

Comparing means (or medians) between 2 groups

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Flow-chart for popularly used statistical tests

Q1, Univariate / Multivariable	Q2, Difference / Correlation	Q3, Paired / related	Q4, Q5 Type of outcome (Normality)	Q6, No. of groups	Q7, sample size	Valid Tests
Univariate	Difference	Independent (un-paired)	Continuous (Normal)	2		Student's t-test
			>2			One-way ANOVA
			Continuous (Non-normal) / Ordered categorical	2		Mann-Whitney U test
			>2			Kruskal-Wallis H test
			Nominal	2	<20	Fisher's exact test
				≥2	≥20	Chi-square test
			Time to Event			Log-Rank test (Kaplan-Meier plot)
	Correlation	Dependent (paired)	Continuous (Normal)	2		Paired-t test
			>2			Repeated measured ANOVA
			Continuous (Non-normal) / Ordered categorical	2		Mixed effect Regression
			>2			Wilcoxon signed-rank test
			Nominal	2		Friedman test
				2		McNemar's test
Multivariable		Independent (un-paired)	Continuous (Normal)			Pearson's correlation (r)
			Continuous (Non-normal) / ordered			Spearman's correlation (rs)
			Nominal (2 levels)	2		Spearman/Kappa (Agreement)
			Continuous (Normal residuals)			Linear Regression
			Continuous (Non-normal residuals)			Linear Regression*
			Ordered categorical			Ordered Logistic Regression
		Dependent (paired)	Nominal (>2 levels)			Binary Logistic Regression
			>2			Multinomial Logistic Regression
			Time to Event			Cox Proportional Hazard Regression
			Continuous (Normal residuals)			Linear Mixed Effect Regression
			Continuous (Non-normal residuals)			Linear Mixed Effect Regression*
			Ordered categorical			Generalized Estimation Equation (GEE)
			Nominal (>2 levels)			Generalized Estimation Equation (GEE)

* Transform outcome variables for normalizing residuals

Created based on Publishing Your Medical Research Paper, by Daniel Byrne, Williams and Wilkins (1998)

Motivating Example: 2-arm RCT of Type 2 diabetes patients

The American Journal of Medicine (2005) 118, 276–284



THE AMERICAN
JOURNAL of
MEDICINE®

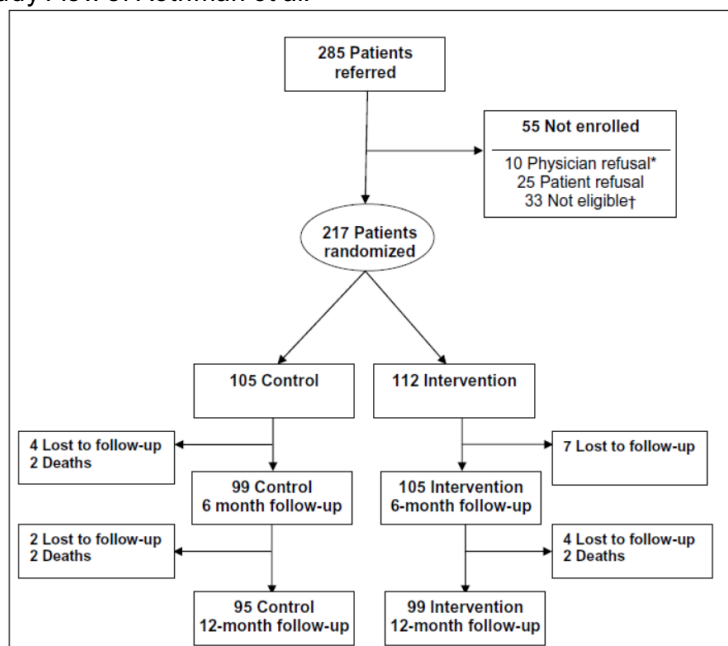
CLINICAL RESEARCH STUDY

A randomized trial of a primary care–based disease management program to improve cardiovascular risk factors and glycated hemoglobin levels in patients with diabetes

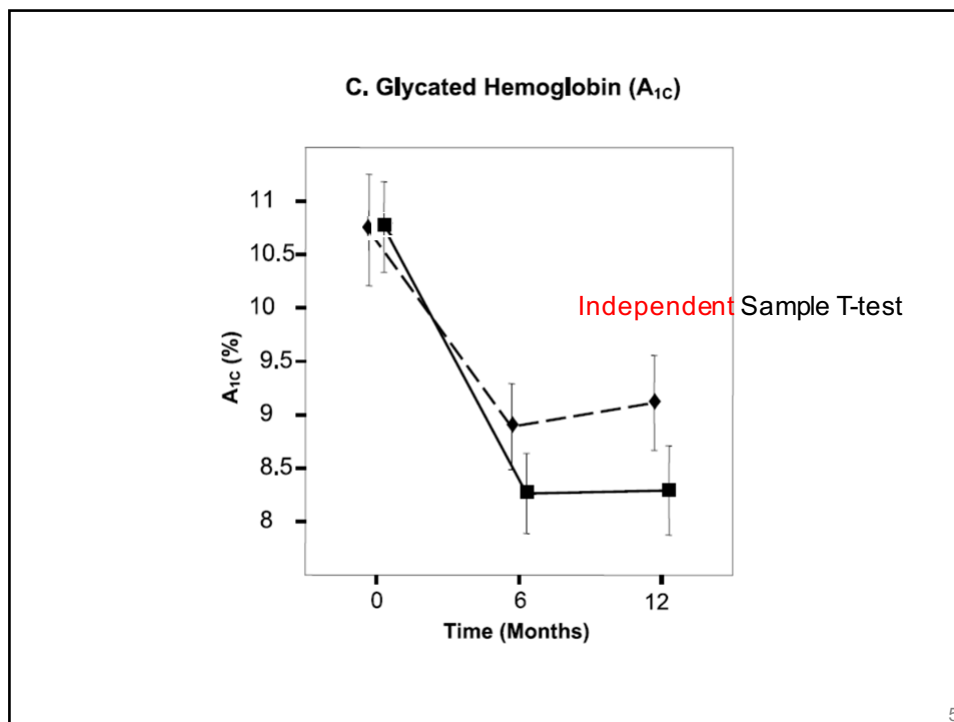
Russell L. Rothman, MD, MPP,^{a,b} Robb Malone, PharmD, CDE,^c
Betsy Bryant, PharmD, CDE,^c Ayumi K. Shintani, PhD, MPH,^{a,b} Britton Crigler,^c
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Michael P. Pignone, MD, MPH^c

