



Methodological insights

# Risk and treatment effect heterogeneity: re-analysis of individual participant data from 32 large clinical trials

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### Abstract

**Background:** Risk of the outcome is a mathematical determinant of the absolute treatment benefit of an intervention, yet this can vary substantially within a trial population, complicating the interpretation of trial results.

**Methods**: We developed risk models using Cox or logistic regression on a set of large publicly available randomized controlled trials (RCTs). We evaluated risk heterogeneity using the extreme quartile risk ratio (EQRR, the ratio of outcome rates in the lowest risk quartile to that in the highest) and skewness using the median to mean risk ratio (MMRR, the ratio of risk in the median risk patient to the average). We also examined heterogeneity of treatment effects (HTE) across risk strata.

**Results**: We describe 39 analyses using data from 32 large trials, with event rates across studies ranging from 3% to 63% (median = 15%, 25th–75th percentile = 9–29%). C-statistics of risk models ranged from 0.59 to 0.89 (median = 0.70, 25th–75th percentile = 0.65–0.71). The EQRR ranged from 1.8 to 50.7 (median = 4.3, 25th–75th percentile = 3.0–6.1). The MMRR ranged from 0.4 to 1.0 (median = 0.86, 25th–75th percentile = 0.80–0.92). EQRRs were predictably higher and MMRRs predictably lower as the c-statistic increased or the overall outcome incidence decreased. Among 18 comparisons with a significant overall treatment effect, there was a significant interaction between treatment and baseline risk on the proportional scale in only one. The difference in the absolute risk reduction between extreme risk quartiles ranged from -3.2 to 28.3% (median = 5.1%; 25th–75th percentile = 0.3–10.9).

**Conclusions:** There is typically substantial variation in outcome risk in clinical trials, commonly leading to clinically significant differences in absolute treatment effects. Most patients have outcome risks lower than the trial average reflected in the summary result. Risk-stratified trial analyses are feasible and may be clinically informative, particularly when the outcome is predictable and uncommon.

Key words: Risk prediction, heterogeneity of treatment effect, subgroup analysis, personalized medicine, patientcentered outcomes research

#### **Key Messages**

- Outcome risk is a mathematical determinant of the treatment effect yet can vary substantially across a trial population, making it unclear how treatment effects might vary in the trial population.
- Using simple risk models based on baseline patient characteristics, among a sample of trials from publicly available sources, we found that outcome rates in the highest risk quartile were as high as 50-fold those in the lowest risk quartile; in fully a quarter of the trials, this ratio exceeded 6.
- Because outcome risk in the trials was generally skewed (log-normal or logistic-normal), with a small group of highrisk patients accounting for a large number of outcomes, the outcome risk in most patients was almost always less than that reflected by the trial summary results.
- Whereas we did not often detect treatment effect heterogeneity on the proportional scale across patients at different baseline risk in this set of trials, substantial differences in absolute treatment effects were common; differences in absolute treatment effects between the extreme quartiles of risk exceeded 10% in a quarter of trials that showed benefit.
- Displaying results across subgroups defined by risk is feasible and can lead to clinically important findings.

#### Introduction

A fundamental incongruity in evidence-based medicine (EBM) is that evidence is derived from groups of people yet medical decisions are made for individuals. Popular approaches to EBM have encouraged the direct application of average effects estimated in clinical trials to guide decision making for individuals, as though all patients meeting trial inclusion criteria are likely to experience similar effects from treatments. This simplistic attitude has proven remarkably durable and compelling, despite the variation in patient characteristics and outcomes seen in clinical practice.<sup>1</sup>

The most commonly used method of examining whether treatment effects vary in a trial population is to serially divide patients into subgroups based on potentially relevant pre-treatment characteristics. The main problem with this conventional approach is that there are too many potentially influential characteristics. This leads to myriad 'one-variable-at-a-time' subgroup analyses, which are typically both underpowered and vulnerable to false-positive results due to multiple comparisons.<sup>2,3</sup> It can also be difficult to understand how to apply such analyses to individuals in clinical practice, because patients have multiple characteristics that vary from one another simultaneously.

