

Meta Analysis of Rectal Cancer Studies

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Biometrician: Svenja Seide
seide@imbi.uni-heidelberg.de

Contacts: Laura Seifert
lauraseifert96@gmail.com

Karl Kowalewski
karl.kowalewski@googlemail.com

Data source: rectal cancer statistic_neu.xlsx

Data supplied by: Karl Kowalewski

Abstract: [ToDo]

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New names:

- * 'Abdominal pain post op lap' -> 'Abdominal pain post op lap...34'
- * 'Abdominal pain post op lap (SD)' -> 'Abdominal pain post op lap (SD)...35'
- * 'Abdominal pain post op rob' -> 'Abdominal pain post op rob...36'
- * 'Abdominal pain post op rob (SD)' -> 'Abdominal pain post op rob (SD)...37'
- * 'Abdominal pain post op lap' -> 'Abdominal pain post op lap...80'
- * ... and 3 more problems

New names:

- * 'LAP directly postop (wert)' -> 'LAP directly postop (wert)...18'
- * 'LAP directly postop (wert)' -> 'LAP directly postop (wert)...19'
- * 'Postop 3 months rob (wert)' -> 'Postop 3 months rob (wert)...20'
- * 'Postop 3months rob (Abweichung)' -> 'Postop 3months rob (Abweichung)...21'
- * 'Postop 3 months rob (wert)' -> 'Postop 3 months rob (wert)...24'
- * ... and 1 more problem

1 Urinary Retention

Robotic treatment (experimental group) compared to laproscopic treatment (control group). Results on the OR-scale (non-logarithmic). In case of 0 events in a study-arm, 0.5 is added to each arm of this study for continuity correction. Assuming that "urinary retention" is an undesirable event, $OR < 1$ favors Rob, while $OR > 1$ favors Lab. Studies are sorted by total sample size (from small to large) in the forest plot.

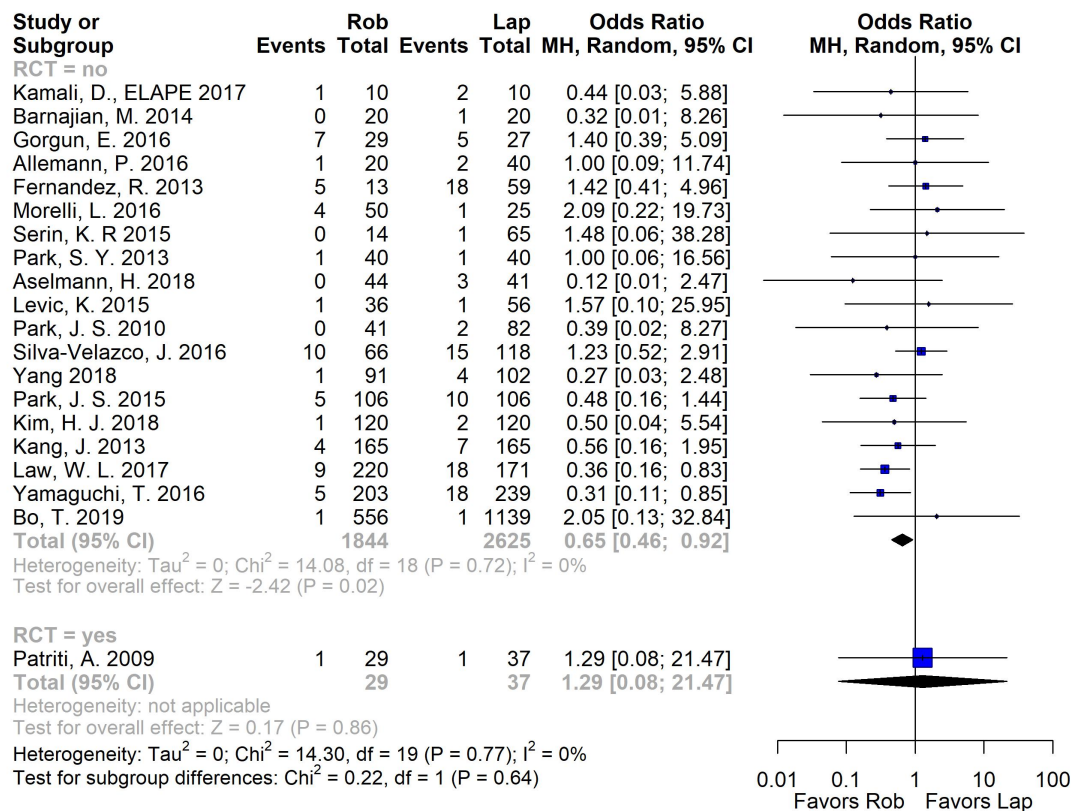


Figure 1: Forest plot for meta-analysis of urinary retention

Number of studies combined: $k = 20$

	OR	95%-CI	z	p-value
Random effects model	0.6561	[0.4636; 0.9286]	-2.38	0.0174

Quantifying heterogeneity:

