

ORIGINAL ARTICLE

Data-driven methods distort optimal cutoffs and accuracy estimates of depression screening tools: a simulation study using individual participant data

Parash Mani Bhandari, MSc^{a,b}, Brooke Levis, PhD^{a,b,c}, Dipika Neupane, MSc^{a,b},
Scott B. Patten, PhD^d, Ian Shrier, MD^{a,b,e}, Brett D. Thombs, PhD^{a,b,f,g,h,i,j,*},
Andrea Benedetti, PhD^{a,b,f,*}, the Depression Screening Data (DEPRESSD) EPDS Group[#]

^aLady Davis Institute for Medical Research, Jewish General Hospital, Montreal, Quebec, Canada

^bDepartment of Epidemiology, Biostatistics and Occupational Health, McGill University, Montreal, Quebec, Canada

^cCentre for Prognosis Research, School of Medicine, Keele University, Staffordshire, UK

^dDepartment of Community Health Sciences, University of Calgary, Calgary, Alberta, Canada

^eDepartment of Family Medicine, McGill University, Montréal, Québec, Canada

^fDepartment of Medicine, McGill University, Montréal, Québec, Canada

^gDepartment of Psychiatry, McGill University, Montréal, Québec, Canada

^hDepartment of Psychology, McGill University, Montréal, Québec, Canada

ⁱDepartment of Educational and Counselling Psychology, McGill University, Montréal, Québec, Canada

^jBiomedical Ethics Unit, McGill University, Montréal, Québec, Canada

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Abstract

Objective: To evaluate, across multiple sample sizes, the degree that data-driven methods result in (1) optimal cutoffs different from population optimal cutoff and (2) bias in accuracy estimates.

Study design and setting: A total of 1,000 samples of sample size 100, 200, 500 and 1,000 each were randomly drawn to simulate studies of different sample sizes from a database ($n = 13,255$) synthesized to assess Edinburgh Postnatal Depression Scale (EPDS) screening accuracy. Optimal cutoffs were selected by maximizing Youden's J (sensitivity+specificity-1). Optimal cutoffs and accuracy estimates in simulated samples were compared to population values.

Results: Optimal cutoffs in simulated samples ranged from ≥ 5 to ≥ 17 for $n = 100$, ≥ 6 to ≥ 16 for $n = 200$, ≥ 6 to ≥ 14 for $n = 500$, and ≥ 8 to ≥ 13 for $n = 1,000$. Percentage of simulated samples identifying the population optimal cutoff (≥ 11) was 30% for $n = 100$, 35% for $n = 200$, 53% for $n = 500$, and 71% for $n = 1,000$. Mean overestimation of sensitivity and underestimation of specificity were 6.5 percentage point (pp) and -1.3 pp for $n = 100$, 4.2 pp and -1.1 pp for $n = 200$, 1.8 pp and -1.0 pp for $n = 500$, and 1.4 pp and -1.0 pp for $n = 1,000$.

Conclusions: Small accuracy studies may identify inaccurate optimal cutoff and overstate accuracy estimates with data-driven methods. © 2021 Elsevier Inc. All rights reserved.

Keywords: Optimal cutoff; Accuracy estimates; Bias; Cherry-picking; Data-driven methods; Depression

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[#] DEPRESSD EPDS Collaboration listed at end of article.

* Corresponding author. Brett D. Thombs: Tel. (514) 340-8222 ext. 25112; Andrea Benedetti: Tel. (514) 934-1934 ext. 32161.

E-mail addresses: brett.thombs@mcgill.ca (B.D. Thombs), andrea.benedetti@mcgill.ca (A. Benedetti).

What is new?

Key findings

- Optimal EPDS cutoffs identified in samples of different sizes varied widely, ranging from ≥ 5 to ≥ 17 for studies with $n = 100$ and ≥ 8 to ≥ 13 for $n = 1,000$.
- Mean overestimation of sensitivity and underestimation of specificity, respectively, were 6.5 percentage point (pp) and -1.3 pp for $n = 100$ and 1.4 pp and -1.0 pp for $n = 1,000$.

What this adds to what is known?

- This is the first study to use real patient data to estimate the degree that data-driven methods result in selection of inaccurate optimal cutoffs and bias in accuracy estimates.

What is the implication?