In part for these reasons, subgroup analyses are usually 'exploratory' and rarely actionable, leaving the clinician to assume that all patients meeting trial inclusion criteria should be similarly treated. EBM is thus methodologically canalized to 'one-size-fits-all' recommendations, a problem increasingly recognized even as EBM has become the dominant paradigm.<sup>4–6</sup> This remains a central challenge to be addressed if EBM is to become more personalized and patient-centred.<sup>4–6</sup>

We recently proposed a framework for assessing heterogeneity of treatment effect (HTE) that seeks to address these issues.<sup>7</sup> The framework prioritizes the analysis and reporting of multi-variable risk-based HTE and suggests that other subgroup analyses should be explicitly labelled either as primary subgroup analyses (well-motivated by prior evidence and intended to produce clinically actionable results) or secondary (exploratory) subgroup analyses (performed to inform future research). Whereas other recommendations or guidance documents have (appropriately) emphasized the risks of overinterpreting the results of subgroup analyses,<sup>8,9</sup> and the different goals of such analyses,<sup>10</sup> our framework is novel in that it also suggests that presenting summary results without examining and reporting how treatment effects change across subgroups with heterogeneous outcome risk is under-utilizing trial data and tantamount to incompletely reporting trial results.

Despite compelling theoretical arguments, a riskmodelling approach is rarely applied. Empirical evidence for its importance remains anecdotal and there are concerns about the feasibility of routine and broad application of this analytical approach in datasets collected in typical randomized trials. To address these concerns, we examined the distribution of outcome risk across a broad range of trials and examine how the effects of therapy were related to this risk.

#### Methods

We searched for publicly available individual participant datasets of randomized clinical trials from the National Heart, Lung, and Blood Institute (NHLBI),<sup>11</sup> the National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK),<sup>12</sup> the journal *Trials* and GlaxoSmithKline.<sup>13</sup> We required that eligible studies had enrolled at least 1000 participants (some subcohorts entered in our analyses had fewer than 1000 participants) randomized to at least two treatment groups, and had a binary (or time-to-event) clinical (i.e. not surrogate) outcome.

#### Predicting outcome risk using baseline covariates

Risk modelling for each trial was informed by examining previously developed published predictive models 'matched' to each trial on the basis of the index condition of the population and the primary outcome.<sup>14</sup> We identified risk predictors that had been used in the published models and the corresponding variables in the trial datasets. Because trial datasets were often not fully compatible with externally developed predictive models, we developed 'internal models' on the trial data using risk predictors that were as close as possible to those in published models. To verify that the use of internal models would not bias estimates of HTE across risk groups, we performed a series of simulations described in a separate publication.<sup>15</sup> Briefly, the simulations revealed that, across a range of scenarios, analyses based on internal models developed on trial participants yield results similar to analyses based on external models developed on non-trial participants sampled from the same population.

All available established risk predictors were entered into a regression model to predict the primary outcome for all patients in the trial. Both trial arms were used in model development, without using the treatment assignment indicator, to avoid differential model fit between the trial arms, potentially inducing a spurious risk-by-treatment interaction.<sup>15</sup> To minimize model complexity for trials for which there were many established predictors, nonsignificant risk predictors were ranked in order of significance and removed until no more than 20 variables were entered into the model (this was needed in only 3 of the 32 trials). No other formal variable selection process or attempt at model re-specification was performed.

In trials with non-statistically significant overall treatment effects for the primary outcome and a statistically significant treatment effect for a binary (or time-to-event) clinical secondary outcome, an additional regression model was fit to predict the secondary outcome. When treatment effects for multiple secondary outcomes were statistically significant, we selected the outcome identified as most clinically relevant in the published trial report.

To minimize bias due to missing data, multivariate normal multiple imputation was used when a complete case analysis would exclude more than 5% of trial participants. Risk factors with missing information from more than 20% of trial participants were not used in analyses.

The statistical analysis model (Cox proportional hazards regression for time-to-event outcomes or logistic regression for binary outcomes) was selected on the basis of the primary analysis of the clinical trial and determined by the nature of the trial data. In general, we included variables as main effects in their original scale, unless published predictive models specified the use of interactions or variable transformations.

Model performance was assessed with respect to discrimination, calibration and overfitting. Discriminatory ability was quantified using the c-statistic.<sup>16</sup> Calibration was assessed visually using calibration plots. Overfitting was assessed with bootstrap validation.<sup>17</sup> We report the number of events per variable in each trial as an indicator for the risk of overfitting.

We evaluated the distribution of predicted risk in the overall study population and separately in each treatment arm. Visual examination of the risk distribution was facilitated by the use of box plots of the predicted risk of the outcome. In addition, we plotted histograms of the empirical distribution of predicted risk in each study to assess how closely the distribution conformed to the truncated log-normal (for risk predicted by proportional hazard models) or the logistic-normal distribution (for risk predicted by logistic regression models).

