

Statistical Inference Course Project - Part 2

In this project we will analyze the ToothGrowth data library(datasets) data(ToothGrowth)

This datasets consists of 60 observations and three variables. Variables are len (numeric: tooth length increase), supp (categorical: VC or OJ) and dose (numerical: dose level - 0.5mg, 1mg or 2mg) look at summary(ToothGrowth) summary(ToothGrowth) table(ToothGrowthsupp, as.factor(ToothGrowthdose))

We will plot boxplots for each suppliment type (red are OJ, blue are VC) and dosage vs tooth length increase boxplot(len supp*dose, data = ToothGrowth, col = c("red", "blue"), main = "Tooth Growth", xlab ="Suppliment type and dose mg", ylab = "tooth length increase in mm")

```
95for(i in levels(ToothGrowthsupp))for(j in unique(ToothGrowthdose)) x <-
ToothGrowthlen[ToothGrowthsupp==i ToothGrowthdose == j]print(paste0("Supplimenttype", i, "and", j,
-1*qnorm(0.975)*sd(x)/sqrt(length(x)), 2), "", round(mean(x)+1*qnorm(0.975)*
sd(x)/sqrt(length(x)), 2)))
```

We can see that as dosage increases, tooth length also increases We can also prove it performing t-tests

```
ToothGrowth.dose0.5 = subset(ToothGrowth, dose == 0.5) ToothGrowth.dose1.0=
subset(ToothGrowth, dose == 1) ToothGrowth.dose2.0= subset(ToothGrowth,
dose == 2) doseEffect0.5to1.0 < -t.test(ToothGrowth.dose0.5len, ToothGrowth.dose1.0len)doseEffect0.5to1.0
doseEffect1.0to2.0 < -t.test(ToothGrowth.dose1.0len, ToothGrowth.dose2.0len)doseEffect1.0to2.0.095
```

Now lets compare the tooth length increase between different suppliment types. ToothGrowth.typeOJ = subset(ToothGrowth, supp == "OJ") ToothGrowth.typeVC = subset(ToothGrowth, supp == "VC") typeEffect <- t.test(ToothGrowth.typeOJlen, ToothGrowth.typeVClen) typeEffect The 95However p-value is only 6

```
ToothGrowth.typeOJ.dose0.5 = subset(ToothGrowth, supp == "OJ" dose==0.5)
ToothGrowth.typeOJ.dose1.0 = subset(ToothGrowth, supp == "OJ" dose==1.0)
ToothGrowth.typeOJ.dose2.0 = subset(ToothGrowth, supp == "OJ" dose==2.0)
ToothGrowth.typeVC.dose0.5 = subset(ToothGrowth, supp == "VC" dose==0.5)
ToothGrowth.typeVC.dose1.0 = subset(ToothGrowth, supp == "VC" dose==1.0)
ToothGrowth.typeVC.dose2.0 = subset(ToothGrowth, supp == "VC" dose==2.0)
typeEffectat0.5 < -t.test(ToothGrowth.typeOJ.dose0.5len, ToothGrowth.typeVC.dose0.5len)typeEffectat1.0
-t.test(ToothGrowth.typeOJ.dose1.0len, ToothGrowth.typeVC.dose1.0len)typeEffectat2.0 <
-t.test(ToothGrowth.typeOJ.dose2.0len, ToothGrowth.typeVC.dose2.0len)
```

For the tests below: Null Hypothesis: True difference in means is equal to 0
Alternative hypothesis: True difference in means is not equal to 0 Significance level: 5

typeEffectat0.595

typeEffectat1.095

typeEffectat2.095Weconcludethatat2.0mgdosagethereisnodifferencebetweensupptypesintoothlengthincro