Probability & Distributions

Multivariate Distributions & Moments

if X_1, \ldots, X_n are **i.i.d.** with *CDF* F(x) then:

$$F_{X_{\min}}(x) = 1 - [1 - F(x)]^n$$

$$F_{X_{\text{max}}}(\mathbf{x}) = [F(\mathbf{x})]^n$$

For random vector $X \in \mathbb{R}^p$, covariance matrix:

$$\Sigma = \text{Cov}(X) = \mathbb{E}[(X - \mathbb{E}[X])(X - \mathbb{E}[X])^T] \in \mathbb{R}^{p \times p}$$

Properties:

- if a is constant vector then: $Var(a^TX) = a^T\Sigma a$
- if $X \perp \!\!\! \perp Y$ then Var(X + Y) = Var(X) + Var(Y)

Correlation coefficient:

$$\operatorname{Corr}(X_i, X_j) = \frac{\operatorname{Cov}(X_i, X_j)}{\sqrt{\operatorname{Var}(X_i)\operatorname{Var}(X_j)}}$$

Multivariate Normal (MVN):

 $X \sim N_p(\mu, \Sigma)$ means X has density

$$f_X(x) = \frac{1}{(2\pi)^{p/2} |\Sigma|^{1/2}} \exp\{-\frac{1}{2}(x-\mu)^T \Sigma^{-1}(x-\mu)\}.$$

Properties:

- Any linear combination of components of *X* is normal.
- If $A \in \mathbb{R}^{k \times p}$, then $Y = AX \sim N_k(A\mu, A\Sigma A^T)$.
- Marginals of a MVN are normal: any subset X_I is $N_{|I|}(\mu_I, \Sigma_{II})$.

• If
$$X = \begin{pmatrix} X_1 \\ X_2 \end{pmatrix} \sim N(\begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, \begin{pmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{21} & \Sigma_{22} \end{pmatrix})$$
, then the conditional distribution $X_1 \mid X_2 = x_2$ is:

$$N(\mu_1 + \Sigma_{12}\Sigma_{22}^{-1}(x_2 - \mu_2), \Sigma_{11} - \Sigma_{12}\Sigma_{22}^{-1}\Sigma_{21}),$$

$$E[X_1|X_2] = \mu_1 + \Sigma_{12}\Sigma_{22}^{-1}(X_2 - \mu_2).$$

• The **MGF** of $X \sim N_p(\mu, \Sigma)$ is:

$$M_X(t) = \exp(t^T \mu + \frac{1}{2} t^T \Sigma t).$$

Independence:

Components of X are independent $\iff \Sigma$ is diagonal (for MVN, uncorrelated ⇒ independent).

Mahalanobis: If $X \sim N_p(\mu, \Sigma)$, then

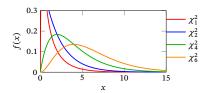
$$(X - \mu)^T \Sigma^{-1} (X - \mu) \sim \chi_p^2$$
 and $\Sigma^{-1/2} (X - \mu) \sim N_p(0, I_p)$.

Multivariate Chi-squared Distribution:

If $X \sim N_p(0, I_p)$, then $\chi^2 = X^T X = \sum_{i=1}^p X_i^2 \sim \chi_p^2$. For non-central case, if $X \sim N_p(\mu, I_p)$, then $X^T X \sim \chi_p^2(\lambda)$ with non-centrality $\lambda = \mu^T \mu$.

- $\bullet E[\chi_p^2] = p, Var(\chi_p^2) = 2p$
- For independent $\chi_{p_1}^2$, $\chi_{p_2}^2$: $\chi_{p_1}^2 + \chi_{p_2}^2 \sim \chi_{p_1+p_2}^2$ MGF: $M_{\chi_p^2}(t) = (1-2t)^{-p/2}$ for t < 1/2

Connection to quadratic forms: If $Z \sim N_p(0, I_p)$ and A is a symmetric idempotent matrix of rank r, then $\mathbf{Z}^T A \mathbf{Z} \sim \chi_r^2$.



Principal Component Analysis (PCA):

For data with covariance matrix Σ (or correlation R), find eigenvalues $\lambda_1 \geq$ $\lambda_2 \geq \ldots \geq \lambda_p$ and eigenvectors e_1, \ldots, e_p (orthonormal) solving $\Sigma e_i = \lambda_i e_i$.