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Study Flow of *Rothman et al.*



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Flow-chart for popularly used statistical tests						
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			>2			Wilcoxon signed-rank test
			Nominal	2		Friedman test
				2		McNemar's test
Multivariable	Difference	Independent (un-paired)	Continuous (Normal)			Pearson's correlation (r)
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			Continuous (Normal residuals)			Linear Mixed Effect Regression
			Continuous (Non-normal residuals)			Linear Mixed Effect Regression*
			Ordered categorical			Generalized Estimation Equation (GEE)
			Nominal (2 levels)			Generalized Estimation Equation (GEE)

* Transform outcome variables for normalizing residuals

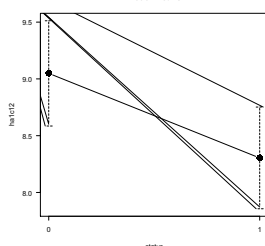
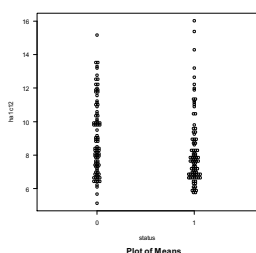
Created based on Publishing Your Medical Research Paper, by Daniel Byrne, Williams and Wilkins (1998)

Independent Sample t-test (Student's t-test)

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Example: In the RCT of pharmacy based intervention of Type 2 diabetes (DiabetesRCT180.xlsx), compare the difference in the improvement in HbA1c at 12 month) **between the intervention and control group.**

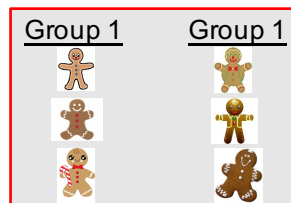
H_0 : Mean of Ha1c in the intervention group (M1) = Mean of ha1c in the control group (M0).
 H_1 : Mean of Ha1c in the intervention group (M1) \neq Mean of ha1c in the control group (M0).



$$\text{Test statistics} = \frac{M1 - M0}{SE(M1 - M0)}$$

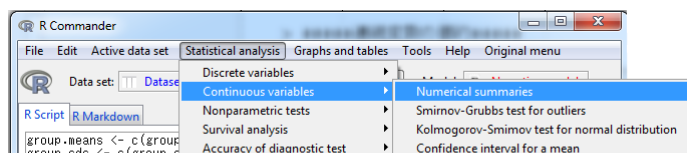
Larger the test statistics, smaller p-value

Difference in means is large \Rightarrow smaller p-value
 Data variation is smaller \Rightarrow smaller p-value



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Descriptive data:



```
> #####Numerical Summaries#####
      Mean    SD   0% 25% 50% 75% 100% data:n
0 9.048315 2.192711 5.1 7.3 8.4 10.5 15.2    89
1 8.305495 2.148765 5.8 6.9 7.6  8.9 16.0    91
```

$$test\ statistics = \frac{M1 - M0}{SE(M1 - M0)} = \frac{M1 - M0}{\sqrt{\left(\frac{s_1^2}{n_1} + \frac{s_0^2}{n_0}\right)}}$$

Where

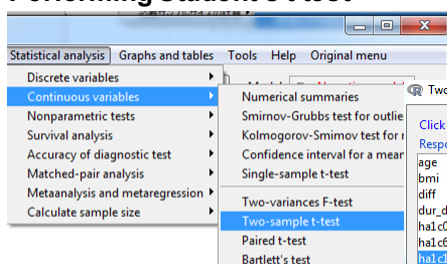
$$s = \sqrt{\left\{\frac{(n_1 - 1)s_1^2 + (n_0 - 1)s_0^2}{n_1 + n_0 - 2}\right\}}$$

S₁=SD for the intervention group, S₀=SD for the control group

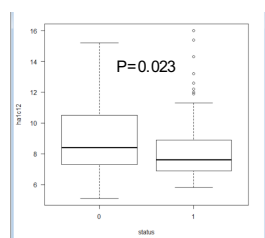
n₁=sample size of the intervention group, n₀=sample size of the control group

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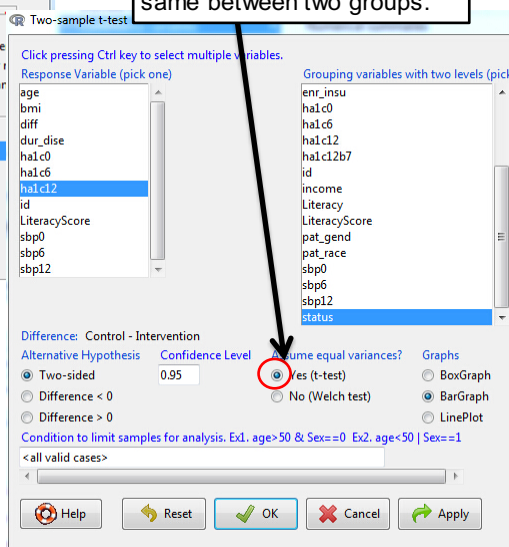
Performing Student's t-test



```
> summary.ttest
      Mean    SD   P-value
status=0 9.048315 2.192711 0.0229
status=1 8.305495 2.148765
```



Select this if variance (SD) of the outcome variable is the same between two groups.