$\tau^2 = 0$; $H = 1.00$ [1.00; 1.20]; $I^2 = 0.0\%$ [0.0%; 30.8%]

Quantifying residual heterogeneity:

$H = 1.00$ [1.00; 1.24]; $I^2 = 0.0\%$ [0.0%; 34.7%]

Test of heterogeneity:

Q	d.f.	p-value
14.30	19	0.7659

Results for subgroups (random effects model):

	k	OR	95%-CI	Q	τ^2	I^2
RCT = no	19	0.6494	[0.4576; 0.9215]	14.08	0	0.0%
RCT = yes	1	1.2857	[0.0770; 21.4725]	0.00	--	--

Test for subgroup differences (random effects model):

	Q	d.f.	p-value
Between groups	0.22	1	0.6370

Details on meta-analytical method:

- Mantel-Haenszel method
- DerSimonian-Laird estimator for τ^2
- Continuity correction of 0.5 in studies with zero cell frequencies

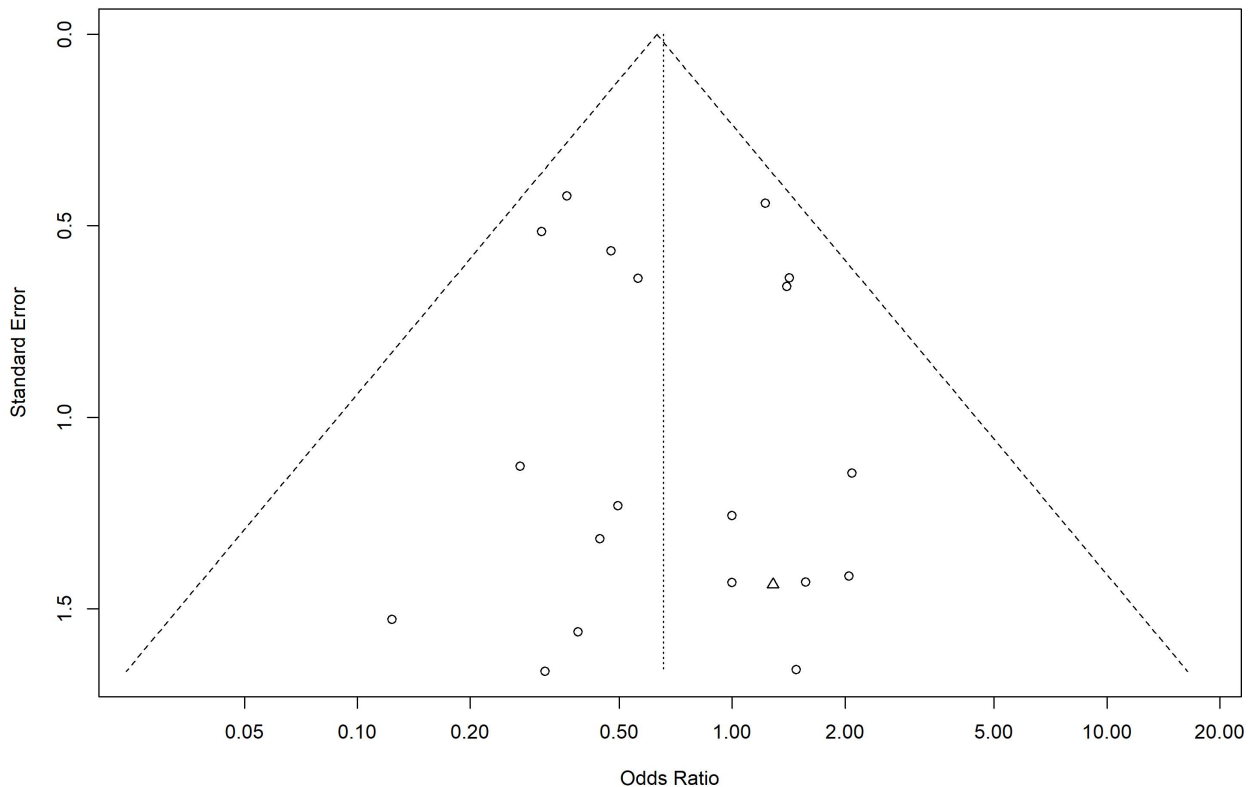


Figure 2: Funnel plot for meta-analysis of urinary retention

2 Ileus

Robotic treatment (experimental group) compared to laproscopic treatment (control group). Results on the OR-scale (non-logarithmic). In case of 0 events in a study-arm, 0.5 is added to each arm of this study for continuity correction. Assuming that "ileus" is an undesirable event, OR < 1 favors Rob, while OR > 1 favors Lab. Studies are sorted by total sample size (from small to large) in the forest plot.

