

## MATH-338 Midterm 2 Cheat Sheet

### THEORY

**Day 14:** probability density function is represented an integral with function  $f(x)$ . Our probability lies within the curve and is always 1. Density curve  $\rightarrow$  bell curve. Z-Score allows us to have a universal standard for density curves with different scales. They are directly proportional to the standard deviation and the delta from the mean of the graph.

**Day 15:** unimodal: one hump, bimodal: two humps. Mean is resistant whereas the mean is subject to change. Density curves decay to histograms (integral  $\rightarrow$  to Reimann Sum). Whisker plots are an effective method to determine if a data set contains outliers (data points not belonging to the sample set)

**Day 16:** error: since there is some error while taking sample data, we do allow for some buffer. We also do not measure exact but to a tolerance which is influenced by the buffer above. Central Limit Theorem: when population size is "large enough"  $\bar{x}$  is an approximation. Higher skew and outliers suggest a larger  $n$  value.

**Day 18:** As  $n \uparrow$ ,  $SEM \downarrow$ . Two-tailed tests take the upper and lower limit of the curve and the significance level ( $\alpha$ ) is the cut off point of being *statistically significant*. Treat as critical region. If in CR, then accept alt. Else accept null.

**Day 19:**  $H_1: \mu < \mu_0 \leftarrow$  left side.  $H_1: \mu \neq \mu_0 \leftarrow$  n  $\sigma$  on both side but no middle.  $H_1: \mu > \mu_0 \leftarrow$  lower.tail = TRUE. Population distribution normality  $\implies$  sample population distribution normality. Matched pairs design: paired subjects receives their respective treatment or an individual gets two treatments. Also a subset of block design. Common hypothesis:  $H_0: \mu_d = 0$  (no difference) and  $H_A: \mu_d \neq 0$  (difference). Matched-pairs t-test reqs: large population, normal distribution,  $\sigma$  is unknown.

**Day 20:** Two Independent Samples t-Test: two unrelated treatments into one numerical response variable measured in two independent groups. Two different  $\mu_1$  and  $\mu_2$ . NHST approach; identify  $\mu_i$

### FORMULAS

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| <ul style="list-style-type: none"><li>• <math>\square = width \times \frac{1}{width}</math> (finite curve)</li><li>• <math>Z = \frac{x-\mu}{\sigma}</math> (z-score)</li><li>• <math>X \sim N(\mu, \sigma)</math></li><li>• <math>\bar{X} \sim N(\mu, \frac{\sigma}{\sqrt{n}})</math></li><li>• <math>SEM = \frac{s}{\sqrt{n}}</math></li><li>• <math>t = \frac{\bar{X}-\mu}{\frac{s}{\sqrt{n}}}</math></li><li>• <math>t = \frac{\bar{x}}{\frac{s}{\sqrt{n}}}</math></li><li>• <math>t = \frac{(\bar{x}_1 - \bar{x}_2)}{\sqrt{\frac{(s_1)^2}{n_1} + \frac{(s_2)^2}{n_2}}} \sim t(K)</math> [NHST]</li><li>• <math>df = n - 1</math></li></ul> | <ul style="list-style-type: none"><li>• <math>IQR = Q_3 - Q_1</math></li><li>• <math>K = 1.5</math></li><li>• Lower fence: <math>Q_1 - K \times IQR</math></li><li>• Upper fence: <math>Q_3 + K \times IQR</math></li></ul> |
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