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Performing Student's t-test

```
Two Sample t-test

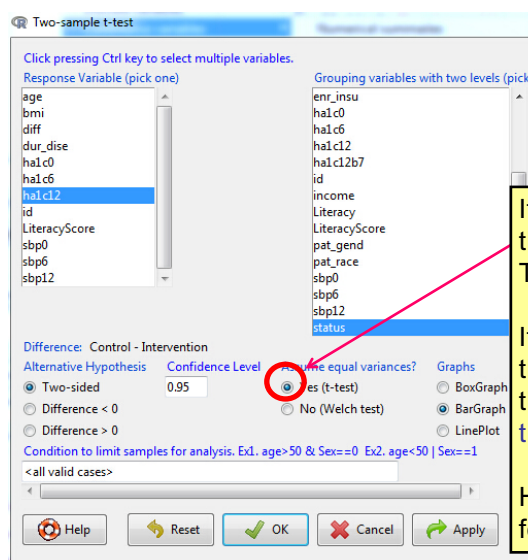
data: ha1c12 by factor(status)
t = 2.2955, df = 178, p-value = 0.02287
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.1042454  1.3813948
sample estimates:
mean in group 0 mean in group 1
 9.048315      8.305495
```

This confidence interval is for the difference in means of group 1 and 0.
Unfortunately R output does not compute this for you.
 $9.048 - 8.305 = 0.743$.

Thus mean difference in Ha1c between the groups is
 0.743 95%CI=(0.104, 1.38) $p=0.023$

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Performing Student's t-test : **Checking equal variances**



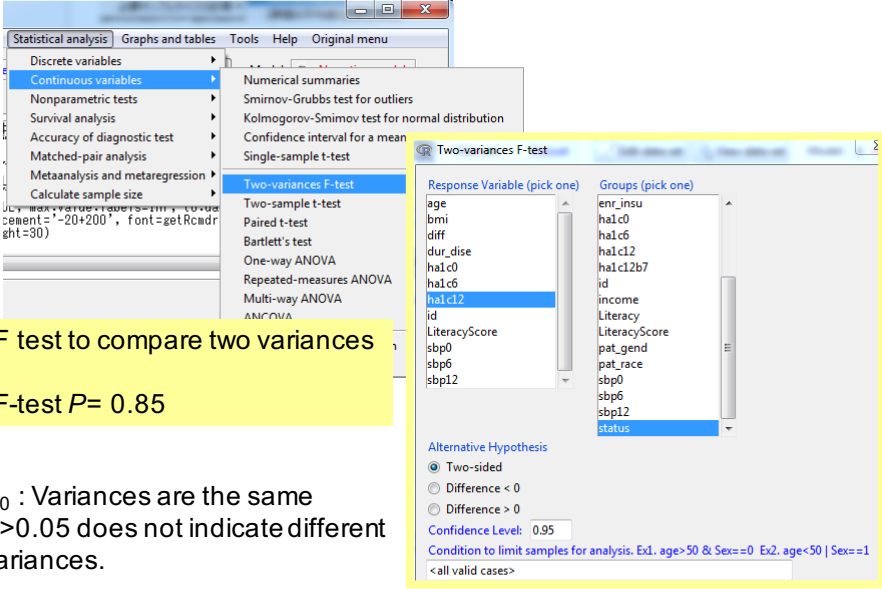
If variance is the same between the groups, use formula on Slide9, Thus click on **Yes**.

If variance is not the same between the groups, use a different formula, thus click on **No** (the formula not in these slides).

How to judge **Yes** or **No** in the following slide.

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Performing Student's t-test : **Checking equal variances**



The image shows the SPSS Statistical analysis menu with 'Continuous variables' selected, leading to 'Two-variances F-test'. The dialog box for 'Two-variances F-test' is open, showing 'Response Variable (pick one)' as 'hal_c12' and 'Groups (pick one)' as 'status'. The 'Alternative Hypothesis' is set to 'Two-sided'. The 'Confidence Level' is 0.95. The 'Condition to limit samples for analysis' is set to '<all valid cases>'. Below the dialog box, a yellow box contains the text: 'F test to compare two variances', 'F-test P= 0.85', and 'H₀ : Variances are the same P>0.05 does not indicate different variances.'

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Assumption of Student's (independent sample) t-test.

