$\mathbf{D}\mathbf{R}\mathbf{D} = \Sigma \quad \square$$

$\square$

### Key Identity

$$\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$$

## 4.3 Element-Wise Illustration

For a  $2 \times 2$  case, the decomposition is explicit:

$$\begin{pmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{pmatrix} \begin{pmatrix} 1 & R_{12} \\ R_{12} & 1 \end{pmatrix} \begin{pmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{pmatrix} = \begin{pmatrix} \sigma_1^2 & \sigma_1 R_{12} \sigma_2 \\ \sigma_1 R_{12} \sigma_2 & \sigma_2^2 \end{pmatrix}$$

where the diagonal elements  $\sigma_i^2$  are the variances and the off-diagonal elements  $\sigma_i R_{ij} \sigma_j$  are the covariances.

## 5 Key Properties

### 5.1 Preservation of Positive Definiteness

**Lemma 1** (PD Preservation). If  $\mathbf{R}$  is positive definite and  $\mathbf{D}$  is diagonal with all positive diagonal entries, then  $\Sigma = \mathbf{DRD}$  is positive definite.

*Proof.* For any non-zero vector  $\mathbf{x}$ :

$$\mathbf{x}^\top \Sigma \mathbf{x} = \mathbf{x}^\top (\mathbf{DRD}) \mathbf{x} = (\mathbf{D}\mathbf{x})^\top \mathbf{R}(\mathbf{D}\mathbf{x})$$

Let  $\mathbf{y} = \mathbf{D}\mathbf{x}$ . Since  $\mathbf{D}$  has positive diagonal entries and  $\mathbf{x} \neq \mathbf{0}$ , we have  $\mathbf{y} \neq \mathbf{0}$ .

Since  $\mathbf{R}$  is positive definite:

$$\mathbf{y}^\top \mathbf{R} \mathbf{y} > 0$$

Therefore:

$$\mathbf{x}^\top \Sigma \mathbf{x} > 0 \quad \square$$

□

**Corollary 1** (Validity of Derived Covariance). If the correlation matrix  $\mathbf{R}$  is positive definite, then the covariance matrix  $\Sigma = \mathbf{DRD}$  is automatically positive definite, regardless of the values in  $\mathbf{D}$  (provided all  $\sigma_i > 0$ ).

### 5.2 Symmetry Preservation

**Lemma 2** (Symmetry Preservation). If  $\mathbf{R}$  and  $\mathbf{D}$  are symmetric, then  $\Sigma = \mathbf{DRD}$  is symmetric.

*Proof.*

$$\Sigma^\top = (\mathbf{DRD})^\top = \mathbf{D}^\top \mathbf{R}^\top \mathbf{D}^\top = \mathbf{DRD} = \Sigma \quad \square$$

□

## 6 Computational Implementation

### 6.1 R Implementation

In R, the efficient computation is:

```
sigma <- outer(standard_deviations, standard_deviations) * correlations
```

where:

- `standard_deviations` is a vector of  $\sigma_1, \dots, \sigma_n$
- `correlations` is the matrix  $\mathbf{R}$
- `outer(x, y)` computes the outer product:  $\mathbf{xy}^\top$
- `*` is element-wise multiplication

## 6.2 Explicit Computation

The `outer()` function computes:

$$\text{outer}(\sigma, \sigma) = \begin{pmatrix} \sigma_1\sigma_1 & \sigma_1\sigma_2 & \cdots & \sigma_1\sigma_n \\ \sigma_2\sigma_1 & \sigma_2\sigma_2 & \cdots & \sigma_2\sigma_n \\ \vdots & \vdots & \ddots & \vdots \\ \sigma_n\sigma_1 & \sigma_n\sigma_2 & \cdots & \sigma_n\sigma_n \end{pmatrix} = \mathbf{D}^2$$

where  $\mathbf{D}^2$  denotes the matrix of all pairwise products.

Then, element-wise multiplication by  $\mathbf{R}$ :

$$\Sigma_{ij} = (\sigma_i\sigma_j) \times R_{ij} = \sigma_i R_{ij} \sigma_j$$

which is exactly the  $(i, j)$ -element of  $\mathbf{DRD}$ .

## 6.3 Computational Advantages

This two-step approach provides several benefits:

1. **Numerical Stability:** Correlation matrices have entries bounded in  $[-1, 1]$ , avoiding overflow/underflow issues when combined with scaling.
2. **PD Validation Efficiency:** Only  $\mathbf{R}$  needs to be checked for positive definiteness. Once  $\mathbf{R}$  is validated (Lemma 1),  $\Sigma$  is automatically PD regardless of  $\sigma_i$  values.
3. **Parameter Separation:** Correlation structure (affecting relationships) and magnitude (affecting variance) are computed independently, simplifying interpretation and parameter selection.
4. **Memory Efficiency:** The computation  $\text{outer}(\sigma, \sigma) \times \mathbf{R}$  avoids explicit construction of the  $n \times n$  matrix  $\mathbf{D}$ .
5. **Reusability:** A single correlation matrix  $\mathbf{R}$  can be scaled with different  $\mathbf{D}$  matrices without recomputation.
6. **Dimension Independence:** For fixed  $\mathbf{R}$ , increasing  $n$  (number of variables) only requires updating the outer product computation.

