

**Handout 1: Symmetric Matrices:** rows and columns are interchangeable.  $A = A^T$ . **Diagonal Matrix:** all off-diagonal elements are zero  $\text{diag}()$  func in R. **0 and I Matrices:**  $AI = IA = A$ . **Equal Matrices:** same dimensions and corresponding elements are equal. **Matrix Multiplication:**  $AB \neq BA$  in general; inner dimensions must match. **Distributive Property:**  $A(B + C) = AB + AC$ . **Transpose of Products:**  $(AB)^T = B^T A^T$ .

**Handout 2: Sum of elements:**  $\sum_{i=1}^n a_i = j^T a$  where  $j$  is a vector of 1s. **Sum of squares:**  $\sum_{i=1}^n a_i^2 = a^T a$  also called the dot product of  $a$  and  $a^T$ . **Length of 2-dim vector:**  $\sqrt{a^T a}$ . **Quadratic Form:**  $y^T A y$  where  $A$  is symmetric matrix and  $y$  is a  $n \times 1$  vector and this returns a scalar. **Linearly Dependent:**  $c_1 v_1 + \dots + c_n v_n = 0$  where  $c_i$  are not all zero. **Rank:** number of linearly independent columns/rows;  $\leq \min(n, p)$ ;  $= \min(n, p)$  if full rank; get to row-echelon form and count pivot rows for rank. **Non-singular: square matrix with full rank; less than full rank is singular; non-singularity = inverse.** **Inverse:**  $A^{-1} A = A A^{-1} = I$ ;  $(AB)^{-1} = B^{-1} A^{-1}$ ;  $(A^T)^{-1} = (A^{-1})^T$ . **Positive Definite Matrix:**  $x^T A x > 0$ ,  $A = T^T T$  then  $T$  is square root matrix of  $A$ ; PDM is also full ranks and has an inverse;  $[x1x2]A[x1x2]^T > 0$  where  $A$  is  $\text{c}(2, 0, 2, 0)$ ,  $\text{ncol}=2$ ,  $\text{nrow}=2$ . **Determinant:** Scalar;  $\det()$  or  $|A|$ ; **computed for var/cov matrix** which is always square, symmetric, and PDM; diagonal elements  $s_i^2$  and off-diagonal elements  $s_{ij}$ ;  $|A| = ad - bc$  for  $\text{c}(a, b, c, d)$ ; for  $3 \times 3$  matrix  $|A| = a(ei - fh) - b(di - fg) + c(dh - eg)$ . **Props of Det:**  $A$  is diagonal then  $|A| = \prod_{i=1}^n a_{ii}$  (product of diagonals); singular then  $\det = 0$ ; non-singular then  $\det \neq 0$  **has an inverse**; PDM then  $\det > 0$ ; **Trace:** sum of diagonal elements for square matrix. **Orthogonal Matrix:** dot product  $= 0$ ; correlation coeff.  $r = 0$  if Orthogonal and linearly independent; numerator of Pearson should be 0, so  $x_c^T y_c^T = 0$ ; **Normalizing a vector:**  $x_c = x / \sqrt{x^T x}$ . **Orthogonal Matrix Properties:** matrix  $A$  is orthogonal matrix if it is a square matrix with orthonormal (unit length) rows and columns;  $A^T A = A A^T = I$  which means  $A^T = A^{-1}$ ; **Eigenvalues and Eigenvectors:**  $Ax = \lambda x$ , where  $A$  defines a transformation and  $x$  is not affected by it;  $A$  is  $n \times n$  then  $n$  distinct eigenvalues; solve  $|A - \lambda I| = 0$ ; Got eigenvalues, plug in and solve for  $x$ , to make it length 1 divide by  $\sqrt{2}$ ; **Eigen Properties:** eigenvalues of a PDM are all positive; eigenvectors of a symmetric matrix are all orthogonal:  $x_i^T x_j = 0$  for  $i \neq j$ ;  $\text{trace}(A) = \text{sum of eigenvalues}$ ;  $\det(A) = \text{product of eigenvalues}$ ; present eigenvalues **in descending order**.

