Re-randomization increased recruitment and provided similar treatment estimates as parallel designs in trials of febrile neutropenia

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Accepted 2 February 2018; Published online 8 February 2018

Abstract

Objective: Re-randomization trials allow patients to be re-enrolled for multiple treatment episodes. However, it remains uncertain to what extent re-randomization improves recruitment compared to parallel group designs or whether treatment estimates might be affected.

Study Design and Setting: We evaluated trials included in a recent Cochrane review of granulocyte colony-stimulating factors for patients with febrile neutropenia. We assessed the recruitment benefits of re-randomization trials; compared treatment effect estimates between re-randomization and parallel group designs; and assessed whether re-randomization led to higher rates of non-compliance and loss to follow-up in subsequent episodes.

Results: We included 14 trials (5 re-randomization and 9 parallel group). The re-randomization trials recruited a median of 25% (range 16–66%) more episodes on average than they would have under a parallel-group design. Treatment effect estimates were similar between re-randomization and parallel group trials across all outcomes, though confidence intervals were wide. The re-randomization trials in this review reported no loss to follow-up and low rates of non-compliance (median 1.7%, range 0–8.9%).

Conclusions: In the setting of febrile neutropenia, re-randomization increased recruitment while providing similar estimates of treatment effect to parallel group trials, with minimal loss to follow-up or non-compliance. It appears to be safe and efficient alternative to parallel group designs in this setting. © 2018 The Authors. Published by Elsevier Inc. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/).

Keywords: Re-randomization; Randomized controlled trials; Clinical trials; Febrile neutropenia; Re-enrolment; Poor recruitment

1. Background

Febrile neutropenia occurs when neutropenic patients (those with abnormally low neutrophil granulocyte counts) develop fever. It is often a complication for patients with cancer who receive chemotherapy regimens which suppress bone marrow activity. Because chemotherapy is usually given in multiple cycles, patients may develop febrile neutropenia multiple times during the course of their cancer treatment, and each episode of febrile neutropenia would require medical intervention. Standard care for febrile neutropenia is broad-spectrum antibiotics [1]. However, it has been suggested that granulocyte colony-stimulating factor (G-CSF) could be useful in this setting, as it regulates the production of the neutrophil lineage [1]. A number of clinical trials have compared the use G-CSF with antibiotics vs. antibiotics alone in patients with febrile neutropenia.

In a parallel group trial, patients would be enrolled for one episode of febrile neutropenia only; if they experienced further episodes of febrile neutropenia, they would no longer be eligible to participate in the trial. This approach can be inefficient, as a large proportion of febrile...
neutropenia episodes may be ineligible for the trial, which can affect recruitment. The majority of trials in this area have recruited fewer than 50 patients per treatment arm [1], which would lead to underpowered analyses for important outcomes such as mortality.

An alternative approach is a re-randomization trial (Fig. 1) [2–4]. In re-randomization trials, patients can be re-enrolled and re-randomized for each new episode of febrile neutropenia they experience. The number of times each patient is enrolled in the trial is not specified in advance, but instead depends on the number of febrile neutropenia episodes they experience during the course of the trial; some patients may be enrolled only once, and others may be enrolled multiple times. Because patients can be enrolled for multiple episodes, re-randomization can increase the recruitment rate compared to parallel group designs, which could facilitate quicker and more efficient trials [2,3].

However, there has been little empirical evaluation of re-randomization trials, and so, it is unclear how much of a recruitment benefit might be expected in practice or whether treatment effect estimates from re-randomization trials might differ to those from parallel group designs. Furthermore, there may be concern that repeated enrollments in re-randomization trials may place undue burden on patients due to increased treatment or follow-up burden and may lead to higher rates of non-compliance or loss to follow-up in subsequent enrollments. We therefore undertook a review of trials in febrile neutropenia to evaluate (1) the impact re-randomization had on recruitment; (2) whether treatment effect estimates from re-randomization trials were different to those from parallel group trials; and (3) whether re-randomization led to higher rates of non-compliance and loss to follow-up in subsequent episodes.

2. Methods

2.1. Overview of re-randomization trials

We begin by providing a brief overview of the re-randomization design (Table 1). This design is appropriate in settings where at least some patients may require treatment on multiple occasions, and in practice, the intervention(s) under study would be used for each new treatment episode that occurred [2,3]. Furthermore, the duration of the intervention and the length of the patient follow-up period must be less than the overall length of the trial recruitment period [2,3]. This design is therefore suitable in the setting of febrile neutropenia, as some patients experience multiple episodes and require treatment for each episode, and the intervention (G-CSF) and patient follow-up duration are typically short-term.