- Optimal cutoffs identified in primary accuracy studies are often incorrect and accuracy estimates are often overstated.
- Researchers should avoid making recommendations about cutoffs to use in practice and accuracy when reporting results from small single studies.
- Clinicians should select cutoffs for specific populations that are generated from well-conducted meta-analyses or identified consistently across multiple large primary studies.

1. Introduction

Depression screening tools are commonly used to identify patients with unrecognized and untreated depression [1,2]. Evidence from studies on the accuracy of depression screening tools is used to select an “optimal” cutoff for use in practice and to estimate accuracy for distinguishing between positive and negative results using that cutoff. Many studies on depression screening tools, however, use data-driven approaches, by which investigators use the same dataset to select an optimal cutoff and estimate screening accuracy at that cutoff. Cutoffs selected in this way may deviate substantially from a true population optimal cutoff that would be selected if a sufficiently large or population database were available. Additionally, accuracy estimates generated in these studies may be optimistic compared to what would occur in clinical practice.

The Edinburgh Postnatal Depression Scale (EPDS) is the most commonly used tool to screen for depression during pregnancy and postpartum [3,4]. Diagnosis of depression in pregnancy and postpartum is particularly challenging, since some symptoms overlap with normal experiences during this period, such as loss of appetite, poor sleep and fatigue [5-7]. Some health care practitioners may not fully

understand the information provided by EPDS; different resources suggest that cutoffs of ≥ 10 and ≥ 13 can be used to identify women with “possible” or “probable” depression [8,9], but only approximately 35% and 60% of women who score above cutoffs of ≥ 10 and ≥ 13 , respectively, will experience a major depressive episode, assuming a prevalence of 10% [10,11].

Misunderstanding about how to interpret results from depression screening tools is compounded by results from primary studies that suggest that different cutoffs identified as “optimal” in their studies should be used in specific populations. These results are often generated using data-driven analytical approaches in small samples. In many of these studies, the abstract, which may be the only part of the article that is read [12,13], only reports accuracy results from a single data-driven optimal cutoff rather than from standard cutoffs. Data-driven methods that are sometimes used for optimal cutoff selection include selecting the cutoff that maximizes Youden’s J (sensitivity + specificity – 1), minimizes Euclidean distance (distance to the corner of the receiver operator characteristic curve) or maximizes diagnostic odds ratio [14]. Youden’s J is the method most frequently used in diagnostic accuracy studies for depression screening tools. Illustrating this, we reviewed recently published primary studies of EPDS accuracy (N participants, range = 118-807; mean = 320) and found that only 1 of 14 studies (7%) reported accuracy results for more than one cutoff in the abstract. The remaining 13 (93%) only reported results from the single best-performing cutoff, which was based on maximizing Youden’s J in 11 of the 13 (85%) studies. Cutoffs identified as optimal in the 14 studies ranged from ≥ 8 to ≥ 13 . In many of the studies, when data-driven optimal cutoffs diverged from more standard cutoffs, authors suggested that this represented a unique optimal cutoff that should be used in the study’s specific target population group. No studies attributed a divergent optimal cutoff to a small sample size or to data-driven cutoff selection methods (Appendix-eMethods1).

We know of only four studies that have investigated the degree to which data-driven selection of cutoff may influence diagnostic accuracy estimates [15-18]. These studies each reported that data-driven cutoff selection produces overly optimistic estimates, particularly in small samples. However, these studies used simulated datasets based on hypothetical test score distributions rather than real participant data. Thus, how widely data-driven optimal cutoffs diverge from population-based optimal cutoffs and how biased estimates of diagnostic accuracy may be based on actual participant data is not known for any depression screening test, including the EPDS.

The objectives of the present study were to illustrate for users of evidence on depression screening tool accuracy, such as the EPDS, across different study sample sizes, the degree to which study-level data-driven cutoff selection: (1) results in the selection of optimal cutoffs that differ from the population optimal cutoff derived from a

“population” dataset and (2) generates biased accuracy results compared to results from the population optimal cutoff.

2. Methods

We used a database originally synthesized for an individual participant data meta-analysis (IPDMA) on the accuracy of the EPDS for depression screening to form a study population from which we simulated studies of different sample sizes [10]. A protocol for the present study was uploaded to the Open Science Framework repository prior to initiating the study (<https://osf.io/qnvzp/>).

