

Outline

- 1 Corrective actions and Nonparametric methods
 - Data transformations
 - Mann-Whitney test

Assessing assumptions

The t-test assuming equal variances is
very **sensitive** to **dependence**,
moderately **robust** against **unequal variance** if $n_1 \approx n_2$, but
much less robust if n_1 and n_2 are quite different (e.g. differ
by a ratio of 3 or more).
robust against **nonnormality**.

Corrective actions for 2 independent samples:

Fundamental changes if problem with independence (...)

Welch t-test if σ_1 and σ_2 differ by 3-fold or more or n_1 and
 n_2 differ by 3-fold or more.

If non-normal distributions:

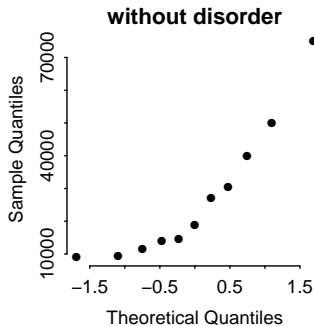
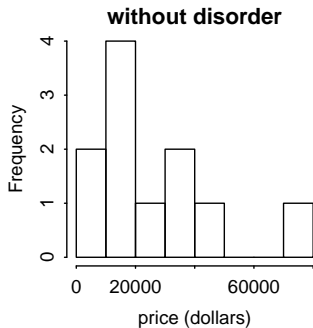
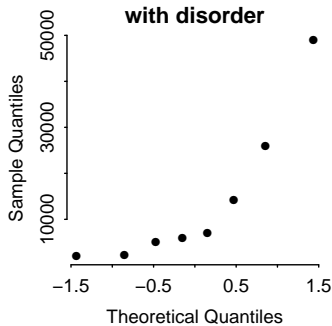
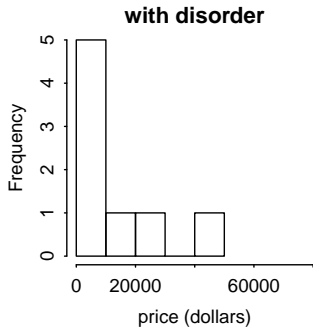
- 1 try a data transformation,
- 2 or switch to a non-parametric test: Mann-Whitney test.

Data transformations

A veterinarian wishes to know if the presence of a certain fetlock disorder in race horses affects their selling price at auction. Data on 8 horses that have the disorder, and 11 that do not (in \$)

With Disorder: 5000, 6000, 14100, 49000, 7000, 26000, 2000, 2200

Without Disorder: 27000, 14000, 11500, 19000, 9500, 40000, 75000, 9000, 14500, 50000, 30500



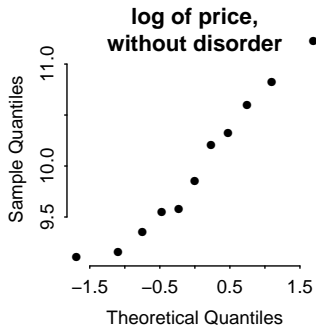
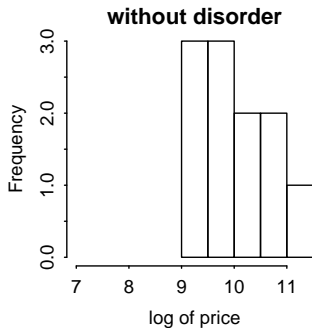
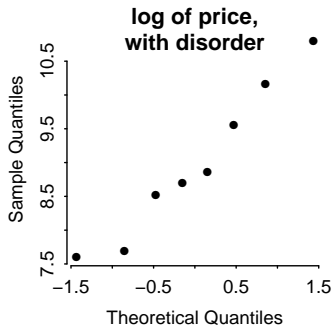
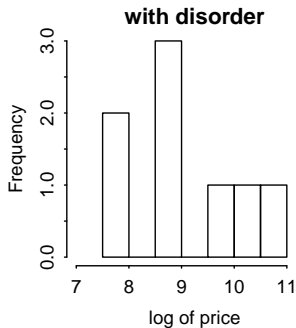
Both samples are skewed right.
Look at the log-values of the prices:

```
> dis
[1] 5000 6000 14100 49000 7000 26000 2000 2200
> log(dis)
[1] 8.52 8.70 9.55 10.80 8.85 10.17 7.60 7.70

> nod
[1] 27000 14000 11500 19000 9500 40000 75000 9000 14500 50000 30500
> log(nod)
[1] 10.20 9.55 9.35 9.85 9.16 10.60 11.23 9.10 9.58 10.82 10.33
```

