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**e-Appendix: Amount of data required for meta-analysis of fully reported outcomes**

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Type of outcome data	Required for meta-analysis
Unpaired continuous data	<ul style="list-style-type: none"> <li>• Sample size in each group;</li> <li>• Magnitude of treatment effect (group means/medians or difference in means/medians); and</li> <li>• Measure of precision or variability (confidence interval, standard deviation or standard error for means; interquartile or other range for medians) or the precise p value*</li> </ul>
Unpaired binary data	<ul style="list-style-type: none"> <li>• Sample size in each group; and</li> </ul> <p>Either</p> <ul style="list-style-type: none"> <li>• Number (or percent) of participants with event for each group; or</li> <li>• Odds ratio or relative risk with measure of precision or variability (confidence interval, standard deviation or standard error) or the precise p value*</li> </ul>
Paired continuous data	<ul style="list-style-type: none"> <li>• Sample size in each group; and</li> </ul> <p>Either</p> <ul style="list-style-type: none"> <li>• Mean difference between groups and a measure of its precision, variability or precise p value; or</li> <li>• Raw data for each participant</li> </ul>
Paired binary data	<ul style="list-style-type: none"> <li>• Sample size in each group; and</li> <li>• Paired numbers of participants with and without events</li> </ul>
Survival data	<p>Either</p> <ul style="list-style-type: none"> <li>• Kaplan-Meier curve or similar, with numbers of patients at risk over time; or</li> <li>• Hazard ratio with a measure of precision, and sample size in each group</li> </ul>

\*Sample sizes, treatment effect and precise p value enable the calculation of a standard error if a measure of precision or variability is not reported.