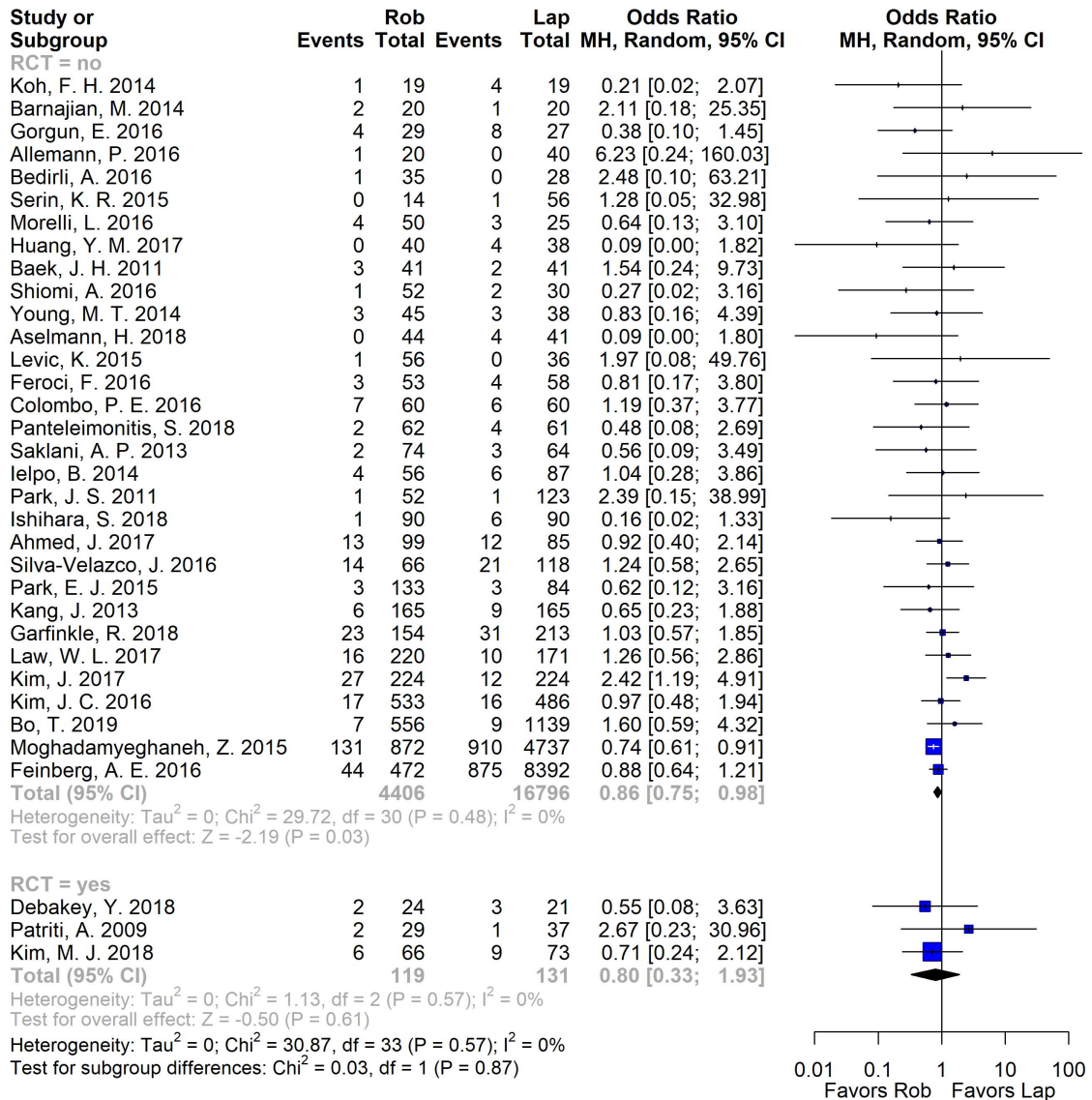


Figure 3: Forest plot for meta-analysis of Ileus

Number of studies combined: k = 34

	OR	95%-CI	z	p-value
Random effects model	0.8578	[0.7503; 0.9807]	-2.24	0.0248

Quantifying heterogeneity:

$\tau^2 = 0$; $H = 1.00$ [1.00; 1.24]; $I^2 = 0.0\%$ [0.0%; 34.5%]

Quantifying residual heterogeneity:

$H = 1.00$ [1.00; 1.26]; $I^2 = 0.0\%$ [0.0%; 36.9%]

Test of heterogeneity:

Q	d.f.	p-value
30.87	33	0.5735

Results for subgroups (random effects model):

	k	OR	95%-CI	Q	τ^2	I^2
RCT = no	31	0.8593	[0.7504; 0.9840]	29.72	0	0.0%
RCT = yes	3	0.7967	[0.3297; 1.9253]	1.13	0	0.0%

Test for subgroup differences (random effects model):

	Q	d.f.	p-value
Between groups	0.03	1	0.8681

Details on meta-analytical method:

- Mantel-Haenszel method
- DerSimonian-Laird estimator for τ^2
- Continuity correction of 0.5 in studies with zero cell frequencies

