

**Additional file 5 –Tables summarising each method by taxonomy classification and type of event suitable for**

**Table 3: Summary of visual approaches to summarise AE data in phase II/III RCTs**

Outcome	Data type	Plot	Reference	Brief Description
<b>Emerging adverse events (multiple)</b>	Binary	Volcano	Zink, Wolfinger & Mann 2013. Xia 2011 first proposed this method for systematic reviews but was not eligible for inclusion in this review. <sup>1 2</sup>	Summarises and compares the incidence of each AE reported by treatment group
	Binary	Dot	Amit, Heiberger & Lane, 2008. Cooper, 2008 also proposed but for pooled trials therefore not eligible for inclusion in this review. <sup>3 4</sup>	Provides an absolute and relative measure compared across treatment group for each AE reported
	Time-to-event	Tendril	Karpefors & Weatherall, 2018. <sup>5</sup>	Provides a summary of time-to-event data by treatment group for each AE reported
	Binary	Heat map	Zink, Marchenko, Sanchez-Kam, Ma & Jiang <sup>6</sup>	Visualises treatment effects for each AE reported
	Binary	Bar chart	Chuang-Stein & Xia, 2013. <sup>7</sup>	Displays frequency of events
	Binary	Venn diagram	Chuang-Stein & Xia, 2013. <sup>7</sup>	Presents frequencies highlighting the prevalence of overlapping events
	Binary	Two-by-two frequencies	Chuang-Stein, Le & Chen, 2001. <sup>8</sup>	Displays two-by-two frequencies graphically
<b>Emerging adverse events (single)</b>	Time-to-event	Kaplan-Meier	Amit, Heiberger & Lane, 2008. <sup>3</sup>	Summarises time-to-event data highlighting absolute differences over time
	Time-to-event	Hazard function	Amit, Heiberger & Lane, 2008. <sup>3</sup>	Summarises time-to-event data highlighting the time at which differences emerge
	Time-to-event	Risk over time	Chuang-Stein & Xia, 2013. <sup>7</sup>	Summarises incidence of an event over time
<b>(Emerging)</b>	Continuous	Cumulative frequency plots/empirical cumulative distribution function	Amit, Heiberger & Lane, 2008. <sup>3</sup>	Provides a summary of the distribution e.g. change for individual participants over time

<b>Laboratory &amp; Vital Signs (single or multiple)</b>	Boxplots	Amit, Heiberger & Lane, 2008. Cooper, 2008 also proposed but for pooled trials therefore not eligible for inclusion in this review. <sup>3,4</sup>	Provides a summary of the distribution e.g. change at specific time points
	Line graphs	Amit, Heiberger & Lane, 2008. <sup>3</sup>	Provides a summary of change at specific time points
	Histograms	Chuang-Stein, Le & Chen, 2001. <sup>8</sup>	Provides a summary of the distribution e.g. change at specific time points
	Scatter plots	Amit, Heiberger & Lane, 2008. Cooper, 2008 also proposed but for pooled trials therefore not eligible for inclusion in this review. <sup>3,4</sup>	Provides a summary of change for individual participants over time
	Scatter plot with regression	Southworth, 2008. <sup>10</sup>	Provides a summary of change for individual participants over time highlighting any outliers
	Delta	Chuang-Stein, Le & Chen, 2001. <sup>8</sup>	Display individual participant changes
	Vector plots	Trost & Freston, 2008. <sup>11</sup>	Simultaneously displays individual participant changes across three laboratory values
	e-Dish	Chuang-Stein & Xia, 2013. <sup>7</sup>	Scatter plot of peak serum ALT and peak bilirubin levels for individual participants to identify drug induced serious hepatotoxicity