# 7 Application to Clinical Trial Simulation

## 7.1 General Framework

In the N-of-1 trial simulation, the complete process is:

1. Define correlation parameters following Hendrickson et al.:  $c_{tv}, c_{pb}, c_{br}, c_{cf1t}, c_{cfct}, c_{bm}$
2. Build the correlation matrix  $\mathbf{R}$  from these parameters (with dimension  $\approx 62$  for 20 timepoints  $\times$  3 factors + 2 baseline variables)

3. Validate  $\mathbf{R}$  is positive definite (eigenvalue check)
4. Define standard deviations  $\sigma_1, \dots, \sigma_n$  from response and baseline parameters
5. Compute  $\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$  using the efficient two-step method
6. Sample from  $\mathbf{Z} \sim \mathcal{N}(\boldsymbol{\mu}, \Sigma)$  using `mvrnorm()`

## 7.2 Why This Matters

The decomposition  $\Sigma = \mathbf{D}\mathbf{R}\mathbf{D}$  is critical because:

- High-dimensional correlation matrices ( $62 \times 62$ ) are at risk of numerical failure
- Separating  $\mathbf{R}$  and  $\mathbf{D}$  isolates the PD problem to  $\mathbf{R}$  only
- This allows systematic validation of correlation structures independently of scales
- The final  $\Sigma$  inherits PD guarantee from  $\mathbf{R}$  via Lemma 1

## 8 Example: 3-Variable Case

For illustration, consider three biomarker components at one timepoint:

$$\mathbf{R} = \begin{pmatrix} 1.00 & 0.20 & 0.15 \\ 0.20 & 1.00 & 0.10 \\ 0.15 & 0.10 & 1.00 \end{pmatrix}$$

with standard deviations  $\sigma_1 = 2.0, \sigma_2 = 1.5, \sigma_3 = 2.5$ .

The covariance matrix is:

$$\Sigma = \begin{pmatrix} 2.0 & 0 & 0 \\ 0 & 1.5 & 0 \\ 0 & 0 & 2.5 \end{pmatrix} \begin{pmatrix} 1.00 & 0.20 & 0.15 \\ 0.20 & 1.00 & 0.10 \\ 0.15 & 0.10 & 1.00 \end{pmatrix} \begin{pmatrix} 2.0 & 0 & 0 \\ 0 & 1.5 & 0 \\ 0 & 0 & 2.5 \end{pmatrix}$$

Computing element (1,2):

$$\Sigma_{12} = 2.0 \times 0.20 \times 1.5 = 0.60$$

and the full covariance matrix:

$$\Sigma = \begin{pmatrix} 4.00 & 0.60 & 0.75 \\ 0.60 & 2.25 & 0.375 \\ 0.75 & 0.375 & 6.25 \end{pmatrix}$$

Note that the diagonal elements are  $\sigma_i^2$ :

$$\Sigma_{11} = 2.0^2 = 4.00$$

$$\Sigma_{22} = 1.5^2 = 2.25$$

$$\Sigma_{33} = 2.5^2 = 6.25$$

## 9 Conclusion

The identity  $\Sigma = \mathbf{DRD}$  is a fundamental decomposition that:

1. **Theoretically** cleanly separates correlation structure from variance scaling
2. **Computationally** enables efficient two-step calculation and validation
3. **Numerically** ensures positive definiteness when  $\mathbf{R}$  is PD
4. **Practically** allows systematic parameter selection and validation before expensive simulations

In the context of N-of-1 trial simulation, this decomposition enables robust construction of high-dimensional covariance matrices while maintaining strict control over positive definiteness constraints.

## A Matrix Algebra Reference

### A.1 Outer Product

For vectors  $\mathbf{u}, \mathbf{v} \in \mathbb{R}^n$ :

$$\text{outer}(\mathbf{u}, \mathbf{v}) = \mathbf{uv}^\top$$

### A.2 Diagonal Matrix Properties

For diagonal matrix  $\mathbf{D}$  with diagonal entries  $d_1, \dots, d_n$  and any matrix  $\mathbf{A}$ :

$$\begin{aligned} (\mathbf{DA})_{ij} &= d_i A_{ij} \\ (\mathbf{AD})_{ij} &= A_{ij} d_j \end{aligned}$$

### A.3 Matrix Transpose Properties

$$\begin{aligned} (\mathbf{AB})^\top &= \mathbf{B}^\top \mathbf{A}^\top \\ (\mathbf{A}^\top)^\top &= \mathbf{A} \\ (\mathbf{D}^\top) &= \mathbf{D} \quad (\text{if } \mathbf{D} \text{ is diagonal}) \end{aligned}$$

### A.4 Eigenvalue Properties

For positive definite matrix  $\mathbf{A}$ :

- All eigenvalues are strictly positive:  $\lambda_i > 0$
- Condition number:  $\kappa(\mathbf{A}) = \frac{\lambda_{\max}}{\lambda_{\min}}$
- Matrix is invertible:  $\det(\mathbf{A}) > 0$