To describe and quantify risk heterogeneity using clinically interpretable metrics, we used two indexes, the extreme quartile risk ratio (EQRR) and the median-to-mean risk ratio (MMRR). To calculate the EQRR, we stratified the trial population into equal-sized quartiles according to the baseline predicted risk from the model.<sup>18</sup> We then calculated the ratio of the predicted outcome risk in the extreme quartiles (high-risk quartile outcome probability divided by the low-risk quartile outcome probability, EQRR<sub>predicted</sub>). We also calculated the same index based on the observed outcome rate (EQRR<sub>observed</sub>) within strata defined by predicted risk. Greater EQRR values indicate greater risk heterogeneity in the risk-stratified patient population. The MMRR is a clinically interpretable measure of skewness calculated as the ratio of the median predicted outcome probability to the mean predicted outcome probability. As the MMRR deviates from one, it reflects the degree to which the summary (average) result may not reflect the effects in the 'typical' patient in the trial. We also calculated Pearson's median skewness coefficient [3\*(mean-median)/standard deviation], a more common measure of skewness.

We also examined the relationship between the outcome prevalence and the c-statistic, and the EQRR and MMRR, visually and using linear regression.

#### Evaluating HTE over predicted outcome risk

Additionally, we analysed the relationship between treatment effect and predicted outcome risk. We estimated treatment effects within each risk quartile on relative and absolute scales. Specifically, we estimated relative treatment effects using logistic regression (using odds ratios as the measure of effect) or Cox regression (using hazard ratios as the measure of effect); we estimated absolute treatment effects using linear probability models for binary outcomes (using absolute risk reduction as the measure of effect). For time-to-event analyses, we calculated absolute risk reduction as the difference in Kaplan-Meier survival probabilities between the intervention and comparator treatment arms.<sup>19</sup> We tested the null hypothesis of no HTE over predicted outcome risk using a product term ('interaction') between the fitted value of the linear predictor (from the risk model) and the treatment assignment indicator. We also compared relative and absolute risk reduction between the extreme risk quartiles in each trial. We summarized these metrics for the subset of trials with statistically significant overall treatment effects, i.e. those trials showing statistically significant benefit or harm on either a primary or a secondary outcome.

Statistical analyses were performed using SAS version 9.3,<sup>20</sup> R open-source software version 3.1.2 (The R Foundation for Statistical Computing) and Stata version 13.1 (Stata Corp., College Station, TX).

#### Results

A total of 32 trials met our inclusion criteria (Table 1). Most trials were in the field of cardiovascular disease, including trials evaluating interventions in atrial fibrillation, coronary heart disease, acute myocardial infarction, heart failure, hypertension and acute stroke. We also included trials of other conditions, such as prediabetes, acute kidney failure, chronic hepatitis C and prostatic hyperplasia. The number of patients in the analysed trial cohorts ranged from 715 to 33 357, and totalled 180 291. Trials had been conducted over a span of several decades; the earliest trial had been published in 1979 and the latest in 2008. Of note, our trials generally did not include interventions with harms anticipated to affect the primary outcome (e.g. as in carotid endarterectomy, which both prevents and causes stroke).

One trial had more than one patient cohort  $(DCCT^{21})$ . one trial had more than one primary outcome (IST<sup>22</sup>) and five trials had non-statistically significant results for their primary outcome but significant results for a secondary outcome (ACCORD,<sup>23</sup> ALLHAT HTN,<sup>24</sup> BEST,<sup>25</sup> DIG,<sup>26</sup> SOLVD<sup>27,28</sup>). Thus, we developed a total of 39 separate risk models. The median number of risk factors used in these models was 10 (average = 10.9; range = 4-20) (Table 2). The median number of events per variable was 51.3 (average = 107.0; range = 12.5-907.1), suggesting that models were unlikely to overfit the data. The median c-statistic was 0.69 (average = 0.70; range = 0.59-0.89). Bootstrap validation produced optimism-corrected c-statistics in the range of 0.58 to 0.88 (median = 0.68, 25th-75th percentile = 0.64-0.70). The difference between original and optimism-corrected c-statistics ranged from 0.001 to (median = 0.007, 25th - 75th percentile = 0.004 -0.02 0.009), again suggesting the absence of substantial overfitting.

# Distribution of predicted outcome risk in large randomized trials

The median overall event rate across the trials was 15% (average = 20%; range = 3–63%). Summary statistics describing the risk heterogeneity of the population are shown in Table 2. The median EQRR<sub>observed</sub> was approximately 4, but more than a quarter of all analyses had an EQRR<sub>observed</sub> over 6 and the range extended to 50. Values of EQRR<sub>predicted</sub> corresponded closely to the observed values. Whereas the median MMRR was 0.86 (indicating that the typical patient was at 86% the outcome risk compared with the average), this index ranged as low as 0.4—and only twice exceeded 1 (ATN,<sup>29</sup> IST<sup>22</sup> 6-month outcome), both times for trials with high outcome rates (52.6% in ATN and 62.6% in IST).