There are two core design requirements for re-randomization trials [2,3]: (1) patients are only re-enrolled and re-randomized after the follow-up period from their previous enrollment is complete (i.e., there cannot be overlapping follow-up periods from different enrollments); and (2) randomizations for the same patient are performed independently (e.g., patients are not forced to crossover from one treatment arm to another between episodes). Analysis of re-randomization trials can be via an “independence” analysis [2], where each episode is analyzed independently (i.e., the correlation between episodes from the same patient is ignored in the analysis). This approach can provide unbiased estimates and correct type I error rates [2]. It will also provide the same power as a parallel group design with an equivalent number of observations in many settings, provided the overall variance is not increased through the use of re-randomization; further details are available in another article [2]. Therefore, in these settings, the same sample size calculation as in a parallel group design could be used; however, instead of recruiting the required number of patients, the re-randomization trial could recruit the required number of treatment episodes. For example, if the sample size calculation for a parallel group trial required 100 patients, a re-randomization trial would require 100 episodes of febrile neutropenia from fewer patients, for example, 100 episodes from 75 patients (where 50 patients...
contributed one episode each, and 25 patients contributed two episodes each) or some other combination.

2.2. Review of trials in febrile neutropenia

We used the Cochrane Collaboration systematic review and meta-analysis conducted by Mhaskar et al. [1] as the basis of our study. Full details of the search strategy, inclusion criteria, and data collection procedure are available in their publication [1]. The review included 14 trials; 9 used a parallel-group design, and 5 allowed re-randomization [5—9]. Information was extracted by two reviewers (B.C.K and T.P.M.), and discrepancies were resolved through discussion.

2.3. Impact of re-randomization on recruitment

For the five trials using re-randomization, we extracted information on the number of individual patients enrolled in the trial and the number of treatment episodes enrolled. Based on this information, we estimated the increased sample size obtained through the use of re-randomization for each trial by dividing the total number of treatment episodes by the number of individual patients. This measure represents the extra number of episodes each trial gained by using a re-randomization design instead of a parallel group design.

2.4. Difference in treatment effect estimates between re-randomization and parallel group trials

For each trial, we extracted the treatment effect estimate and 95% confidence interval (95% CI) for each outcome reported. In our analysis, we only included outcomes that were available for at least two re-randomization and two parallel group trials.

For each outcome, we used the Stata package `metareg` to conduct a random-effects meta-regression model to estimate the difference in treatment effect estimates between re-randomization and parallel group trials [10]. Treatment effect estimates were log(hazard ratio) for time-to-event outcomes, log(risk ratio) for binary outcomes, and standardized mean differences for continuous outcomes. A negative difference in effect sizes indicates that re-randomization trials show a more beneficial treatment effect than parallel group trials.

Two re-randomization trials did not have any events for overall mortality, and two trials (one re-randomization and one parallel group) did not have any events for infection-related mortality. These trials were excluded from the analysis of these outcomes, as it was impossible to estimate either a treatment effect or standard error. One parallel group trial included three treatment arms (two active and one control) and involved two treatment comparisons (both active interventions vs. control). The meta-analysis by Mhaskar et al.


**Table 1. Overview of re-randomization trials**

<table>
<thead>
<tr>
<th>Settings requirements for re-randomization trials</th>
</tr>
</thead>
<tbody>
<tr>
<td>1) Some patients may require treatment on multiple occasions</td>
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<tr>
<td>2) The intervention(s) would be used for each new treatment episode</td>
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<tr>
<td>3) The intervention duration and length of the follow-up period for each treatment episode are less than the overall length of the trial recruitment period</td>
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<table>
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<tr>
<th>Design requirements for re-randomization trials</th>
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<tbody>
<tr>
<td>1) Patients are only reenrolled and rerandomized when they have completed the follow-up period from their previous randomization</td>
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<td>2) Randomizations for the same patient are performed independently</td>
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<tr>
<th>Implementation of re-randomization trials</th>
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<tr>
<td>1) Patients are enrolled as usual, randomized to a treatment group, and followed up until all outcomes have been collected</td>
</tr>
<tr>
<td>2) If patients experience new treatment episodes and require further treatment, they can be reenrolled and rerandomized, provided they have completed the follow-up period from their previous randomization</td>
</tr>
<tr>
<td>3) This process is repeated until the target sample size is met</td>
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![Fig. 2. Percentage increase in number of episodes recruited due to re-randomization.](image-url)
2.5. Treatment compliance and loss to follow-up in re-randomization trials

In order to assess whether re-randomization led to higher rates of non-compliance or loss to follow-up in later episodes, we extracted data on the number of treatment episodes which did not comply with the treatment protocol, and for outcomes reported by two or more re-randomization trials, the number of treatment episodes excluded from the analysis due to missing data. We attempted to extract these data separately for each episode, to assess whether non-compliance and missing data were higher in later episodes. However, this information was not reported in any trial, and so, we extracted the overall rates of non-compliance and loss to follow-up across all episodes.