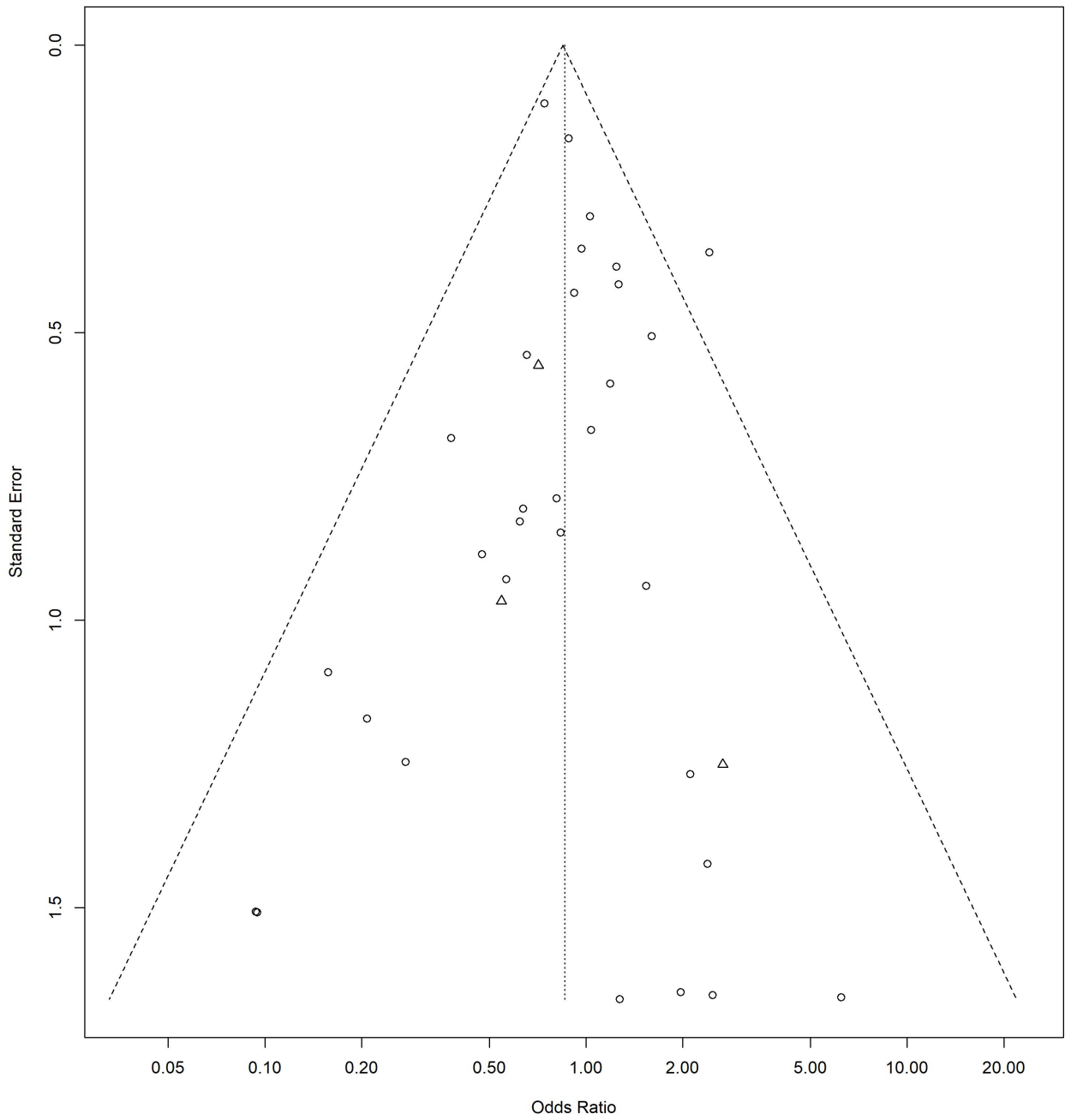


Figure 4: Funnel plot for meta-analysis of Ileus

3 QLQ-Scores

Robotic treatment (experimental group) compared to laproscopic treatment (control group). Results as difference in means. Designations in the forest plot assume a higher mean differences to be better in all cases which translates into a higher score being assumed better.

3.1 QLQ-C-29

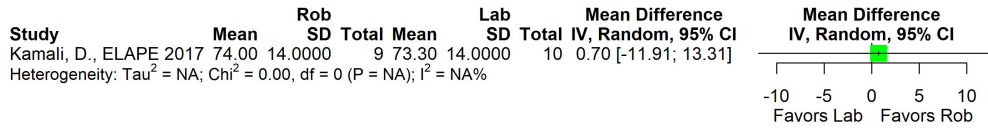


Figure 5: Forest plot for meta-analysis of QLQ-C-29 global health

MD 95%-CI z p-value