**Handout 3: Sample Mean Vector:**  $\text{colMeans}()$ ; estimate of the population mean vector. **Sample Variance:**  $s^2 = \frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2$  so the distance from average, on average;  $\text{cov}()$  for var/cov matrix. **Population Variance:**  $\sigma^2 = E[(y - \mu)^2]$ . **Covariance:**  $s_{jk} = \frac{1}{n-1} \sum_{i=1}^n (y_j - \bar{y}_j)(y_k - \bar{y}_k)$ ; large & positive strong, linear positive association; lose to 0 is a weak linear assoc. **Correlation Coeff and Corr Matrix:** corr coeff is a scaled covariance so it is more interpretable; sample corr coeff:  $r_{jk} = \frac{s_{jk}}{\sqrt{s_{jj}}\sqrt{s_{kk}}}$  and same for population;  $\text{cor}()$  for correlation matrix, has 1 on diagonal,  $\rho_{jk} = \rho_{kj}$ , **between -1 and 1 and no units**;

**Handout 4: Linear Combination:**  $z = c_1 x_1 + c_2 x_2$ ;  $z = c^T x$ ; **Sample Mean :** Vector of  $z$ :  $\bar{z} = c^T \bar{x}$ . **Interpretation:** sample overall grade  $\bar{z}$  across the  $n$  students is the linear combination of the sample means of the various different components. **Population Mean:**  $E(z) = c^T \mu$ . **Variance:**  $\text{Var}(z) = c_1^2 \text{var}(x_1) + c_2^2 \text{var}(x_2) + 2c_1 c_2 \text{cov}(x_1, x_2)$ ;  $s_z^2$  for sample. **Interpretation:** variance of sum of 2 random vars must take into account cov between all pairs of random vars:  $\text{var}(x_1 + x_2) = \text{formula above}$ . **Sample Variance:**  $s_z^2 = A^T S A$  where  $S$  is the var/cov matrix. **Population Variance:**  $\sigma_z^2 = c^T \Sigma c$ . **Several Linear Combinations:**  $Z = CX$ ;  $E(Z) = C\mu$ ;  $\text{Var}(Z) = C\Sigma C^T$ ;  $z = Ay$  and same equations as above. **Partitioning:** sample covariance matrix has transpose of each other within the partitioned matrix. **Number of Elements:**  $\frac{n(n+1)}{2}$  unique elements in a cov or corr matrix of  $n$  variables. **Generalized Sample Variance:**  $|S|$  is product of eigenvalues of  $S$ ,  $|S| = 0$  if any eigenvalue is 0 (multi co-linearity). **Mahalanobis distance:** multivariate version of a z-score;  $\delta^2 = (y_i - \mu)^T \Sigma^{-1} (y_i - \mu)$  is in quadratic form and return s a scalar; **variables that have a high degree of variation will contribute less to the overall Mahalanobis distance**; how far a measurement is from the mean vector relative to what a typical deviation from the mean is. **Multivariate Normal Distribution:**  $f(y) = \frac{1}{(2\pi)^{p/2} |\Sigma|^{1/2}} \exp(-\frac{1}{2}(y - \mu)^T \Sigma^{-1} (y - \mu))$ ;  $y \sim N_p(\mu, \Sigma)$ ; returned value should be a scalar. **Properties of Multivariate Normal:**  $y$  is  $N_p(y_i, \Sigma)$ ;  $z \sim N(a^T \mu, a^T \Sigma a)$  and individual  $y$ 's must be normal too;  $\text{rank}(A) = q \leq p$  then  $z \sim N_q(A\mu, A^T \Sigma A)$ ; if  $\mu = 0$  and  $\Sigma = I$  then  $Ay \sim N(0, I)$ . **Independent Random Variables:**  $y_j$  and  $y_k$  are independent if and only if covariance  $s_{jk} = 0$  or stated in terms of correlation of row  $jk$ ; **Implication goes both ways since property of MV normal distribution**.