3. Results

3.1. Impact of re-randomization on recruitment

Results are shown in Fig. 2. Among the five trials using re-randomization, the median number of individual patients recruited was 40 (range 28–112) and the median number of episodes of febrile neutropenia enrolled was 58 (35–186). The median increase in the sample size obtained through

![Table showing differences in effect sizes between re-randomization and parallel group trials.](image)

**Fig. 3.** Difference in effect sizes between re-randomization and parallel group trials. The blue lines represent the estimated treatment effect and 95% CI from the re-randomization and parallel group trials. The red lines represent the difference in the treatment effect estimates between re-randomization and parallel group trials (and a 95% CI for this difference). If the red line is close to 0, it means that re-randomization and parallel group trials are providing similar estimates of treatment effect; if it is far away from 0, then re-randomization and parallel group trials are giving different estimates of treatment effect. The x-axis shows the size of the effect for both the blue and red lines; however, the x-axis text (whether re-randomization or parallel group trials show more beneficial effects) applies only to the red line. RR, re-randomization; PG, parallel group; CI, confidence interval; Std. Mean Diff., standardized mean difference. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)
the use of re-randomization was 25% (range 16—66%), indicating that using a re-randomization design allowed trials to recruit between 16% and 66% more episodes of febrile neutropenia than they would have under a parallel group design.

3.2. Difference in treatment effect estimates between re-randomization and parallel group trials

Differences in treatment effect estimates between re-randomization and parallel group trials are shown in Fig. 3. Treatment effect estimates for the two designs were similar for each of the five outcomes, and none of the differences were statistically significant. However, CIs were wide, indicating that differences were possible.

3.3. Treatment compliance and loss to follow-up in re-randomization trials

Results are shown in Table 2. The median percentage of episodes which were not compliant with the protocol was 1.7% (range 0—8.9%). None of the five outcomes we assessed excluded any episodes from the analysis due to missing data.

3.4. Characteristics of re-randomization trials

Results are shown in Table 3. None of the five trials explicitly stated whether randomizations for the same patient were independent or that patients were only re-enrolled once the follow-up period from their previous enrollment was complete.

Of the trials that reported a sample size calculation, each based their calculation on a parallel group design. All trials analyzed the data on a per-episode basis and all used an independence analysis (i.e., ignored correlation between episodes from the same patient), and none of the trials adjusted for any factors associated with re-randomization in the analysis (e.g., whether it was the patient’s first or second time enrolled in the trial).

4. Discussion

We found that using re-randomization in trials of febrile neutropenia increased the number of episodes recruited by between 16% and 66%. Furthermore, we found that re-randomization trials provided similar estimates of treatment effect to parallel group designs. Given that many trials face challenges in recruitment [11–14], re-randomization may be a viable option in appropriate settings to facilitate more efficient recruitment than parallel group designs, while providing similar results. We found that re-randomization trials had minimal non-compliance and no loss to follow-up, indicating that increased patient burden due to trial re-enrollment was not an issue in this setting. This may be because most outcomes were recorded by the trial team, rather than by the patients themselves.

Most re-randomization trials were designed and analyzed using very simple approaches; they used the same
sample size calculation and analysis method as would be used in a parallel group trial, except instead of recruiting and analyzing patients, they recruited and analyzed episodes. This approach will lead to adequately powered trials in many settings [2] and can provide unbiased estimates of treatment effect and correct type I error rates [2]. Therefore, using a re-randomization design rather than a parallel group trial design does not necessarily require additional methodological complexity and can be done in a very simple way.

This is the first review to assess the use of re-randomization trials in practice. However, there were some limitations. There were only a small number of trials available. This led to wide CIs for the differences in treatment effect estimates, meaning that we could not rule out differences between designs. We also focused only on one clinical area, and so, these results may not be generalizable to other settings. Furthermore, reporting of key trial characteristics in re-randomization trials was often inadequate, which may in part reflect a lack of guidance on good reporting practice.

5. Conclusions

In the setting of febrile neutropenia, re-randomization increased recruitment while providing similar estimates of treatment effect to parallel group trials, with minimal loss to follow-up or non-compliance. It appears to be safe and efficient alternative to parallel group designs in this setting.

Acknowledgments

Table 1 was reproduced from Kahan BC. Using re-randomization to increase the recruitment rate in clinical trials—an assessment of three clinical areas. Trials 2016;17:595. doi:10.1186/s13063-016-1736-z with permission.

Authors’ contributions: B.C.K. concepted the study, developed review, performed data extraction, and wrote the manuscript. T.P.M. developed review, performed data extraction, and contributed to the manuscript. E.H. developed review and contributed to the manuscript. R.P., R.H., and S.E. contributed to the manuscript.

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