 0.7000 [-11.9076; 13.3076] 0.11 0.9133

Details:

- Inverse variance method

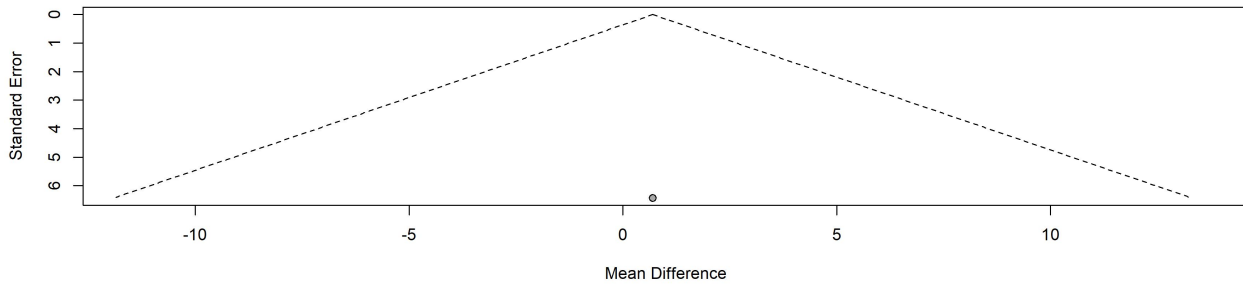


Figure 6: Funnel plot for meta-analysis of QLQ-C-29 global health

3.2 QLQ-C-30

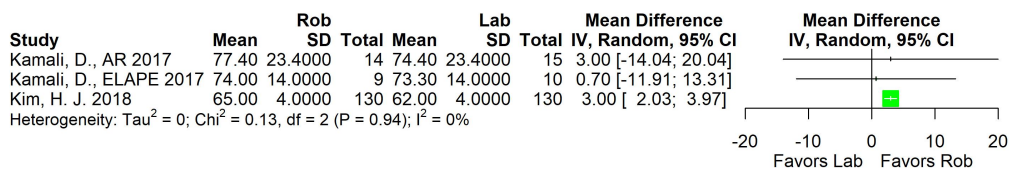


Figure 7: Forest plot for meta-analysis of QLQ-C-30 global health

Number of studies combined: k = 3

MD 95%-CI z p-value
Random effects model 2.9864 [2.0185; 3.9544] 6.05 < 0.0001

Quantifying heterogeneity:
 $\tau^2 = 0$; $H = 1.00$ [1.00; 1.00]; $I^2 = 0.0\%$ [0.0%; 0.0%]