**Table 4: Summary of hypothesis tests to analyse prespecified harm outcomes in phase II/III RCTs**

<b>Outcome</b>	<b>Data type</b>	<b>Model</b>	<b>Reference</b>	<b>Brief Description</b>
<b>Prespecified safety outcome</b>	Binary	Logit	Bolland & Whitehead <sup>12</sup>	Alpha-spending function for sequential monitoring
	Binary, count, time-to-event or continuous	Not applicable	Fleishman & Parker <sup>13</sup>	Redefine significance threshold for sequential monitoring
	Binary, count, time-to-event or continuous	Not applicable		Conditional power at each interim analysis for sequential monitoring
	Time-to-event	Exponential		Alpha-spending function for sequential monitoring
	Binary or incidence rate	Binomial or Poisson	Lieu et al. <sup>14</sup>	Maximised sequential probability ratio test for sequential monitoring
	Binary, count - incident rate	Poisson	Shih, Lai, Heyse & Chen <sup>15</sup>	Sequential generalised likelihood ratio test for sequential monitoring
	Binary, count, time-to-event or continuous	Not applicable	Liu <sup>16</sup>	Non-inferiority test for final analysis

**Table 5: Summary of hypothesis tests to analyse emerging AE data in phase II/III RCTs**

Outcome	Data type	Model	Reference	Brief Description
<b>Emerging Adverse events (multiple)</b>	Binary	Not applicable	Mehrotra & Heyse <sup>17</sup>	P-value adjustment
	Binary	Not applicable	Mehrotra & Adewale <sup>18</sup>	P-value adjustment
<b>Emerging Adverse events (single)</b>	Time-to-event	Poisson	Huang, Zalkikar & Tiwari <sup>19</sup>	Likelihood ratio test to compare relative risk for time-to-first event
	Time-to-event	Poisson		Likelihood ratio test to compare relative risk allowing recurrent events
<b>Overall adverse event profile*</b>	Binary	Multivariate - Markov chain	Bristol & Patel <sup>20</sup>	Multivariate likelihood ratio test with Markov chain of order one
	Binary	Multivariate - chi-squared	Chuang-Stein Mohberg & Musselman <sup>21</sup>	Multivariate test with chi-squared distribution
	Binary	Multinomial - logit	Agresti & Klingenberg <sup>22</sup>	Likelihood ratio test using a logit models to test for equality of two vectors for the marginal distributions
	Binary	Exact permutation distribution		Likelihood ratio test using the exact permutation distribution to test joint distributions
	Binary	Multinomial - logistic normal random intercept model		Likelihood ratio test using a logistic normal random intercept model to compare marginal distributions whilst modelling the joint distributions

\* Where the overall adverse event profile describes multiple events that are somehow combined for evaluation.

**Table 6: Summary of estimation approaches to analyse emerging AE data in phase II/III RCTs**

Outcome	Data type	Estimate	Reference	Brief Description
<b>Emerging Adverse events (multiple)</b>	Ordinal	Posterior probability	Leon-Novelo, Zhou, Nebiyou Bekele & Muller <sup>23</sup>	Posterior probability of each grade of an AE (participant maximum grade used) allowing multiple different events per participant
<b>Emerging Adverse events (single)</b>	Binary	Frequencies & percentage	Evans & Nitsch <sup>24</sup>	Standard estimates for AE analysis including frequencies, percentages, risk differences and odds ratios
	Binary	Regression models	Evans & Nitsch <sup>24</sup>	Regression based approaches for AE analysis e.g. Poisson regression
	Binary	Confidence interval	O'Gorman, Woolson, Jones <sup>25</sup>	Two methods to estimate CIs for risk-difference when combing data across multiple sites
	Count	Confidence interval	Liu, Wang, Liu & Snavely <sup>26</sup>	Four methods to estimate CIs for exposure adjusted incident ratios
	Binary	Confidence interval	Borkowf <sup>27</sup>	Alternative to the Clopper-Pearson CI for proportions
	Binary	Mean cumulative function	Siddiqui <sup>28</sup>	Non-parametric estimate of mean cumulative number of recurrent events
	Binary	Prevalence	Lancar, Kramar & Haie-Meder <sup>29</sup>	Non-parametric estimate of prevalence of event allowing for recurrence
	Binary	Mean frequency function	Gong, Tong, Strasak & Fang <sup>30</sup>	Non-parametric estimate of mean cumulative number of recurrent events in presence of competing risks
	Binary	Mean cumulative duration	Wang & Quartey, 2012 <sup>31</sup>	Non-parametric estimate of mean cumulative duration for recurrent events
	Binary	Mean cumulative duration	Wang & Quartey, 2013 <sup>32</sup>	Semi-parametric estimate of mean cumulative duration and prevalence of recurrent events
	Binary	Dependence between AEs and discontinuation	Rosenkranz <sup>33</sup>	Three methods to estimate the level of dependence between AE and discontinuation by treatment group that corrects for any dependence in the treatment effect estimate
	Time-to-event	Hazard ratio	Henglebrock, Gillhaus, Kloss & Leverkus <sup>34</sup>	Two methods to estimate hazard ratio for recurrent events