- The *i*th PC is $Z_i = e_i^T(X \bar{X})$ with $Var(Z_i) = \lambda_i$
- PCs are uncorrelated (orthogonal directions of maximal variance)
- Proportion of variance explained by first k PCs: $\frac{\lambda_1 + \dots + \lambda_k}{\lambda_1 + \dots + \lambda_n}$

Quadratic Forms & Idempotent Matrices:

Idempotent Matrices: A matrix *H* is *idempotent* if $H^2 = H$.

- Idempotent H has eigenvalues 0 or 1 only
- \cdot rank(H) = tr(H) (number of eigenvalues = 1)

Quadratic Forms: If $Z \sim N(0, I_n)$ and P is symmetric idempotent of rank r,

$$Q = Z^T P Z \sim \chi_r^2$$

$$Z \sim N(0, \sigma^2 I_n) \implies Q/\sigma^2 \sim \chi_r^2$$

Independence: If P_1 and P_2 are symmetric idempotent with $P_1P_2=0$ (projections onto orthogonal subspaces), then $Z^T P_1 Z$ and $Z^T P_2 Z$ are independent.

Linear Model Setup:

Assume $Y = X\beta + \varepsilon$ where Y is $n \times 1$, X is $n \times p$ (full rank p), β is $p \times 1$ of unknown parameters, and $\varepsilon \sim N(0, \sigma^2 I_n)$ (errors independent, homoscedastic, normal). Assumptions: linear relationship, $E[\varepsilon] = 0$, $Var(\varepsilon) = \sigma^2 I$, independent errors (normality for inference). Under these assumptions,

$$Y \sim N(X\beta, \sigma^2 I)$$
.

The ordinary least squares (OLS) estimator is:

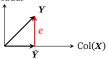
$$\begin{split} \hat{\beta} &= (X^T X)^{-1} X^T Y, \quad \begin{cases} E[\hat{\beta}] &= \beta & \text{(unbiased)} \\ \text{Var}(\hat{\beta}) &= \sigma^2 (X^T X)^{-1} \\ \text{SE}(\hat{\beta}) &= \sqrt{\text{diag}(\text{Var}(\hat{\beta}))} = \sigma \sqrt{(X^T X)_{jj}^{-1}} & \text{(std. error)} \end{cases} \end{split}$$

solving the normal equations $X^{T}(Y - X\hat{\beta}) = 0$. The *fitted values* are,

$$\hat{Y} = X\hat{\beta} = HY.$$

Hat matrix $H = X(X^TX)^{-1}X^T$ is symmetric and idempotent (rank p). The residuals are $e = Y - \hat{Y} = (I - H)Y$, where (I - H) is symmetric idempotent (rank

Residual



Gauss-Markov theorem: $\hat{\beta}$ is the best linear unbiased estimator (minimal variance). If ε is normal, then $\hat{\beta} \sim N(\beta, \sigma^2(X^TX)^{-1})$.

$$\hat{\mathbf{Y}} = H\mathbf{Y} \sim N(\mathbf{X}\boldsymbol{\beta}, \ \sigma^2 H), \quad e = (I - H)\mathbf{Y} \sim N(0, \ \sigma^2 (I - H)).$$

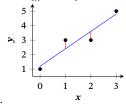
Moreover, $\hat{\mathbf{Y}}$ and e are independent (since H(I-H)=0). If an *intercept* (β_0) is included,

$$SST = \sum_{i} \underbrace{(\hat{\mathbf{Y}}_{i}^{i} - \bar{\mathbf{Y}})^{2}}_{SSE} + \sum_{i} \underbrace{(\hat{\mathbf{Y}}_{i}^{i} - \hat{\mathbf{Y}}_{i}^{i})^{2}}_{SSE} = \sum_{i} (\mathbf{Y}_{i}^{i} - \bar{\mathbf{Y}})^{2}$$
(1)

$$R^2 = \frac{SSR}{SST} = 1 - \frac{SSE}{SST}, \quad R_{adj}^2 = 1 - \frac{n-1}{n-p-1} (1 - R^2)$$
 (2)

$$\mathrm{df}_{\mathrm{SSR}} = p-1, \quad \mathrm{df}_{\mathrm{SSE}} = n-p, \quad \mathrm{df}_{\mathrm{SST}} = n-1$$
 (DoF)

Under normal errors, $SS_{err}/\sigma^2 \sim \chi^2_{n-p}$ then SS_{err} is independent of $\hat{\beta}$. An



unbiased estimator of σ^2 is $\hat{\sigma}^2 = \frac{1}{n-n} SS_{err}$.