We found the overall outcome rate in the trial and the c-statistic were strong predictors of the risk distribution. In linear regression, the outcome rate and c-statistic were shown to strongly predict the EQRR ( $R^2 = 0.86$ ) and the MMRR ( $R^2 = 0.78$ ) (Table 3). As discrimination improved, and as the overall outcome rate was lower, EQRR

Trial acronym	Year of publication	Patients randomized ( <i>n</i> )	Patient population/ index condition	Intervention	Comparator	Primary outcome <sup>a</sup>	Secondary outcome <sup>a</sup>
ACCORD <sup>b23</sup>	2008	10251	Type 2 diabetes mellitus	Intensive strategy	Standard treatment	First occurrence of a major	All-cause mortality (-)
AFFIRM <sup>51</sup>	2002	4060	Atrial fibrillation/risk of stroke or death	Rate control therapy	Rhythm control therapy	All-cause mortality	Not assessed
ALLHAT HTN <sup>b,c24</sup>	2002	33357	Hypertension	Amlodipine or lisinopril	Chlorthalidone	Fatal CHD or nonfatal MI comhined	Combined CVD events (-)
ALLHAT LLT <sup>52</sup>	2002	10355	Hypercholesterolaemia/	Pravastatin	Usual care	All-cause mortality	Not assessed
A MIS <sup>30</sup>	1 980	4524	hypertension Mvocardial infarction	Aspirin	Placeho	All-cause mortality	Not assessed
$ATN^{29}$	2008	1124	Acute kidney failure/sepsis	Intensive renal-replacement	Less intensive renal-	All-cause mortality (60-day)	Not assessed
				therapy	replacement therapy		
BARI <sup>53</sup>	1996	1829	Coronary artery disease/se-	Percutaneous transluminal	Coronary artery bypass	Cardiac mortality	Not assessed
				angioplasty	Summer S		
BEST <sup>b25</sup>	2001	2708	Advanced heart failure/con- gestive heart failure	Bucindolol hydrocholoride	Placebo	All-cause mortality	Death due to cardiovascular causes (+)
BHAT <sup>54</sup>	1982	3837	Acute myocardial infarction	Propranolol	Placebo	All-cause mortality (+)	Not assessed
CAST <sup>55</sup>	1 9 9 1	1498	Mvocardial infarction	Class I and Ib ant-jarrhyth-	Placeho	All-cause mortality or	Not assessed
				mic agents		cardiac arrest (-)	
CPPT <sup>56</sup>	1984	3806	Hypercholesterolaemia	Cholestyramine	Placebo	CHD death and/or definite nonfatal myocardial infarc-	Not assessed
DCCT <sup>b21</sup>	1993	Prevention: 726 intervention: 715	Type 1 diabetes mellitus	Intensive diabetes therapy	Conventional diabetes therapy	Appearance and/or progres- sion of retinopathy and other comblications (+)	Not assessed
DIG <sup>b26</sup>	1997	6800	Heart failure	Digoxin	Placebo	All-cause mortality	Hospitalization for worsen- ing heart failure (+)
D.D.033	100 L	1000	A = = = = = = = = = = = = = = = = = = =	()	DI	D ]	Mart manute (⊤)
DIT	7007	1070	AUTION TOL UTADELES INCLINUS	1) metror mut 2) mutansive lifestyle intervention	r lacedo	Development of maderes $(\pm)$	INUL assessed
ENRICHD <sup>57</sup>	2003	2481	Acute myocardial	Cognitive behaviour ther-	Usual medical care	All-cause mortality or recur-	Not assessed
			infarction	apy-based intervention		rent myocardial infarction	
FAVORIT <sup>58</sup>	2011	4110	Stable kidney transplant	Multivitamin plus folic acid, vitamin B <sub>12</sub> and	Treatment with an identical multivitamin alone	Arteriosclerotic cardiovascu- lar disease outcome	Not assessed
;				vitamin B <sub>6</sub>			
FUTURA	2010	2026	Unstable angina/	Low-dose unfractionated heparin	Standard-dose unfractio- nated heparin	Peri-PCI major bleeding, minor bleeding, major vas- cular access site comnlications	Not assessed
			Non-STEMI				
HALTC <sup>60</sup>	2008	1050	Chronic hepatitis c	Indefinite pegylated inter- feron alpha-2a (beyond 3.5 years of use)	Pegylated interferon alpha- 2a, discontinue use after 3.5 years	Progression to cirrhosis	Not assessed
							(Continued)