	Time-to-event	Cumulative incidence function	Allignol, Beyersmann & Schmoor <sup>35</sup>	Two methods to estimate the probability of an event in presence of competing risks
	Time-to-event	Conditional cumulative incidence function	Nishikawa, Tango & Ogawa <sup>36</sup>	Probability of a recurrent event in presence of competing risks
<b>(Emerging) Laboratory &amp; vital signs (multiple)</b>	Continuous	GENIE score	Sogliero-Gilbert, Ting, & Zubkoff <sup>37</sup>	Weighted linear combination of absolute normalised deviations from the reference range to indicate abnormalities

**Table 7: Summary of decision making probability methods to analyse prespecified harm outcomes in phase II/III RCTs**

Outcome	Data type	Model	Prior	Reference	Brief Description
<b>Predefined safety outcome</b>	Binary	Beta-Binomial	Beta	Berry <sup>38</sup>	Posterior probability that event rate or incidence rate (incorporating exposure time) is greater in the treatment group compared to control group
	Time-to-event	Exponential	Not specified		
	Binary	Beta-Binomial	Beta	Yao, Zhu, Jiang & Xia <sup>39</sup>	Beta-binomial model to give posterior probability that predefined risk difference threshold is exceeded
	Count	Gamma-Poisson	Gamma	Zhu, Yao, Xia & Jiang <sup>40</sup>	Gamma-Poisson model to give posterior probability that predefined risk difference (incorporating exposure time) threshold is exceeded
	Binary	Logit model	Normal	French, Thomas and Wang <sup>41</sup>	Logit model and a piecewise exponential model to give posterior probabilities that predefined risk difference threshold is exceeded
	Time-to-event	Piecewise exponential	Normal & Gamma		

**Table 8: Summary of decision making probability methods to analyse emerging AE data in phase II/III RCTs**

Outcome	Data type	Model	Prior	Reference	Brief Description
<b>Emerging Adverse events (multiple)</b>	Binary	Logit	Mixed	Berry & Berry <sup>42</sup>	Bayesian hierarchical logit model to give posterior probability that event rate greater in treatment compared to control group
	Binary	Logit	Normal	Xia, Ma & Carlin <sup>43</sup>	Bayesian hierarchical logit and log-linear (incorporating exposure time) models to give posterior probability that event rate greater in treatment compared to control group
	Count	Log (Poisson)	Mixed		
	Count	Log (Poisson)	Normal		
	Binary	Logit	Mixed	Chen <sup>44</sup>	Sequential method. Bayesian hierarchical logit model to give posterior probability that event rate greater in treatment compared to control group for interim analysis
	Binary	Beta-Binomial	Isling	McEvoy <sup>45</sup>	Multivariate approach to give posterior probability of difference in event rates based on indicator functions
	Binary	Beta-Binomial	Beta	Gould <sup>46</sup>	Posterior probability that AEs in treatment group produced by a larger process than AE in control group
	Count	Gamma-Poisson	Poisson	Gould <sup>47</sup>	Posterior probability that AEs in treatment group produced by a larger process than AE in control group accounting for exposure time



## References for tables 3-8

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