Details on methodology for the original IPDMA used in this study are published elsewhere [10], and are provided in Appendix-eMethods2.

2.1. Simulation of study samples and statistical analyses

Unlike many other depression screening tools, EPDS does not have a clearly recognized standard cutoff for depression screening. The original validation study which included 84 participants and 24 cases of definite or probable major depression based on Research Diagnostic Criteria suggested that cutoffs of ≥ 10 or ≥ 13 could be used [6]. However, many studies report using different cutoffs between ≥ 10 and ≥ 13 to identify major depression [19,20], with ≥ 13 being the most common [20]. A recent IPDMA using an updated and slightly larger version of the dataset used in the present study found that a cutoff of ≥ 11 maximized Youden’s J overall and for subgroups.

For the present study, we used our IPDMA dataset to represent a hypothetical “population” of women, and defined population sensitivity and specificity values for EPDS cutoffs to be those estimated in this population. To do this, we analyzed the IPDMA dataset, ignoring sampling weights as well as study-level clustering of observations. We ignored sampling weights and clustering to have a defined population from which we could draw samples that represented simulated primary studies and to be able to use the same analytical approach when analyzing the population data and the simulated primary study data. As a result, we generated accuracy estimates that differed slightly from those reported in the full IPDMA, which used sampling weights and study-level clustering and a slightly larger sample. We verified that a cutoff of ≥ 11 maximized Youden’s J for the unweighted population.

From the population IPDMA dataset, we sampled with replacement to generate 1,000 random samples of sample size 100, 200, 500, 1,000 each. For each sample, we defined the sample-specific optimal cutoff as the cutoff that maximized Youden’s J in the sample. If there was a tie in maximum Youden’s J between multiple cutoffs, we selected the higher cutoff. For each sample size, across the 1,000 samples, we (1) graphically illustrated the variability

in sample-specific optimal cutoffs and the variability in accuracy of the sample-specific optimal cutoffs; (2) estimated the mean difference (bias) and associated 95% confidence interval (CI) between sensitivity and specificity based on sample-specific optimal cutoffs versus the population sensitivity and specificity based on the population optimal cutoff of ≥ 11 , and (3) estimated the mean difference (bias) and 95% CI for sensitivity and specificity based on a cutoff of ≥ 11 in each sample versus the population sensitivity and specificity also based on a cutoff of ≥ 11 . CIs for the variability in optimal cutoffs and the unweighted accuracy estimates were computed using a one sample proportion test with continuity correction. For all analyses, sensitivity and specificity were estimated using crude 2×2 table counts. In additional analyses, we stratified results by the optimal cutoff value identified in each sample.

2.2. Deviations from protocol

We initially specified that we would also compare accuracy of the optimal cutoff in each sample with that of cutoff ≥ 13 , which is the cutoff most commonly used in practice [19,20]. We subsequently determined that a population optimal cutoff of ≥ 11 maximizes Youden’s J in our IPDMA “population”, which was established in the main IPDMA database and confirmed in the present study. Thus, we used a cutoff of ≥ 11 only and not ≥ 13 , since the purpose was to determine how data-driven results would diverge from similar analyses done with population data.

3. Results

The original IPDMA database included 49 primary studies with 13,255 participants (1,625 major depression cases, 12.3%), which constituted the “population” for the present study. Characteristics of the primary studies included in the IPDMA database are provided in Appendix-eTable1. The sample sizes of the primary studies ranged from 40 to 2,634 (mean = 271, median = 190). The mean number of cases of major depression was 34 (median = 25), and 20 studies included < 20 cases of major depression. Frequencies of EPDS scores for cases and non-cases in the IPDMA database are shown in Appendix-eTable2. As shown in Appendix-eFigure1, study-specific optimal cutoffs that maximized Youden’s J ranged from ≥ 5 to ≥ 19 . For the “population” of 13,255 participants and using a cutoff of ≥ 11 , the unweighted sensitivity and specificity were 78.7% (95% CI: 76.6, 80.7) and 83.4% (95% CI: 82.7, 84.0).

