

About the Pediatric Pain data

This data set consists of up to four observations on 64 elementary school children. The response is the length of time the child can tolerate keeping their arm in very cold water.

Two trials were made during a first visit, and two more trials were made during a second visit two weeks later. During each visit, the first trial used the dominant arm, and the second trial used the non-dominant arm.

There is missing data on six students because of arms in casts and similar reasons; some are missing data on one arm for both visits, some for only one visit; some are missing data from the second visit altogether.

Subjects were asked what they were thinking about during the first two trials, and their response was classified into two coping style groups: those thinking about the experiment (attenders) and those thinking about other things (distracters).

Before the fourth trial, a randomized treatment was assigned, consisting of a ten-minute counseling session, where advice was given either to attend or to distract, or no advice was given.

Scientific interest lies both in the main effects of treatment and coping strategy and in possible interactions between treatment and coping strategy. In particular, subjects for whom the advice treatment matched the coping style were expected to do better (have longer times) than those with mismatched advice treatment or no advice treatment.

- How could you test for overall differences in the mean response between coping style (ignoring treatment) using methods you already know about? You may have to take a subset of the data or use summary statistics instead of the raw data. How might you handle the missing data?

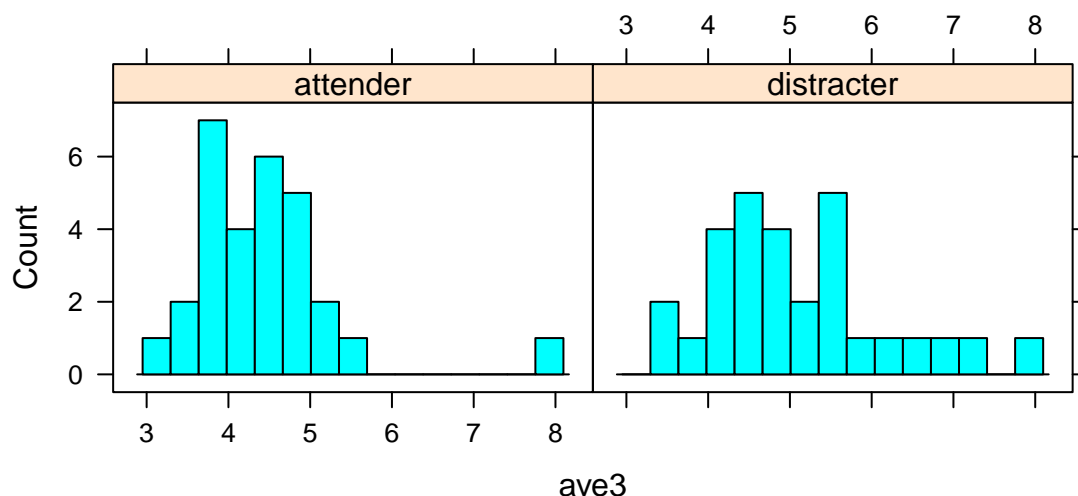
Testing means, by coping style

Read in data and remove any with missing data

```
> pp <- read.delim("http://rem.ph.ucla.edu/rob/mld/data/tabdelimiteddata/pain.txt")
> pp <- subset(pp, !is.na(l2paintol))
> ppl <- cast(id ~ ., value = "l2paintol", length, data = pp)
> names(ppl)[2] <- "n"
> id.ok <- ppl$id[ppl$n == 4]
> pp <- subset(pp, id %in% id.ok)

> pp3 <- subset(pp, trial <= 3)
> ppw <- cast(id + cs + treatment ~ ., value = "l2paintol", data = pp3,
+   fun.aggregate = mean)
> names(ppw)[4] <- "ave3"

> plot(histogram(~ave3 | cs, data = ppw, nint = 15, type = "count"))
```



```
> cast(cs ~ ., data = ppw, value = "ave3", function(x) c(n = length(x),
+   mean = mean(x), sd = sd(x)))
```

	cs	n	mean	sd
1	attender	29	4.472964	0.879483
2	distracter	29	5.076170	1.093887

```
> t.test(ave3 ~ cs, data = ppw)
```

```
Welch Two Sample t-test
```

```
data: ave3 by cs
```

```
t = -2.3143, df = 53.531, p-value = 0.02452
```

```
alternative hypothesis: true difference in means is not equal to 0
```

```
95 percent confidence interval:
```

```
-1.12586495 -0.08054762
```

```
sample estimates:
```

```
mean in group attender mean in group distracter
```

```
4.472964
```

```
5.076170
```

Testing slopes, by coping style

How do we get a linear model fit and the slope?

```
> coef(lm(l2paintol ~ trial, data = pp))
```

```
(Intercept)      trial
```

```
4.85179622 -0.03150331
```

We split the data by id and do that to each part, merging the result with the original data to keep the coping style information.

```
> ppid <- split(pp, pp$id)
```

```
> regs <- sapply(ppid, function(x) coef(lm(l2paintol ~ trial, data = x)))
```

```
> regs <- as.data.frame(t(regs))
```

```
> names(regs) <- c("intercept", "slope")
```

