Additional file 1: Meta-regression and Additional figures.

Meta-regression

1. Sepsis mortality

Meta-regression	Number of obs = 14							
REML estimate of between-study variance				ta	= .0113			
% residual variation due to heterogeneity				I -squared_res = 91.99%				
Proportion of between-study variance explained				Adj R-squared = -61.65%				
Joint test for all covariates				Model $F(5,8) = 0.13$				
With Knapp-Hartung modification				Prob > F = 0.9825				
_ES	Coef.	Std. Err.	t	P> t	[95% Conf. I	nterval]		
+								
Definition	.019325	.0350875	0.55	0.597	0615869	.1002369		
Study design	.0098267	.0719412	0.14	0.895	15607	.1757235		
Study location	0290154	.0558661	-0.52	0.618	1578428	.099812		

0.38

-0.27

1.30

0.713

0.798

0.230

-.156752

-.1803497

-.2208817

.2188418

.1431608

.7898267

.0814382

.0701453

.2191471

.0310449

.2844725

2. Septic shock mortality

_cons |

Study centers | -.0185944

Endpoints |

Meta-regression	Number of obs $=$ 10
REML estimate of between-study variance	tau2 = .01358
% residual variation due to heterogeneity	$I-squared_res = 87.42\%$
Proportion of between-study variance explained	Adj R-squared = 24.45%
Joint test for all covariates	Model $F(4,5) = 1.53$
With Knapp-Hartung modification	Prob > F = 0.3220

_ES	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
Definition	.0245329	.1276626	0.19	0.855	3036343	.3527
Study design	2249434	.1081076	-2.08	0.092	5028429	.052956
Study location	0291704	.1532626	-0.19	0.857	4231445	.3648036
Study centers	.0133103	.1025027	0.13	0.902	2501813	.2768019
_cons	.7553199	.3030357	2.49	0.055	0236582	1.534298

Figure S1



Domains:

- D1: Bias due to confounding.
 D2: Bias due to selection of participants.
 D3: Bias in classification of interventions.
 D4: Bias due to deviations from intended interventions.
- D5: Bias due to missing data. D6: Bias in measurement of outcomes.
- D7: Bias in selection of the reported result.

Judgement

Critical

Serious

Moderate

Low

No information

Figure S2

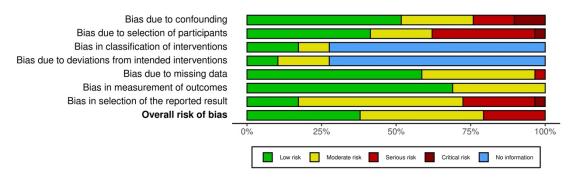


Figure S3