3.1. Variability of sample-specific optimal cutoffs in simulated samples

Figure 1 shows the variability of sample-specific optimal cutoffs for each sample size. Optimal cutoffs in indi-

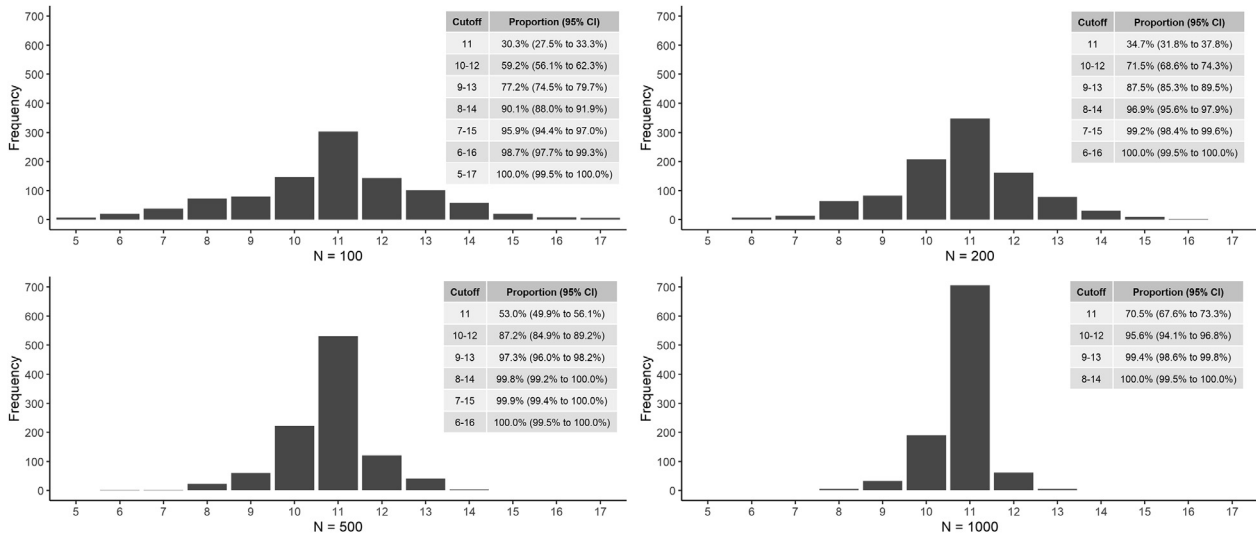


Fig. 1. Variability in optimal cutoffs in 1,000 simulated samples of sample size 100, 200, 500 and 1,000. Optimal cutoff was defined as the cutoff that maximized Youden’s J (sensitivity + specificity – 1) in the study sample. 26 out of the 4,000 simulated samples had a tie in maximum Youden’s J, and the higher cutoff was selected as the optimal cutoff.

Table 1. Bias in accuracy estimates (in percentage point) in 1,000 simulated samples of sample size 100, 200, 500 and 1,000

	Mean Difference (95% CI)							
	Sensitivity	Specificity	Sensitivity	Specificity	Sensitivity	Specificity	Sensitivity	Specificity
	Sample size = 100		Sample size = 200		Sample size = 500		Sample size = 1,000	
Sample-specific optimal cutoff values – Population values with cutoff ≥ 11	6.5 (5.8, 7.2)	-1.3 (-1.9, -0.7)	4.2 (3.6, 4.7)	-1.1 (-1.6, -0.7)	1.8 (1.4, 2.1)	-1.0 (-1.3, -0.7)	1.4 (1.1, 1.6)	-1.0 (-1.2, -0.8)
Sample cutoff ≥ 11 values – Population values with cutoff ≥ 11	0.0 (-0.7, 0.8)	0.1 (-0.2, 0.3)	0.2 (-0.3, 0.8)	0.1 (-0.1, 0.3)	-0.2 (-0.6, 0.1)	0.0 (-0.1, 0.1)	0.1 (-0.1, 0.3)	-0.1 (-0.1, 0.0)

Optimal cutoff refers to the cutoff that maximized Youden’s J (sensitivity + specificity – 1) in the sample. Sample values are estimated from the simulated samples. Population values are estimated from the full dataset.

vidual samples ranged from ≥ 5 to ≥ 17 for $n = 100$, ≥ 6 to ≥ 16 for $n = 200$, ≥ 6 to ≥ 14 for $n = 500$, and ≥ 8 to ≥ 13 for $n = 1,000$. There was a tie in maximum Youden’s J between multiple cutoffs in 26 of the 4,000 samples. The percentage of samples that identified the true population optimal cutoff of ≥ 11 was 30.3% (95% CI: 27.5, 33.3) for $n = 100$, 34.7% (95% CI: 31.8, 37.8) for $n = 200$, 53.0% (95% CI: 49.9, 56.1) for $n = 500$, and 70.5% (95% CI: 67.6, 73.3) for $n = 1,000$.