Test of heterogeneity:
Q d.f. p-value
0.13 2 0.9384

Details on meta-analytical method:
- Inverse variance method
- DerSimonian-Laird estimator for τ^2

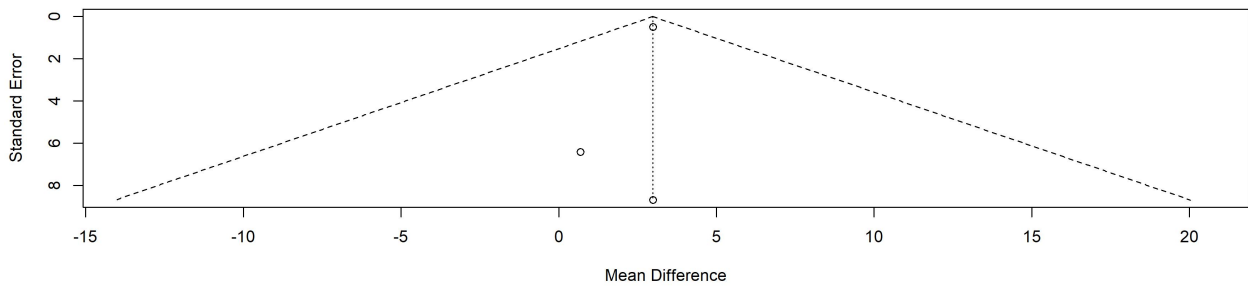


Figure 8: Funnel plot for meta-analysis of QLQ-C-30 global health

4 IIEF Total Scores

Robotic treatment (experimental group) compared to laparoscopic treatment (control group). Results as mean differences and assuming a higher score being better. Assuming that the data is asymptotically normal (especially symmetric) in the studies reporting median and range, mean and standard deviation are estimated.

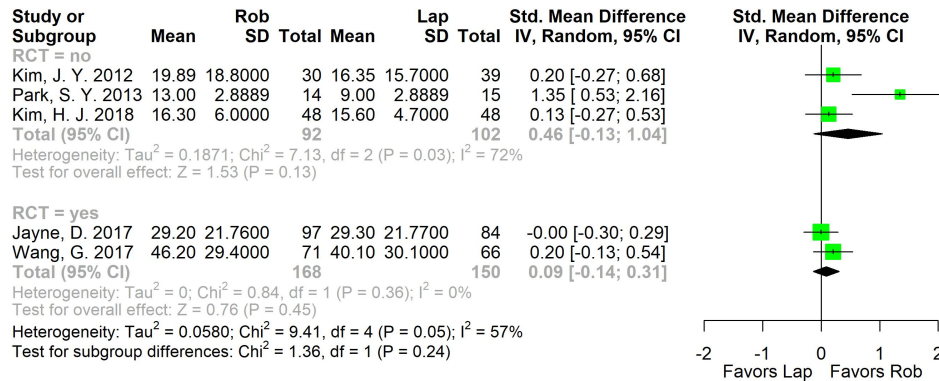


Figure 9: Forest plot for meta-analysis of IIEF at latest time point available, sorted by time of measurement from earliest to latest

Number of studies combined: $k = 5$

	SMD	95%-CI	z	p-value
Random effects model	0.2375	[-0.0491; 0.5242]	1.62	0.1043

Quantifying heterogeneity:

$\tau^2 = 0.0580$; $H = 1.53$ [1.00; 2.52]; $I^2 = 57.5\%$ [0.0%; 84.2%]

Quantifying residual heterogeneity:

$H = 1.63$ [1.00; 2.81]; $I^2 = 62.4\%$ [0.0%; 87.4%]

Test of heterogeneity:

Q	d.f.	p-value
9.41	4	0.0517

Results for subgroups (random effects model):

	k	SMD	95%-CI	Q	τ^2	I^2
RCT = no	3	0.4571	[-0.1287; 1.0429]	7.13	0.1871	72.0%
RCT = yes	2	0.0852	[-0.1353; 0.3057]	0.84	0	0.0%