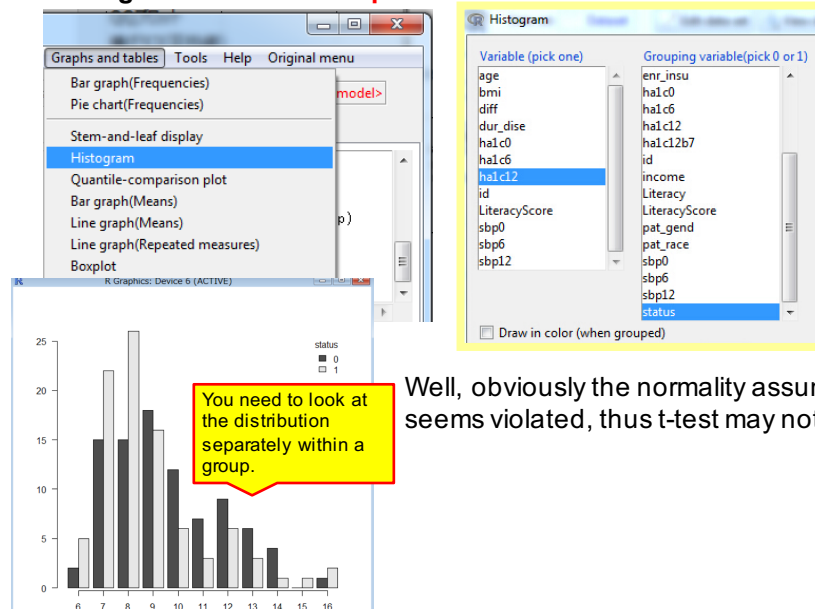
1. Distribution of outcome variable is **normal** within a group.
2. All observations are **independent**.

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**How to deal with a violation of normality assumption:
Log-transforming outcome variables**

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Checking the normal assumption



Well, obviously the normality assumption seems violated, thus t-test may not be valid.

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How to deal with a violation of normality assumption (1): Log-transforming the outcome variable

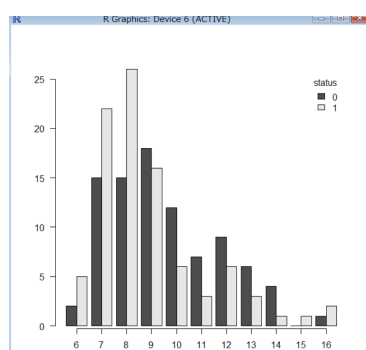
The screenshot shows the R Commander interface. The 'Variables' menu is open, and 'Create new variable' is selected. A dialog box titled 'Create new variable' is displayed. In the 'Current variables (double-click to expression)' list, 'ha1c12' is selected. The 'New variable name' field contains 'loge.ha1c12'. The 'Expression to compute' field contains 'log(ha1c12)'. The 'OK' button is highlighted.

Below the dialog box, a 'Data' window shows a table with columns 'id', 'ha1c12', and 'loge.ha1c12'. The data is as follows:

id	ha1c12	loge.ha1c12
1	131	8.1
2	140	12.6
3	159	11.0
4	173	6.7
5	98	6.9
6	205	5.8
7	213	12.0
8	61	7.0
9	16	6.6
10	99	7.6
11	198	8.0
12	56	6.5
13	134	13.2
14	190	6.6
15	58	7.4
16	84	10.1
17	129	11.9
18	171	9.6
19	189	6.9
20	34	9.3
21	79	9.1
22	161	7.0
23	72	7.5
24	104	5.8
25	53	8.8
26	127	6.7

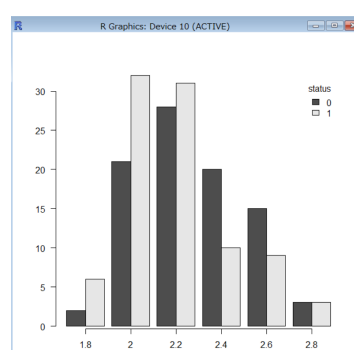
17

Checking for assumption for the model using Log (e) transformed outcome variable, Ha1c12



Ha1c12

Before Log(e) transformation



Ha1c12

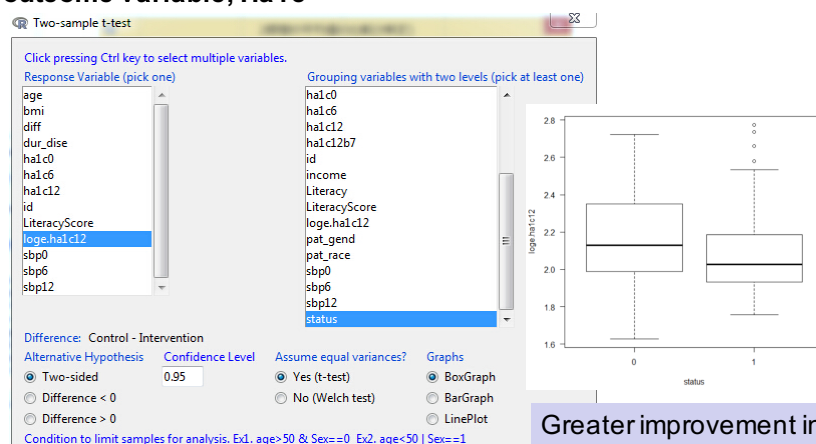
After Log(e) transformation

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**How to deal with a violation of normality assumption:
Performing student's t-test using the log-transformed outcome variable**

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Solution to the example: student's t-test for log (e) transformed outcome variable, Ha1c



```
> summary.ttest
      Mean      SD      P-value
status=0 2.174374 0.2378323 0.0151
status=1 2.088704 0.2305144
```

Greater improvement in
The normality would often
lead to gaining analytical
power, resulting in smaller p-
value.

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MED101x

Introduction to Applied Biostatistics

```
> (res <- t.test(logc.ha1c12~factor(status), alternative='two.sided', conf.level=0.95,
+ var.equal=TRUE, data=Dataset))
Two Sample t-test

data: logc.ha1c12 by factor(status)
t = 2.4541, df = 178, p-value = 0.01509
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.0167815 0.1545582
sample estimates:
mean in group 0 mean in group 1
 2.174374      2.088704
```

This confidence interval is for the difference in means of group 1 and 2.

2.174374 - 2.088704

= 0.08567

Exp(0.08567) = 1.089 = 9.05 / 8.30

= Ratio of mean Ha1c12 between groups

= mean Ha1c12 in the control group is 1.089 times of Mean Ha1c12 in the intervention group.