Inference in Linear Model:

For each parameter β_i , under H_0 : $\beta_i = 0$,

$$t = \frac{\hat{\beta}_j}{\operatorname{SE}(\hat{\beta}_j)} \sim t_{n-p}.$$

Thus $(1 - \alpha)$ CI for β_i is:

$$\hat{\beta}_j \pm t_{n-p, \, \alpha/2} \, \text{SE}(\hat{\beta}_j)$$

For a new predictor value x_0 , the predicted response $\hat{y}_0 = x_0 \hat{\beta}$ has $\text{Var}(\hat{y}_0) = \sigma^2 x_0 (X^T X)^{-1} x_0^T$. A $(1-\alpha)$ CI for the mean at x_0 is $\hat{y}_0 \pm$ $t_{n-p, \alpha/2}$ $\hat{\sigma}\sqrt{x_0(X^TX)^{-1}x_0^T}$, and a prediction interval for a new Y at x_0 is $\hat{y}_0 \pm t_{n-p, \alpha/2} \hat{\sigma} \sqrt{x_0 (X^T X)^{-1} x_0^T + 1}$.

Hypothesis Testing:

F-test:

For general linear hypothesis $H_0: L\beta = 0$ where L is a $d \times p$ matrix of rank d:

$$F = \frac{(L\hat{\beta})^T [L(\boldsymbol{X}^T\boldsymbol{X})^{-1}L^T]^{-1}(L\hat{\beta})}{\hat{\sigma}^2} \sim F_{d,n-p}$$

- Testing $H_0: \beta_j = 0: t^2 = F_{1,n-p}$
- Global test $H_0: \beta_1 = \cdots = \beta_{p-1} = 0: F = \frac{\text{SSR}/(p-1)}{\text{CSE}/(p-p)} \sim F_{p-1,n-p}$

Reject H_0 if $F > F_{d,n-p,\alpha}$.

Model Selection:

Common criteria: AIC = $n \ln(SSE/n) + 2p$, BIC = $n \ln(SSE/n) + p \ln n$ (smaller is better). Mallows' $C_p \approx \frac{\text{SSE}_{\text{model}}}{\sigma_{(\text{full})}^2} - (n-2p)$, targeting $C_p \approx p$.

Diagnostics:

Residual standard deviation: $\hat{\sigma} = \sqrt{SSE/(n-p)}$. Check residual plots for constant variance (no patterns in residuals vs fitted) and for normality (e.g. Q-Q plot). High-leverage points: $h_{ii} = H_{ii}$; large h_{ii} (relative to p/n) indicates an plot). High-leverage points, $n_{ii} - n_{il}$, $n_{ij} = n_{il}$, $n_{ij} = n_{il}$ outlier in predictor space. Standardized residual $r_i = \frac{e_i}{\hat{\sigma} \sqrt{1 - h_{il}}}$ (should be

 $\approx N(0,1)$ under the model). Outliers may have $|r_i| > 2$ or 3. Influence can be assessed by Cook's distance: $D_i = \frac{e_i^2}{p \, \sigma^2} \, \frac{h_{ii}}{(1 - h_{ii})^2}$ (values $D_i > 1$ are often considered large). If assumptions are violated (nonlinearity, heteroscedastic-

ity, non-normal errors), consider remedies such as transforming variables or using a different model. For variance stabilization, choose g(y) such that Var(g(Y)) is roughly constant. E.g. for Poisson $Y(Var \mu)$, use $g(Y) = \sqrt{Y}$; for Binomial proportion (Var $\approx \mu(1-\mu)$), use $g(Y) = \arcsin \sqrt{Y}$; for Var $\propto \mu^2$, use $\log Y$ (Box–Cox power transform can find an optimal λ).

Multiple Testing:

When performing m hypothesis tests, the family-wise error rate (FWER) at level α is Pr(any false rejection) $\leq \alpha$.