Table 1. Description of trials

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l rial acronym	Year of publication	Patients randomized $(n)$	Patient population/ index condition	Intervention	Comparator	Primary outcome"	Secondary outcome"
HDFP <sup>61</sup>	1979	10940	Hypertension	Stepped care antihyperten- sive therany	Referred care	All-cause mortality (+)	Not assessed
HEMO <sup>c62</sup>	2002	1846	Haemodialysis	High-dose/high-flux	Standard-dose/low-flux	All-cause mortality	Not assessed
IST <sup>b22</sup>	1997	19435	Acute stroke	atatysis Unfractionated heparin, 	dialysis Placebo	Death within 14 days, death/	Not assessed
MAGIC <sup>63</sup>	2002	6213	Acute myocardial	aspirin Intravenous magnesium	Placebo	dependency at 6 months All-cause mortality	Not assessed
MRFIT <sup>64</sup>	1982	12866	infarction At risk for coronary heart disease	sulphate Stepped-care treatment, counselling, dietary	Usual care	Death from coronary heart disease	Not assessed
MT OP S <sup>d 32</sup>	2003	3047	Benign prostatic hyperplasia	advice 1)doxazosin, 2) finasteride or 3) combination	Placebo	Clinical progression of benign prostatic hyperplasia (+)	Not assessed
OAT <sup>65</sup>	2006	2166	Congestive heart failure	therapy Routine PCI and stenting with optimal medical	Optimal medical therapy alone	Mortality, recurrent MI, and hospitalization for CHF	Not assessed
PEACE <sup>66</sup>	2004	8290	Coronary artery disease	therapy Trandolapril	Placebo	Death from cardiovascular	Not assessed
ROC <sup>b,c67,68</sup>	HS: 2011	HS:895	Hypovolaemic shock	Hypertonic saline solution	Normal saline solution	causes or non-fatal MI 28-day survival, 6-month neurological outcome based on the extended	Not assessed
SHEP <sup>69</sup>	TBI: 2010 1991	TBI: 1331 4736	Traumatic brain injury Hypertension	Chlorthalidone/atenolol antihypertensive drug	Placebo	Glasgow Outcome Scale Non-fatal and fatal stroke (+)	Not assessed
SOLVD <sup>e27,28</sup>	Prevention: 1992 intervention: 1991	Prevention: 4228 intervention: 2569	Congestive heart failure	regimen Enalapril	Placebo	All-cause mortality (T+)	Death or hospitalization for heart failure (P+)
TIMI-II <sup>70</sup> 32 trials (33 cohorts)	1989	3262 180291	Acute myocardial infarction	Invasive strategy	Conservative strategy	All-cause mortality or non- fatal MI	Not assessed 18 positive treatment effects From 14 trials

admike έ. trials have one cohort; PCI, percutaneous coronary intervention.

<sup>a</sup>Summary treatment effect on outcome is statistically insignificant unless indicated by sign: (+) indicates positive treatment effect, (-) indicates treatment harm.

<sup>b</sup>Indicates two risk distributions.

<sup>c</sup>Indicates two treatment arms. <sup>d</sup>Indicates three treatment arms. <sup>e</sup>Indicates three risk distributions (if not specified, assume one risk distribution).

increased and MMRR decreased in a predictable fashion. Indeed, we found that knowing these two parameters (overall outcome incidence and c-statistic) essentially determine the full distribution of predicted risk, because the risk distributions were close to the log-normal (for risk predicted using Cox models) or logistic-normal shape (for risk predicted using logistic regression models) (Figure 1). This can be seen by comparing the histograms and kernel densities of the predicted values (in black) against the lognormal (red) or logistic-normal densities (blue) fit to the same values via maximum likelihood, which were fairly similar in most studies.

#### HTE over-predicted outcome risk

Among the 18 trials with statistically non-significant results, two trials showed statistically significant HTE over the estimated linear predictor from the risk model. In the AMIS trial,<sup>30</sup> high-risk patients with acute myocardial infarction appeared to get more benefit from aspirin than low-risk patients (P = 0.02) on the proportional scale; in IST,<sup>22</sup> for the combined outcome of death or dependency at 6 months, low-risk patients appeared to obtain more benefit than high-risk patients (P = 0.04) on the proportional scale.