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Flow-chart for popularly used statistical tests

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			Continuous (Non-normal) / Ordered categorical	2		Mann-Whitney U test
			> 2			Kruskal-Wallis H test
			Nominal	2	< 20	Fisher's exact test
				≥ 2	≥ 20	Chi-square test
			Time to Event			Log-Rank test (Kaplan-Meier plot)
	Correlation	Dependent (paired)	Continuous (Normal)	2		Paired-t test
			> 2			Repeated measured ANOVA
			Continuous (Non-normal) / Ordered categorical	2		Mixed effect Regression
			Nominal	2		Wilcoxon signed-rank test
Multivariable	Difference	Independent (un-paired)	Continuous (Normal)	2		Friedman test
			> 2			McNemar's test
			Continuous (Non-normal) / Ordered categorical	2		Pearson's correlation (r)
			Nominal (2 levels)	2		Spearman's correlation (rs)
						Spearman/Kappa (Agreement)
			Continuous (Normal residuals)			Linear Regression
	Correlation	Dependent (paired)	Continuous (Non-normal residuals)			Linear Regression*
			Ordered categorical			Ordered Logistic Regression
			Nominal (2 levels)			Binary Logistic Regression
			> 2			Multinomial Logistic Regression
	Difference	Independent (un-paired)	Time to Event			Cox Proportional Hazard Regression
			Continuous (Normal residuals)			Linear Mixed Effect Regression
			Continuous (Non-normal residuals)			Linear Mixed Effect Regression*
			Ordered categorical			Generalized Estimation Equation (GEE)
			Nominal (2 levels)			Generalized Estimation Equation (GEE)
	Correlation	Dependent (paired)	Continuous (Normal residuals)			Linear Regression
			Continuous (Non-normal residuals)			Linear Regression*
			Ordered categorical			Ordered Logistic Regression
			Nominal (2 levels)			Binary Logistic Regression

* Transform outcome variables for normalizing residuals

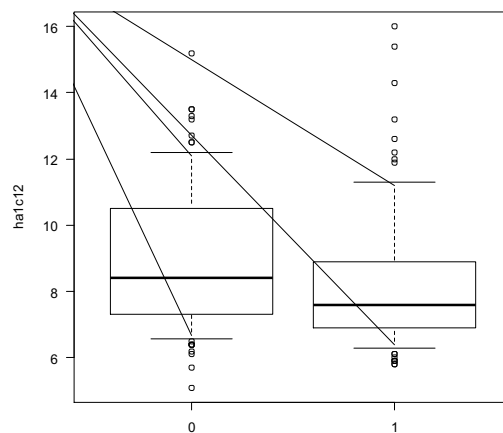
Created based on Publishing Your Medical Research Paper, by Daniel Byrne, Williams and Wilkins (1998)

How to deal with a violation of normality assumption (2):
Use non-parametric test

Mann-Whitney U test
Wilcoxon Rank-Sum test

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Comparing two independent distributions of continuous outcome:
Wilcoxon Rank-Sum test / Mann-Whitney U test



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Mann-Whitney U test / Wilcoxon rank-sum test

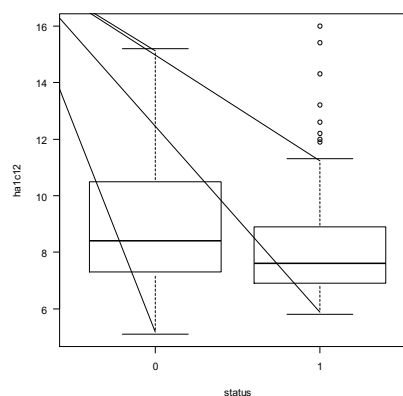
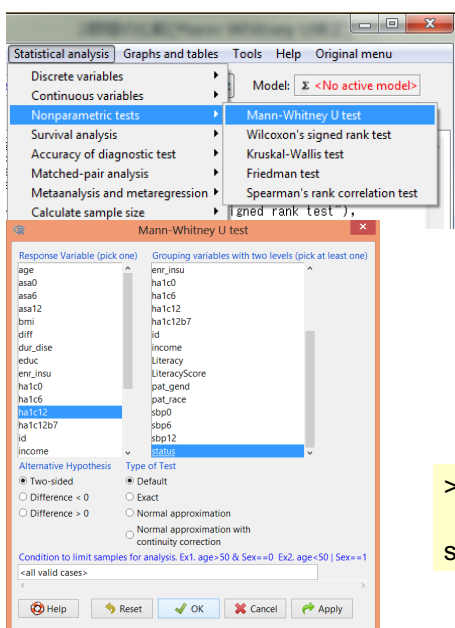
id	status	ha1c12	rank_ha1c12
131	0	8.1	93.0
159	0	11.0	149.0
16	0	8.6	24.5
198	0	8.0	88.0
56	0	6.5	21.0
134	0	13.2	172.5
84	0	10.1	139.0
129	0	11.9	161.0
79	0	9.1	121.0
181	0	7.0	47.0
72	0	7.5	67.5
53	0	8.8	113.0
127	0	6.7	30.0
174	0	6.4	17.5
...
140	1	12.6	170.0
173	1	8.7	30.0
98	1	6.9	39.5
205	1	5.8	3.5
213	1	12.0	163.5
81	1	7.0	47.0
99	1	7.8	73.0
190	1	6.6	24.5
58	1	7.4	63.5
171	1	9.6	127.5
189	1	6.9	39.5
34	1	9.3	123.0
104	1	5.9	6.0
28	1	11.0	149.0

- (1) Combine the data from the two groups, and order the values from the lowest to the highest.
- (2) Assign ranks to the individual values from the lowest rank to the highest rank.
- (3) If a group of observations has the same value, then compute the range of ranks for the group, and then assign the average rank for each observation in the group.