Bonferroni correction: $\alpha = \frac{\alpha}{m}$ for each test (controls FWER $\leq \alpha$).

Šidák correction: $\alpha = 1 - (1 - \alpha)^{1/m}$.

Variance-Stabilising Transformations: Step-by-Step Guide

Given a random variable Y with mean $\mu = \mathbb{E}[Y]$ and variance $\text{Var}(Y) = v(\mu)$, a variance–stabilising transformation g is obtained as follows:

$$\mu = \mathbb{E}[Y], \quad v(\mu) = \text{Var}(Y)$$

$$Var[g(Y)] \approx [g'(\mu)]^2 v(\mu) = C$$
(4)

$$g'(\mu) = \frac{\sqrt{C}}{\sqrt{v(\mu)}}\tag{5}$$

(3)

$$g(y) = \sqrt{C} \int \frac{dy}{\sqrt{v(y)}} \tag{6}$$

Typically choose C = 1 and drop the additive constant C_0 .

ANOVA (Analysis of Variance):

One-way ANOVA with a groups (levels) and total N observations: model $Y_{ij} = \mu + \tau_i + \varepsilon_{ij}$, for $i = 1, \ldots, a$ and $j = 1, \ldots, n_i$, where τ_i is the effect of group i (with $\sum_i \tau_i = 0$) and $\varepsilon_{ij} \sim N(0, \sigma^2)$. Null hypothesis $H_0: \tau_1 = \tau_2 = \cdots = \tau_a = 0$ (all group means equal) is tested by

$$F = \frac{\text{SSB}/(a-1)}{\text{SSW}/(N-a)} \sim F_{a-1,N-a},$$

where SSB = $\sum_{i=1}^a n_i (\bar{Y}_i - \bar{Y})^2$ (between-group SS) and SSW = $\sum_{i=1}^a \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_i)^2$ (within-group SS). Total SST = SSB + SSW with df N-1=(a-1)+(N-a). If H_0 is rejected, follow-up with multiple-comparison tests (adjusting for multiple testing). Two-way ANOVA (two factors) and factorial designs partition variability into main effects and interaction effects similarly, each tested with an F-ratio of mean squares.

Design of Experiments (DOE):

In a 2^k full factorial design, k factors each have 2 levels (often coded -1 and +1). All 2^k combinations are run (possibly with replication). The *main effect* of factor A is the difference in average response between high and low levels of A; an *interaction* effect (e.g. AB) is the difference in A's effect between the

two levels of B. For example, A effect = $\bar{Y}(A^+) - \bar{Y}(A^-)$, and AB interaction = $[(\bar{Y}_{A+B+} - \bar{Y}_{A-B+}) - (\bar{Y}_{A+B-} - \bar{Y}_{A-B-})]$. Effects can be estimated via a regression model $Y = \beta_0 + \sum \beta_i x_i + \sum \beta_{ij} x_i x_j + \cdots$ with $x_i = \pm 1$. (In this coding, β_i equals half the main effect for factor i, etc.) Orthogonal designs: in a full factorial, the design matrix columns for each effect are orthogonal, simplifying estimation and interpretation (no confounding among effects). Blocking: to account for nuisance variables, experiments may be divided into blocks. In a 2^k design with 2 blocks, one effect (usually a high-order interaction) is confounded with the block factor (indistinguishable from a block effect). E.g. in a 2³ design, to run in 2 blocks we can confound the ABC interaction with blocks by assigning all runs with ABC = +1 to Block 1 and ABC = -1 to Block 2. Then any systematic block difference will appear as an ABC effect (and vice versa). Fractional factorial 2^{k-p} designs run a fraction of the 2^k runs. Specified by *p generators* (defining relations). E.g. a 2^{3-1} half-fraction with generator I = ABC means we run only combinations where ABC = +1. This yields aliasing: A is aliased with BC, B with AC, C with AB. (Defining relation I = ABC; resolution III design since smallest alias involves 3 factors.) Design resolution R: no effect involving < R factors is aliased with any other effect with < Rfactors. Higher resolution designs reduce confounding (e.g. resolution IV: no main effect aliased with any other main or 2-factor effect; resolution V: no main or 2-factor aliased with any other up to 2-factor effect, etc.). To de-alias key effects, one can run fold-over (the complementary fraction) or combine fractional designs to a higher resolution.