**Handout 5: Disprove p vars are jointly MV normally distributed is to show that at least 1 y is not uni-variate normal.**; using qq-plot or rigorous tests; null hypothesis of these tests is that variables do follow a MV normal distribution. **Univariate CLT:** Typically  $\bar{y}$  is within  $\frac{\sigma}{\sqrt{n}}$  of  $\mu$  and follows a normal distribution. **Multivariate CLT:**  $\bar{y} \sim N_p(\mu, \frac{\Sigma}{n})$  for a large enough sample size  $n$ . **Sample Variance and Cov Matrix Distribution:** univariate follows a chi-squared distribution; sample var/cov matrix follows a Wishart distribution:  $(n-1)S \sim \text{Wishart}(n-1, \Sigma)$  if  $y$  follows a MV normal distribution. **Univariate 1-Sample T Test:**  $t = \frac{\bar{y} - \mu_0}{s/\sqrt{n}}$  nominator is how far  $\bar{y}$  is from the hypothesized pop mean if  $H_0$  were true; denominator is relative to a typical deviation of  $\bar{y}$  from  $\mu$ . **Hotelling's 1-Sample  $T^2$  Test:**  $T^2 = n(\bar{y} - \mu_0)^T S^{-1} (\bar{y} - \mu_0)$ ; measures how far observed  $\bar{y}$  is from its expected value, if the null were true, while taking into account the variance/cov of the sample mean vector. **Hotelling's  $T^2$  Density:** no upper bound; reject when  $T^2$  is large. **P-value accuracy:** data values must have been sampled from MV normal dist,  $S$  must be non-singular, and  $n > p$  (observations  $>$  variables). **Steps to Take After Rejecting Null:** check multivariate normality of data, conduct univariate tests on each variable. **Benefits of MV Tests:** using  $p$  univariate tests inflates the type I error rate (rejecting null when it is true);  $p = 4$  and  $\alpha = 0.05$  then  $1 - (1 - 0.05)^4 = 0.19$  probability and quickly increases with  $p$ ; does not ignore correlation structure between  $p$  vars; more power (prob of rejecting null when null is true) and high power is good; small deviations may be statistically significant when combined. **Limitations of MV Tests:** interpretations difficult without univariate tests when statistically significant MV test; **non-directional** so null hypothesis for MV is 2-tailed.

**Handout 6:**

- **Univariate Two-Sample T Test:** null and alt hypoth; t stat and pooled variance; assuming equal variances; validity of p-val is sampled from normal dist with equal var but possibly diff means
- **Multivariate Two-Sample Hotelling's T Test:** null and alt; t stat (maha dist); pooled sample cov; validity pval is MV normal dist with equal cov matrices but diff means;  $y_{ij}$  is the  $j$ th obs in group  $i$  and vectors with  $p$  entries; individual t test to see where differences lie.
- **Paired Multivariate Data:** Difference = Treatment - Control; multivariate and paired (same subject); apply 1-sample Hotelling's T test; null is mean diff 0, alt is mean diff not 0; t stat is maha between  $\bar{d}$  and 0; validity is mv normal; follow up univariate analysis to see which dist from 0 is significant;

Handout 7:

- **Univariate 1-Way ANOVA:**  $y_i$  is sum across all measurements in sample  $i$ ,  $\bar{y}_i$  is sample mean of all measurements in sample  $i$ ,  $\bar{y}$  is (grand) sample mean of all measurements across all samples; statistical model for data, assume ...; null and alt; estimate of  $\sigma^2$  using within variation MSE (within group); MSE thought of as pooled estimate of  $\sigma^2$ ; between sample variation is MSH; F stat eq, explained var / unexplained var (within is noise and between is signal); Reject when F large; look at graphs, sd of within group (noise).
- **1-Way MANOVA:** vector  $y_{ij}$  has  $y_{ij1} \dots y_{ijp}$ ; statistical model with  $y_i$  being estimate of  $\mu$ ; null and alt hypoth; between var is H eq, within var is E eq;  $E + H$  is total sample covariance; dets of each give generalized vars; Wilk's  $\Lambda$  stat which is unexplained or noise / total; reject when  $\Lambda$  is small; get F by transforming and when large reject null. `manova()` then summary with Wilk's test, `summary$SS` gives H and `$Residuals` gives E; Follow up with univariate anovas to see which var has diff means across groups; `TukeyHSD()` with Bonferroni correction finds which groups are different, p adj is bumped up little so hard to reject null, protect from making type I error, compare p-val to alpha / num of vars to see which are significant, can compare sample means this way.

Handout 8:

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