Then sum of these ranks is compared between groups.

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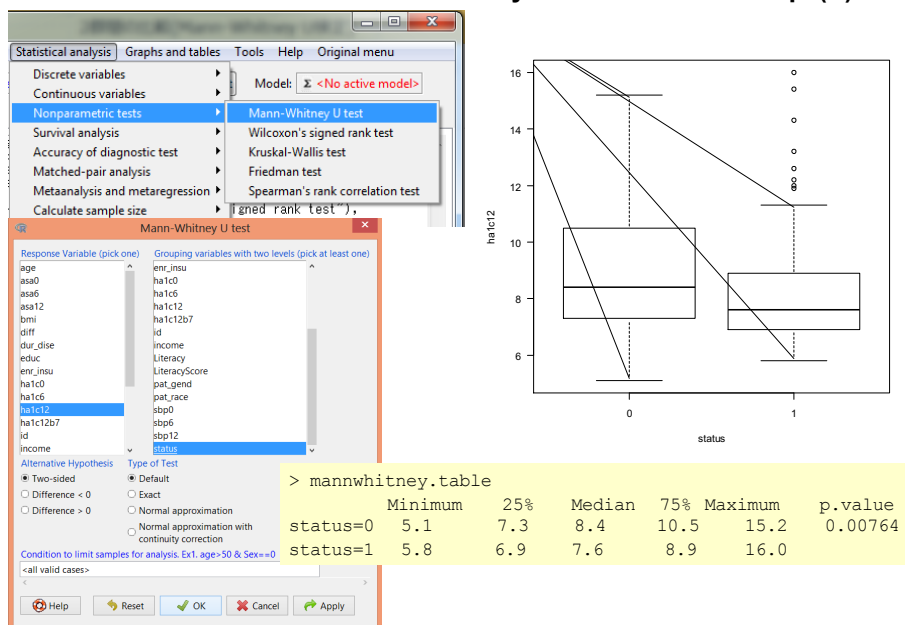
Wilcoxon Rank-Sum test / Mann-Whitney U test: Basic Concept (2)



> mannwhitney.test
p.value
status 0.00764

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Wilcoxon Rank-Sum test / Mann-Whitney U test: Basic Concept (2)



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			>2		≥20	Chi-square test
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	Correlation	Dependent (paired)	Continuous (Normal)	2		Paired-t test
			>2			Repeated measured ANOVA
			Continuous (Non-normal) / Ordered categorical	2		Mixed effect Regression
			>2			Wilcoxon signed-rank test
Multivariable	Difference	Independent (un-paired)	Ordered categorical	2		Friedman test
			Nominal	2		McNemar's test
			Continuous (Normal)			Pearson's correlation (r)
			Continuous (Non-normal) / ordered			Spearman's correlation (rs)
			Nominal (2 levels)	2		Spearman/Kappa (Agreement)
			Continuous (Normal residuals)			Linear Regression
	Correlation	Dependent (paired)	Continuous (Non-normal residuals)			Linear Regression*
			Ordered categorical			Ordered Logistic Regression
			Nominal (2 levels)			Binary Logistic Regression
			>2			Multinomial Logistic Regression
	Difference	Independent (un-paired)	Time to Event			Cox Proportional Hazard Regression
			Continuous (Normal residuals)			Linear Mixed Effect Regression
			Continuous (Non-normal residuals)			Linear Mixed Effect Regression*
			Ordered categorical			Generalized Estimation Equation (GEE)
			Nominal (2 levels)			Generalized Estimation Equation (GEE)
	Correlation	Dependent (paired)	Continuous (Normal residuals)			Linear Regression
			Continuous (Non-normal residuals)			Linear Regression*
			Ordered categorical			Ordered Logistic Regression
			Nominal (2 levels)			Binary Logistic Regression
			>2			Multinomial Logistic Regression

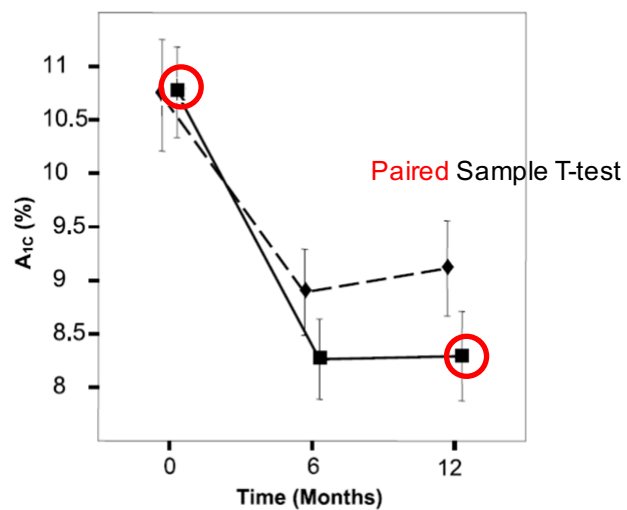
* Transform outcome variables for normalizing residuals

Created based on Publishing Your Medical Research Paper, by Daniel Byrne, Williams and Wilkins (1998)

Paired t-test

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C. Glycated Hemoglobin (A_{1c})



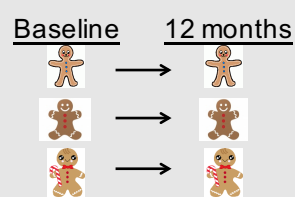
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Example (slide 6): Randomized Controlled Clinical Trial.