Could we do the **t-test on log-values** instead?

If the price tends to go down with the fetlock disorder, then the $\log(\text{price})$ also tends to be lower with the disorder than without (and vice versa).



T-test on the log-transformed prices

The distribution of log-prices looks beautifully normal for both samples! Welch t-test on the log-transformed prices:

```
dis = c(5000, 6000, 14100, 49000, 7000, 26000, 2000, 2200)
nod =c(27000, 14000, 11500, 19000, 9500, 40000, 75000, 9000,
      14500, 50000, 30500)

> t.test(log(dis), log(nod))

      Welch Two Sample t-test

data:  log(dis) and log(nod)
t = -2.1955, df = 10.951, p-value = 0.05059
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -1.988749799  0.003048454
sample estimates:
mean of x mean of y
 8.985856  9.978706
```

Conclusion: moderate evidence that the auction prices tend to be lower with the fetlock disorder than without ($p=0.051$).

Transformations

Log transformation:

- helps when the distributions are skewed right,
- only when all values are positive

Square-root transformation:

- helps when distributions are moderately skewed right,
- only when all values are ≥ 0 (zeros are okay)

Apply the **same** transformation (here: take the log) to **all** values in **both** samples.

Choose the transformation in order to **satisfy assumptions**, **not** based on the resulting p-value.

Confidence intervals on the original scale (not log, not transformed) are more difficult to get.

What if...

The data are too skewed and no transformation can help?

For instance: a transformation might help make one sample look normally distributed but make the other sample look worse.

Third option: use a 'non parametric' test, here test that does not assume the normal distribution: the Mann-Whitney test.

Mann-Whitney test (aka Wilcoxon rank sum test)

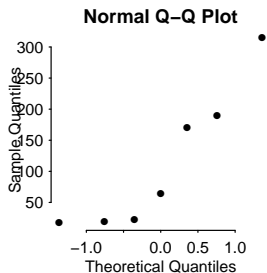
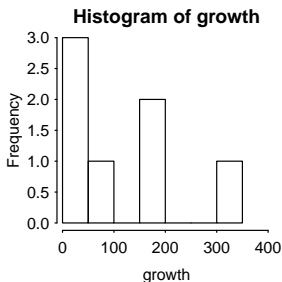
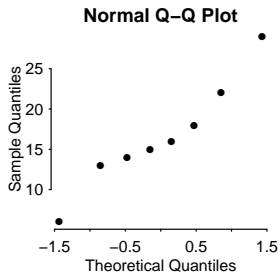
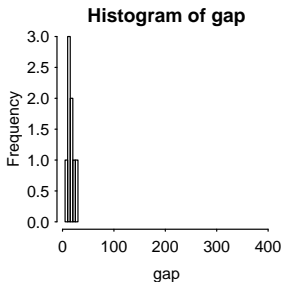
Analogous to the Wilcoxon signed-rank test (for paired samples) but here for **two independent samples**.

No distribution assumption, but still assume independence.

Main idea: look at the **ranks** of the observations

Example: Does soil respiration affect plant growth? Soil cores taken from 2 locations in a forest: under an opening in the forest canopy (“gap”) and at a nearby area under heavy tree growth (“growth”). Measured: amount of carbon dioxide given off by each soil core (mol CO₂/g soil/hr). Data:

Gap	22	29	13	16	15	18	14	6
Growth	17	20	170	315	22	190	64	



Gap data: distribution has normal shape,

Growth data: skewed right.