	Median	25th–75th percentile	Mean	Range
Overall event rate	0.15	0.09-0.29	0.20	0.03-0.63
Model risk predictors	10	7–16	10.9	4-20
Events per variable	51.3	32.3-84.7	107.0	12.5-907.1
c-statistic	0.69	0.65-0.71	0.70	0.59-0.89
EQRR observed	4.3	3.0-6.1	6.1	1.8-50.7
EQRR predicted	4.0	3.1-5.4	5.3	1.9-35.2
MMRR	0.86	0.80-0.92	0.84	0.42-1.04
PMSC	0.74	0.60-0.86	0.70	-0.24-1.56

EQRR, extreme quartile risk ratio; MMRR, median-to-mean risk ratio; PMSC, Pearson's median skewness coefficient.

Table 3.	Regression	model	results
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In the 14 trials with statistically significant results, 18 unique treatment comparisons were analysed. Although the relative treatment effects appeared to decrease over risk quantiles in some trials (e.g. BEST, CPPT and MTOPS [Figure 2a]) and increase over risk quantiles in others (ACCORD, CAST and DPP [Figure 2a]), overall there was no apparent relationship between baseline risk and the hazard (or odds) ratio of treatment across trials. The median ratio of the hazard or odds ratio in the fourth quartile over that in the first quartile was 1.02 (25th-75th percentile = 0.70-1.21) (Table 4). We found a statistically significant interaction between treatment and the estimated linear predictor on the proportional scale only in one of 18 analyses -(DPP, metformin vs placebo; high-risk patients experienced greater benefit than low-risk patients; P = 0.0008). Despite the absence of 'statistically significant' HTE on the proportional scale, absolute risk reduction estimates varied substantially over predicted outcome risk and were generally higher in high-risk strata, ranging from -1.4% to 18.3% (median = 4.7%; 25th-75th percentile = 0.8-6.1%) in the first quartile of predicted risk and from 0.8% to 35.0% (median = 9.0%; 25th-75th percentile = 3.3-19.8%) in the fourth quartile. The difference in the absolute risk reduction between the extreme-risk quartiles ranged from -3.2% to 28.3% (median = 5.1%;  $25^{\text{th}}$ -75th percentile = 0.3-10.9) across studies. Figure 2b displays these absolute effects graphically.

#### Discussion

Our results show that clinically significant risk heterogeneity is common even in phase III 'efficacy' trials, which are often characterized as enrolling relatively homogeneous populations. Whereas statistically significant HTE on the proportional scale was unusual in this set of trials, in which interventions generally did not have anticipated harms on the primary outcome, variability in risk often gave rise to substantial HTE on the absolute risk scale. Though it is most common to test for heterogeneity on the proportional scale, absolute risk reduction (and its inverse, the number needed to treat) are generally considered the most relevant

	log	EQRR predicted			MMRR	
	Estimate (SE)	t-Value	P-value	Estimate (SE)	t-Value	P-value
Intercept	-3.88 (0.39)	-10.05	< 0.0001	1.80 (0.12)	15.47	< 0.0001
Overall event rate	-1.94(0.24)	-7.98	< 0.0001	0.66 (0.07)	9.06	< 0.0001
c-statistic	8.27 (0.57)	14.41	< 0.0001	-1.57(0.17)	-9.05	< 0.0001
R-square	0.86			0.78		

EQRR, extreme quartile risk ratio; MMRR, median-to-mean risk ratio; SE, standard error.



Figure 1. Risk distributions. The histograms show the distribution of the predicted risk for the outcome of interest. Curves shown in red are fitted to the distribution of predictions generated by Cox models; curves shown in blue are fitted to the distribution of predictions generated by logistic models. Fitted log-normal curves and fitted logistic-normal curves are also shown for the Cox- and logistic regression-generated curves, respectively. As can be seen, these log-normal and logistic-normal curves approximate very well the red and blue fitted curves. Note: The FUTURA Trial is not included in this figure since we could not export individual-level patient predictions from the site in which the data were housed.

scales for clinical decision making.<sup>31</sup> We did not use formal criteria to assess clinically important HTE, but it is note-worthy that, among treatment comparisons with statistically significant overall results, 25% showed differences in absolute risk differences greater than 10% between the extreme quartiles of predicted risk. We considered our analysis of two trials (MTOPS<sup>32</sup> and DPP<sup>33</sup>), encompassing 5 of our 18 treatment comparisons, to be of sufficient clinical interest to report in separate clinical manuscripts.<sup>34,35</sup> These papers join a growing list of papers showing clinically important variation in benefits when trial results are risk stratified, typically showing that an identifiable subgroup of higher-risk patients often account for most of the treatment benefit.<sup>36–46</sup>

Another consistent finding was that the median predicted outcome risk in these trials was lower than the mean predicted risk (i.e. MMRR < 1). Because the summary results of trials reflect the arithmetical mean risk, rather than the median risk, this implies that the typical patient is often at somewhat lower risk—and sometimes at much lower risk—than one might infer from the overall result. When proportional effects are similar across risk groups, summary results may have a tendency to overestimate the degree of benefit on the absolute scale.<sup>5,47</sup> These concerns are especially germane when outcomes rates are predictable and outcome rates relatively low.

