# Challenges in NMA Methodology Based on Re-Randomization and Treat-through

content

# Trial Designs

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# **OBJECTIVE**

This study explores the methodological challenges affecting network meta-analyses (NMAs) involving re-randomized (RR) and treat-though (TT) trial designs and potential methodological solutions, using ulcerative colitis (UC) and Crohn's disease (CD) as an example

### **DISCUSSION AND CONCLUSIONS**

- TT and RR trials differ in their purpose and research approach. As a result, they often generate maintenance outcomes that cannot be used in an NMA without prior alignment
- Evidence from active arms can generally be aligned to some degree using different adjustment approaches, while evidence for maintenance placebo arms remains substantially heterogeneous and may not be suitable for inclusion in mixed design NMAs
- Different methodologies have been utilized to combine evidence from RR and TT trials. Approaches incorporating adjustment to individual patient data (IPD), generally, required fewer assumptions than approaches using aggregate data and external evidence, and may be more robust
- The presence of differing active treatment carry-over effects in RR trials may pose challenges to meeting necessary consistency assumptions to enable NMA and should be considered in future NMA feasibility assessments of emerging UC and CD treatments

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<sup>a</sup>Patients who continue on active treatment are having maintenance dosing of active treatment Note: Blue lines illustrate the path of patients receiving placebo in maintenance

#### RESULTS

Re-Randomized trial to Treat-through trial					
Induction Dosing Period and Response Assessment	Maintenance Dosing Period	Imputation	Data used in NMA		
	Continue active treatment	No imputation			

#### BACKGROUND

- RR and TT are trial designs to assess outcomes for treatments with an induction phase aiming for a rapid reduction of disease symptoms and severity, and a maintenance phase maintaining or further improving outcomes
- NMAs enable a comprehensive comparison of treatment options, when head-to-head trials are not available. To perform NMA, similarity assumptions (amongst others) need to be met, including on study design, population, and outcomes
- As shown in Figure 1, RR and TT trial designs have key differences. This heterogeneity presents challenges when NMAs include both trial designs, especially in the maintenance phase
  - In RR trials, maintenance patients have a history of response to active treatment, this means patients randomized to placebo during induction usually do not enter maintenance and instead get followed-up outside the maintenance trial
  - In TT trials, induction patients on active treatment continue treatment in the maintenance phase independent of induction response. In the placebo arm, usually, only induction responders continue on placebo into maintenance, non-responders are followed-up outside maintenance trial

#### **METHODS**

- A targeted literature review was conducted to identify recent submissions to the National Institute for Health and Care Excellence (NICE) and NMA publications in UC and CD where TT and RR trials are included, as well as methodological publications exploring key sources of uncertainty and approaches addressing these challenges
- Based on the literature identified, we summarized findings, considering different scenarios of data availability
- The review identified 7 recent NICE submissions assessing treatments of UC<sup>1-7</sup> and 4 submissions for CD<sup>8-11</sup>, and a further 3 NMA publications (2 for UC<sup>12,13</sup> and 1 for CD<sup>14</sup>) providing information of interest
- A summary of methodological challenges and potential solutions discussed in the identified sources is presented in Table 1



	treatment induction responders re-randomised in maintenance phase)	Lack of comparable placebo maintenance arm when	No respective adjustments were identified to mimic active treatment withdrawal in RR trials, even when IPD is available for TT trial. To connect active arms adjusted from TT to RR into an evidence
<ul> <li>Induction non-responders</li> <li>Continue active treatment</li> <li>OOOO</li> <li>OOO</li> <li>OO</li></ul>		adjusting TT to RR	network, a second active arm needs to be present in the TT trial and in one of the RR trials.
R Placebo — No comparable approach was identified to replicate a RR placebo arm, only active treatments can be used in an NMA		<b>Carry-over effects from</b> <b>induction to maintenance in RR</b> <b>placebo arms</b>	Network meta-regression can be used to investigate potential carry- over effects by using induction outcomes as covariates in estimating maintenance outcomes. <sup>15</sup> However, induction data needs to be available for maintenance trials, and a strong carry-over effect may
With treat-through trial aggregated data Based on above TT trial example: Induction responders = 10/20 (50%) Maintenance responders = 8/20 (40%) Hence TT to RR using aggregated data would be assuming 8/10 (80%) patients responded in maintenance Assumptions required: 1) Induction response/remission only occurs during the induction phase (no delayed response)			be needed to identify a significant coefficient. Further, this requires a very strong assumption that the carry-over effect is proportional to induction outcomes. Treatment sequence approaches utilizing one placebo-to-placebo arm to replace the placebo arms of all trials have been tested.
<ul> <li>2) Maintenance response/remission occurs only in patients with induction response/remission occurs on the patients with induction response/remission occurs on the patients with induction response/remission occurs on the patients with induction response with induction response with remission occurs on the patients with induction response with remission occurs on the patients with remains occurs of the patients with remains occurs on the patients with r</li></ul>	nse/remission = patient entering the trial		However, the approach breaks randomization and enables confounding due to differences in settings, treatments, and disease severity across trials. <sup>9</sup>
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