Welch t-test not recommended, but there is another way!

```
pdf("lec15-01.pdf",width=5,height=5)
gap    =c(22,29,13,16,15,18,14,6)
growth=c(17,20,170,315,22,190,64)
layout(matrix(1:4,2,2))
par(mar=c(3.1,3.1,1.5,.5), mgp=c(1.8,.4,0), tck=-0.01, las=1,bty="n")
hist(gap,      xlim=c(5,400))
hist(growth,  xlim=c(5,400), breaks=10)
qqnorm(gap    ,pch=16)
qqnorm(growth,pch=16)
dev.off()
```

Mann-Whitney test

- 1 H_0 : the 2 populations have the same distribution.
Soil respiration has the same distribution in the 2 locations, with $\mu_1 = \mu_2$ in particular.

H_A : soil respiration does not have the same distribution in the 2 populations. Test most sensitive to a shift between the 2 distributions, so it's usually assumed that H_A is: 'the 2 distribution have different means'.

Mann-Whitney test

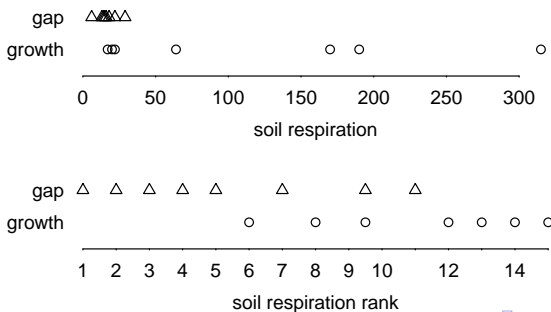
- 2 Rank the observations, calculate:

$U_1 = \#$ of observations in group 2 that are smaller

$U_2 = \#$ of observations in group 1 that are smaller

and summarize the data by $U = \max\{U_1, U_2\}$.

If H_0 is true, then U has a Wilcoxon distribution (does not depend on the common distribution of the data).



Mann-Whitney test

- ③ We got $U = 49.5$, more extreme = larger, so $p\text{-value} = \mathbb{P}\{U \geq 49.5\}$.

Table E, $n_1 = 8$ and $n_2 = 7$: critical (minimum) U is 46 for rejecting at $\alpha = 0.05$, and 50 at $\alpha = 0.01$

So here $.01 < p\text{-value} < .05$

- ④ Conclusion: we have moderate evidence that the soil respiration distribution differs between the two locations.

Soil respiration has a higher mean in the aread under heavy tree growth, than in the area under the opening of the forest canopy.

Note: Table E has no number listed for $n_1 = 3$ and $n_2 = 4$: we can never reject H_0 at $\alpha = 0.05$.

One-sided Mann-Whitney test

H_A : distribution shift with $\mu_1 > \mu_2$ for instance.

First check that the data go in the same direction as H_A , i.e. check that $U_1 > U_2$ if testing $H_A: \mu_1 > \mu_2$.

If not: p-value > 0.50 .

If so: p-value is half as much as what it would be for a two-sided test.

wilcox.test() in R

```
> gap
[1] 22 29 13 16 15 18 14  6
> growth
[1]  17  20 170 315  22 190  64

> wilcox.test(gap, growth)
```

Wilcoxon rank sum test with continuity correction

data: gap and growth

W = 6.5, p-value = 0.015

alternative hypothesis: true location shift is not equal
to 0

Warning message:

In wilcox.test.default(gap, growth) :
cannot compute exact p-value with ties

Warnings and Assumptions

If there are **ties**, the table gives approximation only.

The test does not work well if the **variances** are very different

To interpret H_A as simply $\mu_1 \neq \mu_2$ rather than 'the 2 distributions are different', we actually need to assume that when the 2 distributions differ, they only differ by their means –not variances.

The Mann-Whitney test is **less powerful** than the t-test (when both are applicable) for small sample sizes, but almost as powerful for last sample sizes.

Try transformation + t-test first: more powerful if applicable
Otherwise use Mann-Whitney.