Whereas several trials in our database of trials exhibited large heterogeneity in predicted outcome risk, overall the results of our analyses were somewhat less extreme than previous published examples might have suggested.<sup>36–45</sup> There are several explanations for this observation. First, risk heterogeneity may be somewhat restricted in large phase III randomized studies if they tend to enroll homogeneous patient populations. Second, because we wanted to limit the risk of overfitting models to data, we favoured simpler models, which generally had modest discriminatory ability. Finally, previously published examples might be 'cherry-picked' for extreme results and clinical

#### ACCORD ALLHAT-HTN BEST BHAT CAST 0.2 0.5 · 1 Ŧ Ŧ Ŧ Ŧ 1 1.5 p = 0.5380p = 0.5259p = 0.2519p = 0.3909p = 0.62302 CPPT DIG DPP (lifestyle) DPP (metformin) HDFP 0.2 Ŧ HR (+/- SE) I Ŧ 0.5 Ŧ ł Ŧ Ŧ ٠ Ŧ Ŧ Ŧ 1 1.5 p = 0.2946p = 0.2970 p = 0.1549 p = 0.0008 p = 0.73822 SHEP SOLVD (prevention) SOLVD (treatment) Q4 Q1 Q1 Q2 Q3 Q2 Q3 Q4 0.2 0.5 ł Ŧ • Ŧ Ŧ • Ŧ 1 1.5 p = 0.98110.4870 p = 0.0776D 2 Q1 Q2 03 Q4 Q1 Q2 Q3 Q4 Q1 Q2 Q3 Q4 **Risk Quartile** DCCT (prevention) DCCT (treatment) MTOPS (combination) MTOPS (doxazosin) MTOPS (finasteride) p = 0.4739p = 0.8321p = 0.9253p = 0.3382 p = 0.2918 0.08 HR (+/- SE) 0.2 I Ŧ Ŧ Ŧ Ī ł 0.4 Ŧ Ŧ Ŧ ł Ŧ Ŧ 0.6 1 Q2 Q4 Q1 Q3 Q1 Q2 Q3 Q4 Q1 Q3 Q2 Q3 Q4 Q1 Q2 Q4 Q1 Q2 Q3 Q4 **Risk Quartile**

#### A Hazard ratios

Figure 2. A: Hazard or odds ratios across risk quartiles Hazard ratios are shown for all trials except HDFP, which displays odds ratios. Red markers indicate that the treatment arms were switched (intervention was harmful). The scale for hazard ratio axis is different for DCCT and MTOPS.

significance. It is also important to recognize that expressing heterogeneity of risk using a finer grouping of predicted risk (e.g. quintiles or deciles) would yield ratios that are more extreme than the EQRRs reported here.

The observation that indices that describe the distribution of predicted risk are predictable based on the c-statistic, and the overall event rate of each trial, are as telling as the specific examples in our study. The predictability of the risk distribution derives from the fact that the linear predictor from the risk model conforms fairly closely to a normal distribution,<sup>48</sup> yielding distributions of risk that (to a good approximation) conform to log-normal (for risk estimates derived from Cox models) or logistic-normal distributions (for risk estimates derived from logistic regression models). This relationship permits us to anticipate the degree of risk heterogeneity (i.e. EQRR) and the skewness



#### **B** Absolute risk reduction

Figure 2. B: Absolute risk reduction across risk quartiles Red markers indicate that the treatment arms were switched (intervention was harmful). In Figure 2B, the scale for absolute risk reduction is different for DPP, MTOPS, and DCCT.

**Table 4.** Summary of results for 18 positive treatment comparisons (14 trials)

	Median	IQR	Mean	Range
Hazard (or odds) ratio Q1	0.63	0.52-0.87	0.66	0.16-1.10
Hazard (or odds) ratio Q4	0.69	0.44-0.90	0.64	0.27-0.96
Extreme quartile relative hazard ratio (Q4/Q1)	1.02	0.70-1.21	1.05	0.41-1.82
Absolute risk reduction Q1 (%)	4.73	0.83-6.06	4.50	-1.43-18.27
Absolute risk reduction Q4 (%)	9.04	3.25–19.84	12.01	0.77–34.99
Extreme quartile absolute risk reduction difference (Q4-Q1)	5.10	0.33–10.91	7.51	-3.23-28.33

Q, quartile; IQR, inter-quartile range.