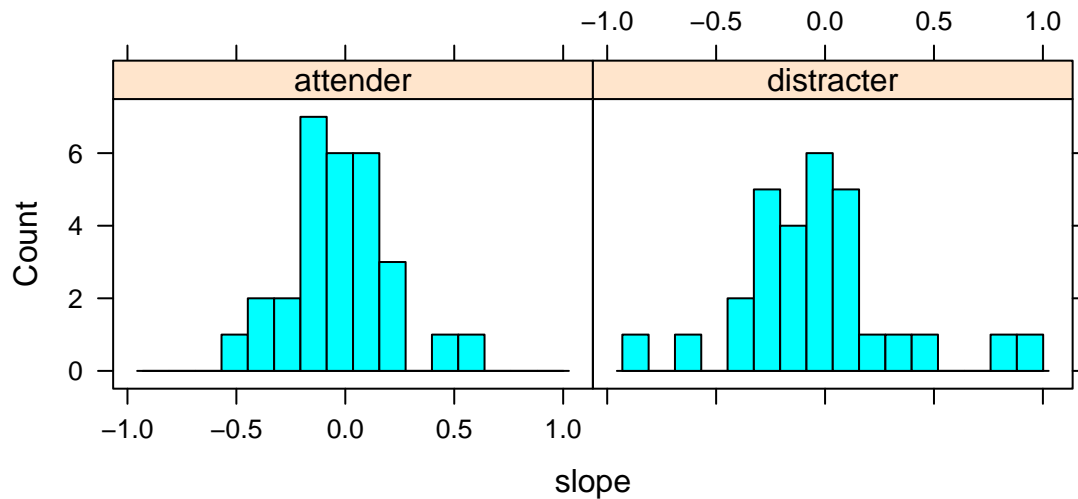
```
> regs$id <- as.numeric(rownames(regs))
```

```
> regs <- merge(regs, ppw[, 1:3])
```

```
> head(regs)
```

	id	intercept	slope	cs	treatment
1	1	5.157234	-0.37392629	attender	attend
2	2	4.941397	-0.20226956	distracter	distract
3	3	3.721983	-0.13184438	attender	no directions
4	5	4.427293	0.14114682	attender	attend
5	6	7.028407	-0.29019996	distracter	attend
6	7	4.255284	0.04324618	distracter	no directions

```
> plot(histogram(~slope | cs, data = regs, nint = 16, type = "count"))
```



```
> cast(cs ~ ., data = regs, function(x) c(n = length(x), mean = mean(x),
+   sd = sd(x)), value = "slope", fill = NA)
```

	cs	n	mean	sd
1	attender	29	-0.01737223	0.2399295
2	distracter	29	-0.04563439	0.3721908

```
> t.test(slope ~ cs, data = regs)
```

Welch Two Sample t-test

data: slope by cs

t = 0.3437, df = 47.845, p-value = 0.7326

alternative hypothesis: true difference in means is not equal to 0

95 percent confidence interval:

-0.1370865 0.1936109

sample estimates:

mean in group attender	mean in group distracter
-0.01737223	-0.04563439

A paired t-test

```
> ppw <- cast(id + cs + treatment ~ trial, value = "l2paintol",  
+ data = pp)  
> head(ppw)
```

	id	cs	treatment	1	2	3	4
1	1	attender	attend	4.361066	5.142005	3.836934	3.549669
2	2	distracter	distract	4.814038	4.598127	3.987321	4.343408
3	3	attender no directions		3.584963	3.321928	3.614710	3.047887
4	5	attender	attend	4.638653	4.549053	4.961160	4.971773
5	6	distracter	attend	6.773073	6.424754	6.099716	5.914086
6	7	distracter no directions		4.306700	4.244126	4.555816	4.346957

Two equivalent ways:

```
> t.test(ppw$`4`, ppw$`2`, paired = TRUE)
```

Paired t-test

data: ppw\$`4` and ppw\$`2`

t = 0.547, df = 57, p-value = 0.5865

alternative hypothesis: true difference in means is not equal to 0

95 percent confidence interval:

-0.1809929 0.3170310

sample estimates:

mean of the differences

0.06801907

```
> ppw$diff <- ppw$`4` - ppw$`2`
```

```
> t.test(ppw$diff)
```

One Sample t-test

data: ppw\$diff

t = 0.547, df = 57, p-value = 0.5865

alternative hypothesis: true mean is not equal to 0

95 percent confidence interval:

-0.1809929 0.3170310

sample estimates:

mean of x

0.06801907

Difference of Differences

```
> ppa <- subset(ppw, cs == "attender" & treatment != "distract")
> t.test(diff ~ treatment, data = ppa, var.equal = TRUE)
```

Two Sample t-test

```
data: diff by treatment
t = 0.4516, df = 17, p-value = 0.6573
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -0.5639146  0.8710289
sample estimates:
 mean in group attend mean in group no directions
           0.18194159                0.02838446
```

```
> ppd <- subset(ppw, cs == "distracter" & treatment != "attend")
> t.test(diff ~ treatment, data = ppd, var.equal = TRUE)
```

Two Sample t-test

```
data: diff by treatment
t = 2.6448, df = 17, p-value = 0.01702
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.2819996 2.5060545
sample estimates:
 mean in group distract mean in group no directions
           0.7153376                -0.6786894
```