DiabetesRCT180.xlsx

Assess if there is a change in HbA1c level **between baseline and 12 months** among patients in the intervention group.

Paired groups: A patient was measured twice at baseline and at 12 months follow-up. Two samples are said to be paired when each data point of the first sample is matched (or related) to a unique data point of the second sample.



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Paired t-test:

$H_0: \mu_d = 0$. Where μ_d = the difference between baseline and follow-up values

This is equivalent with one-sample test for mean which we learned at the last lecture!!!

These 2 observations are related, since they came from the same person.

id	status	ha1c0	ha1c12	diff
140	1	11.0	12.6	-1.60
173	1	15.2	6.7	8.50
98	1	9.0	6.9	2.09
205	1	9.7	5.8	3.89
213	1	14.3	12.0	2.30
61	1	8.9	7.0	1.89
99	1	11.1	7.6	3.50
190	1	11.6	6.6	5.00
58	1	11.6	7.4	4.20
171	1	9.8	9.6	0.19
189	1	8.7	6.9	1.79
34	1	11.3	9.3	2.00
104	1	9.8	5.9	3.90
28	1	10.6	11.0	-0.39
118	1	12.3	7.5	4.80
123	1	12.7	9.6	3.09

Mean diff > 0
↕
Reduction in Ha1c

Mean diff = 0
↕
No change in Ha1c

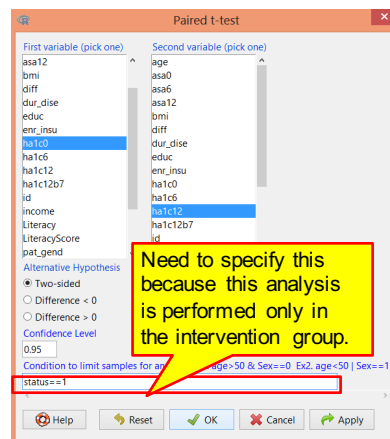
Mean diff < 0
↕
Increase in Ha1c

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Performing paired t-test

Paired t-test
data: subset(Dataset, status == 1)\$ha1c0
and subset(Dataset, status == 1)\$ha1c12
t = 8.242, df = 90, p-value = 1.29e-12
alternative hypothesis: true difference in
means is not equal to 0
95 percent confidence interval:
1.879043 3.072605
sample estimates:
mean of the differences
2.475824

```
> summary.ttest
      mean      sd p.value
ha1c0 10.781319 2.178323 1.29e-12
ha1c12  8.305495 2.148765
```



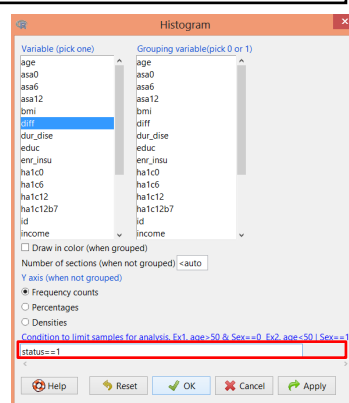
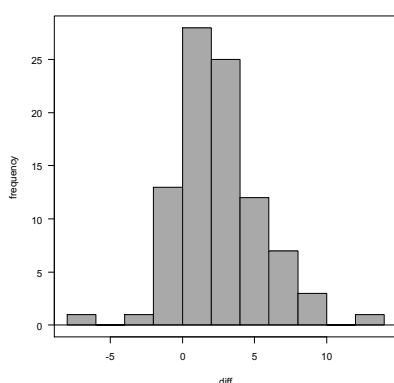
Mean reduction of Ha1c was 2.48 with 95% CI = (1.88, 3.07) $p < 0.0001$. Thus the observed reduction was statistically significant at 2-sided significance level of 5%. Note that the 95% CI for the difference in means **does not include 0**.

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Important

Assumption of paired t-tests is:

The difference between the two related variables are normally distributed.



Remember the assumption of the paired t-test is that the difference between the two related variables are normally distributed. Thus we first create the difference variable.

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Wilcoxon Signed-Rank test

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Flow-chart for popularly used statistical tests

Q1, Univariate / Multivariable	Q2, Difference / Correlation	Q3, Paired / related	Q4, Q5 Type of outcome (Normality)	Q6, No. of groups	Q7, sample size	Valid Tests
Univariate	Difference	Independent (un-paired)	Continuous (Normal)	2		Student's t-test
				>2		One-way ANOVA
			Continuous (Non-normal) / Ordered categorical	2		Mann-Whitney U test
				>2		Kruskal-Wallis H test
			Nominal	2	<20	Fisher's exact test
				≥2	≥20	Chi-square test
			Time to Event			Log-Rank test (Kaplan-Meier plot)
		Dependent (paired)	Continuous (Normal)	2		Paired-t test
				>2		Repeated measured ANOVA
			Continuous (Non-normal) / Ordered categorical	2		Mixed effect Regression
				>2		Wilcoxon signed-rank test
Multivariable	Correlation	Independent (un-paired)	Continuous (Normal)	2		Friedman test
				>2		McNemar's test
			Continuous (Normal)			Pearson's correlation (r)
			Continuous (Non-normal) / ordered			Spearman's correlation (rs)
		Dependent (paired)	Nominal (2 levels)	2		Spearman/Kappa (Agreement)
			Continuous (Normal residuals)			Linear Regression
			Continuous (Non-normal residuals)			Linear Regression*
			Ordered categorical			Ordered Logistic Regression
			Nominal (>2 levels)			Binary Logistic Regression
			Nominal (>2)			Multinomial Logistic Regression
			Time to Event			Cox Proportional Hazard Regression
		Dependent (paired)	Continuous (Normal residuals)			Linear Mixed Effect Regression
			Continuous (Non-normal residuals)			Linear Mixed Effect Regression*
			Ordered categorical			Generalized Estimation Equation (GEE)
			Nominal (2 levels)			Generalized Estimation Equation (GEE)