(i.e. MMRR) based on knowledge of the outcome rate and the discrimination (c-statistic) of the model—provided that the risk model is well calibrated. For example, using our simple linear regression results, we would anticipate that, when the outcome rate is 10% and the c-statistic is 0.8, the EQRR will be approximately 13 and the MMRR will be approximately 0.6. When risk differs 13-fold between large population subsets, the overall treatment effect estimated for the trial population is not clinically interpretable. When the median risk is 40% lower than the mean risk, it also seems likely that the average effects may not be easily translated even to typical patients in the same trial. Higher c-statistics and lower outcome prevalence would lead to even more skewed distributions, implying greater risk heterogeneity.

Thus, it does not take extreme assumptions to yield risk distributions that would make overall clinical trial results misleading for many patients. The relationship also implies that a risk-stratified approach might be especially important and clinically informative when the outcome is predictable, based on easily available clinical information, and the overall outcome rate is low. This conclusion is consistent with clinical intuition, because when the outcome is rare and predictable by baseline covariates, it is possible to identify very-low-risk patients who are unlikely to benefit from therapy. Analyses of HTE over-predicted risk are also more likely to be useful for risky or costly therapies, when identifying patients who are unlikely to benefit may be of especially high interest.

Despite the fact that only one trial (DPP) showed a 'statistically significant' interaction between the linear predictor of risk and the treatment assignment indicator, we would urge caution in interpreting the ostensible consistency of effects on the multiplicative scale. We note that the true relationship between risk and effect is underdetermined by the data. Indeed, trial results may often be statistically consistent with homogeneous effects on both the additive and the multiplicative scales across risk groups—despite the mathematical incompatibility of these models and the potential clinical importance of the different inferences the models may yield. We believe that consistency of effects across any of these scales is unlikely to represent the 'true' relationship between the risk of the outcome and the effect of a therapy.

Our study has several limitations. We acknowledge that the use of quartiles is arbitrary, and tends to underestimate heterogeneity, compared with using finer strata of predicted risk or assuming a smooth function of predicted risk. We present our data in quartiles to facilitate comparisons across analyses, based on a previously suggested framework.<sup>18</sup> Heterogeneity may be slightly overestimated based on model overfitting or underestimated based on underfitting; more careful model building (e.g. exploring non-linearity and interactions in the risk models) could have given the impression of more extreme risk heterogeneity. We did not explore non-linear relationships between risk and treatment effects, which may have revealed additional HTE. Additionally, we tried to standardize our modelling approach but we used only a single model for each trial. Different models may fit the data equally well, yet results regarding HTE may be sensitive to the specific variables included in the models and whether any of these variables are treatment effect modifiers. Whereas different models may yield different results, the degree to which any particular covariate modifies treatment is typically unknown-and when there is a strong a priori reason to believe that a particular covariate is likely to modify a treatment effect (apart from its influence on risk) then the relationship of the covariate with the treatment effect should also be examined separately. Finally, we used a convenience sample of large trials, which does not represent the full spectrum of clinical conditions or, specifically, those conditions for which risk modelling may be most informative. A risk-modelling approach may be especially informative when treatment can both prevent and cause the primary outcome of interest (presumably via different mechanisms).<sup>5,6,39,49</sup> In such conditions, the risks of therapy may outweigh the benefits in very low-risk patients, and more treatment effect heterogeneity would be anticipated.

Despite these limitations, our results suggest that clinically important differences in effect across predicted risk are likely to be common in trials with statistically significant average treatment effects. A common assumption (of unclear validity) is consistency of treatment effects across risk groups on the proportional scale, but the only way of testing this assumption is to actually perform such riskstratified analyses. Even when analyses fail to reject the null of proportional effects across different risk strata, the results of risk-stratified analyses can demonstrate clinically important risk differences which would otherwise be obscured. Nevertheless, risk-stratified analyses of clinical trials are still rarely planned as part of the initial study design; if reviewers, editors and regulators expected (or required) such analyses to be routinely conducted, the approach would be more widely adopted.<sup>50</sup>

In summary, predicted risk distributions from Cox regression and logistic regression are largely determined based on c-statistic and outcome rates. Clinically significant risk heterogeneity is common even in large 'efficacy' trials—particularly when outcome rates are low and c-statistics are high. The median risk in these trials is generally lower than the average risk. Statistically significant HTE on the relative risk scale is unusual, but clinically significant heterogeneity in absolute effects appears to be common. A risk stratified approach to trial analysis is feasible and may be most clinically informative when an uncommon outcome is predictable by baseline covariates.

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#### Supplementary Data

Supplementary data are available at IJE online.

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