Test for subgroup differences (random effects model):

	Q	d.f.	p-value
Between groups	1.36	1	0.2442

Details on meta-analytical method:

- Inverse variance method
- DerSimonian-Laird estimator for τ^2
- Hedges' g (bias corrected standardised mean difference)

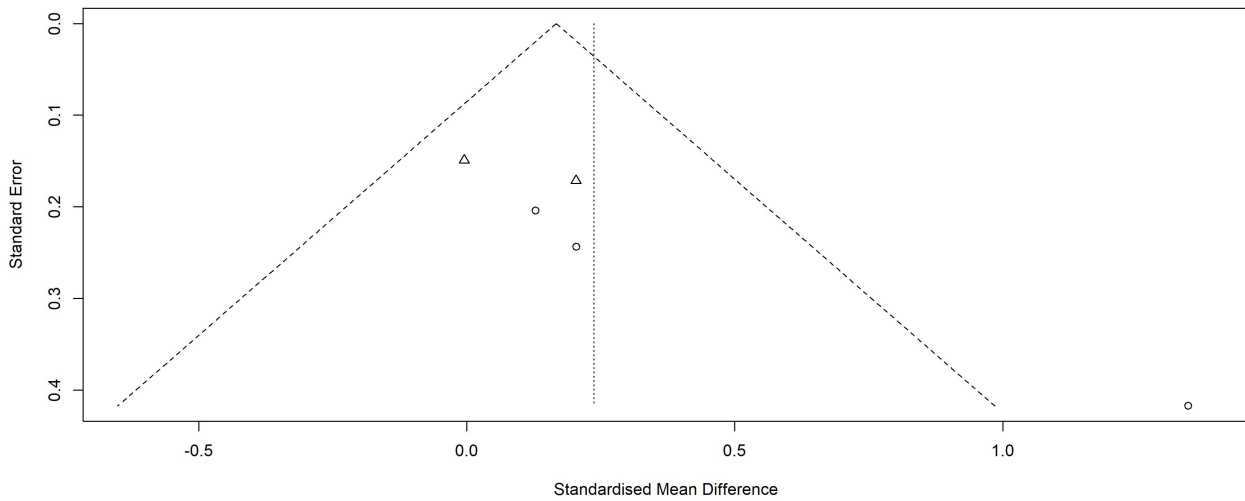


Figure 10: Funnel plot for meta-analysis of IIEF at latest time point available, sorted by time of measurement from earliest to latest

5 IPSS Total Scores

Robotic treatment (experimental group) compared to laproscopic treatment (control group). Results as mean differences and assuming a lower score to be better. Assuming that the data is asymptotically normal (especially symmetric) in the studies reporting median and range, mean and standard deviation are estimated.

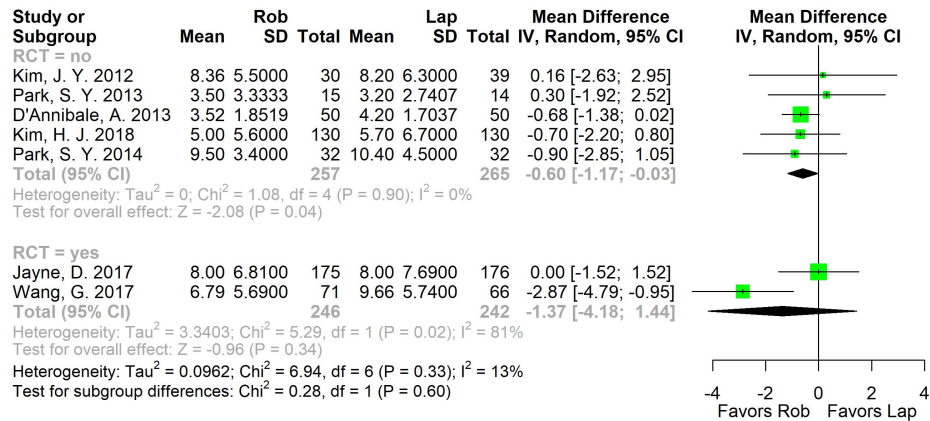


Figure 11: Forest plot for meta-analysis of IPSS at latest time point available, sorted by time of measurement from earliest to latest