* Transform outcome variables for normalizing residuals

Created based on Publishing Your Medical Research Paper, by Daniel Byrne, Williams and Wilkins (1998)

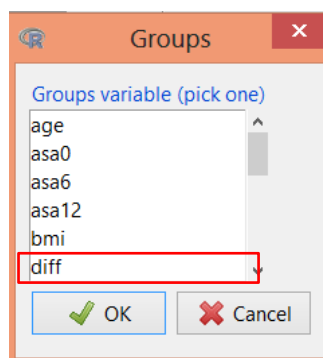
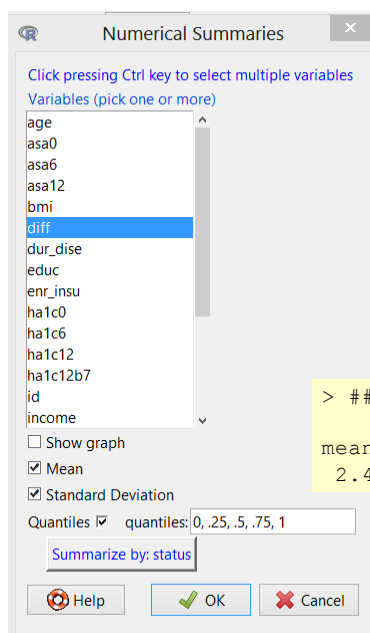
Wilcoxon Signed Rank test (Non parametric test for comparing paired data).

id	status	ha1c0	ha1c12	diff	Sign	Ranks on Absolute Difference
140	1	11.0	12.6	-1.60	-	3
173	1	15.2	6.7	8.50	+	16
98	1	9.0	6.9	2.09	+	7
205	1	9.7	5.8	3.89	+	11
213	1	14.3	12.0	2.30	+	8
61	1	8.9	7.0	1.89	+	5
99	1	11.1	7.6	3.50	+	10
190	1	11.6	6.6	5.00	+	15
58	1	11.6	7.4	4.20	+	13
171	1	9.8	9.6	0.19	+	1
189	1	8.7	6.9	1.79	+	4
34	1	11.3	9.3	2.00	+	6
104	1	9.8	5.9	3.90	+	12
28	1	10.6	11.0	-0.39	-	2
118	1	12.3	7.5	4.80	+	14
123	1	12.7	9.6	3.09	+	9

- (1) Arrange the difference in order of absolute values.
- (2) Count the number of differences with the same absolute value.
- (3) Ignore the observations where differences = 0, and rank the remaining observations from 1 for the observation with the lowest absolute value, up to n for the observation with the highest absolute value.
- (4) If there is a group of several observations with the same absolute values (**ties**), then find the lowest rank in the range = 1 + R and the highest rank in the range = G + R, where R = the highest rank used prior to considering this group and G = the number of differences in the range of ranks for the group. Assign the **average rank** = (lowest rank in the range + highest rank in the range) / 2 as the rank for each difference in the group.

Rank-sum for "+" = 16+7+11+8+5+10+15+...+9
Rank-sum for "-" = 3+2

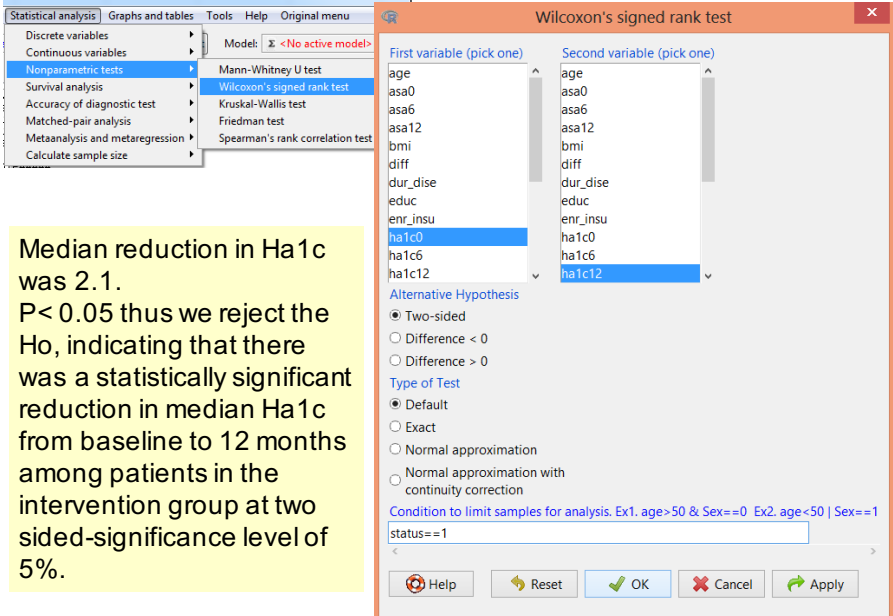
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```
> #####Numerical summaries#####
mean      sd      0% 25% 50% 75% 100% n
2.475824 2.865556 -6.3 0.7 2.1 3.95 12.8 91
```

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Performing a Wilcoxon Signed Rank test



Median reduction in Ha1c was 2.1.
 $P < 0.05$ thus we reject the H_0 , indicating that there was a statistically significant reduction in median Ha1c from baseline to 12 months among patients in the intervention group at two sided-significance level of 5%.