Number of studies combined: k = 7

	MD	95%-CI	z	p-value
Random effects model	-0.7019	[-1.3081; -0.0957]	-2.27	0.0232

Quantifying heterogeneity:

$\tau^2 = 0.0962$; $H = 1.08$ [1.00; 1.99]; $I^2 = 13.5\%$ [0.0%; 74.7%]

Quantifying residual heterogeneity:

$H = 1.13$ [1.00; 1.71]; $I^2 = 21.5\%$ [0.0%; 65.9%]

Test of heterogeneity:

Q	d.f.	p-value
6.94	6	0.3268

Results for subgroups (random effects model):

	k	MD	95%-CI	Q	τ^2	I^2
RCT = no	5	-0.6020	[-1.1706; -0.0335]	1.08	0	0.0%
RCT = yes	2	-1.3733	[-4.1832; 1.4367]	5.29	3.3403	81.1%

Test for subgroup differences (random effects model):

	Q	d.f.	p-value
Between groups	0.28	1	0.5980

Details on meta-analytical method:

- Inverse variance method
- DerSimonian-Laird estimator for τ^2

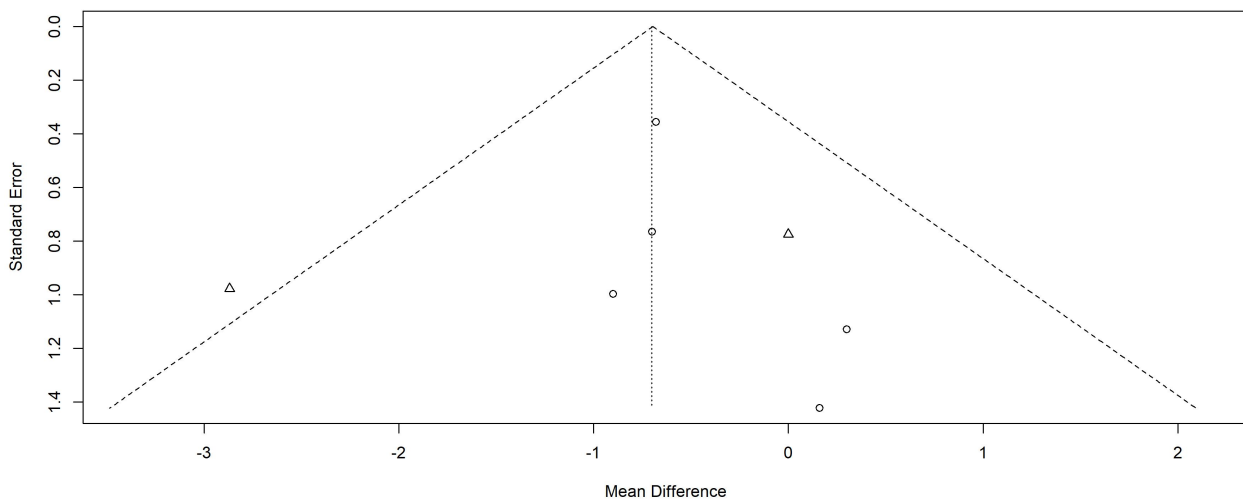


Figure 12: Funnel plot for meta-analysis of IPSS at latest time point available, sorted by time of measurement from earliest to latest

6 Session Information

R version 3.5.1 (2018-07-02)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows 10 x64 (build 17134)

Matrix products: default

locale:

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[3] LC_MONETARY=German_Germany.1252 LC_NUMERIC=C
[5] LC_TIME=German_Germany.1252

attached base packages:

[1] stats graphics grDevices utils datasets methods base

other attached packages:

[1] readxl_1.1.0 meta_4.9-2 forcats_0.3.0 stringr_1.3.1
[5] dplyr_0.8.0.1 purrr_0.2.5 readr_1.1.1 tidyr_0.8.2
[9] tibble_2.1.1 ggplot2_3.1.0 tidyverse_1.2.1

loaded via a namespace (and not attached):

[1] Rcpp_1.0.0 cellranger_1.1.0 pillar_1.3.1 compiler_3.5.1
[5] plyr_1.8.4 tools_3.5.1 digest_0.6.18 lubridate_1.7.4
[9] jsonlite_1.5 evaluate_0.12 nlme_3.1-137 gtable_0.2.0
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[21] xml2_1.2.0 httr_1.3.1 knitr_1.20 hms_0.4.2
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