3.2. Bias from data-driven cutoff selection in simulated samples

As shown in Table 1, based on the overall mean across 1,000 samples, sensitivity based on sample-specific optimal cutoffs was overestimated compared to the sensitivity in the population based on the population optimal cutoff by

6.5 percentage point (pp) (95% CI: 5.8, 7.2) for $n = 100$ [i.e. mean sensitivity of optimal cutoffs in 1,000 simulated samples of $n = 100$ (85.2%) – true population sensitivity (78.7%) = 6.5 pp], 4.2pp (95% CI: 3.6, 4.7) for $n = 200$, 1.8 pp (95% CI: 1.4, 2.1) for $n = 500$ and 1.4 pp (95% CI: 1.1, 1.6) for $n = 1,000$. Specificity was underestimated by 1.3 pp (95% CI: -1.9, -0.7) for $n = 100$, 1.1 pp (95% CI: -1.6, -0.7) for $n = 200$, 1.0 pp (95% CI: -1.3, -0.7) for $n = 500$ and 1.0 pp (95% CI: -1.2, -0.8) for $n = 1,000$. Figure 2 presents quartiles of the accuracy estimates for simulated samples.

Figure 3 and Appendix-eTable3 show that the direction and magnitude of bias in sensitivity and specificity estimates depended on the optimal cutoff identified in each sample. For instance, with $n = 100$, in samples with sample-specific optimal cutoff ≥ 5 to ≥ 8 , sensitivity was overestimated by 16.0 pp (95% CI: 14.8, 17.2), and specificity was underestimated by 19.6 pp (95% CI: -20.8, -

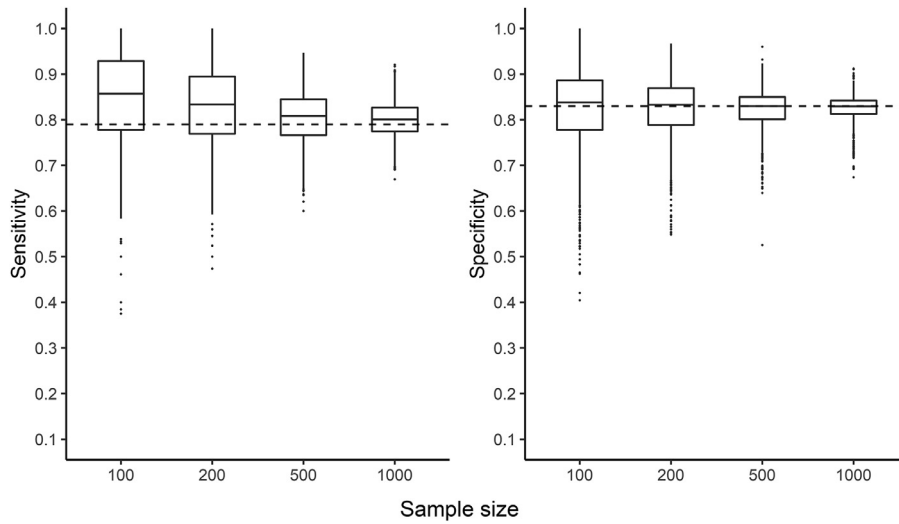


Fig. 2. Boxplots of accuracy estimates of the optimal cutoff in 1,000 simulated samples of sample size 100, 200, 500 and 1,000 compared to the accuracy estimates of cutoff ≥ 11 in the population. Optimal cutoff refers to the cutoff that maximized Youden’s J (sensitivity + specificity – 1) in the study sample. Dotted horizontal line represents the accuracy of cutoff ≥ 11 in the population (full IPDMA dataset). Boxplots present quartiles (first quartile, median and third quartile) of accuracy estimates for the simulated samples.

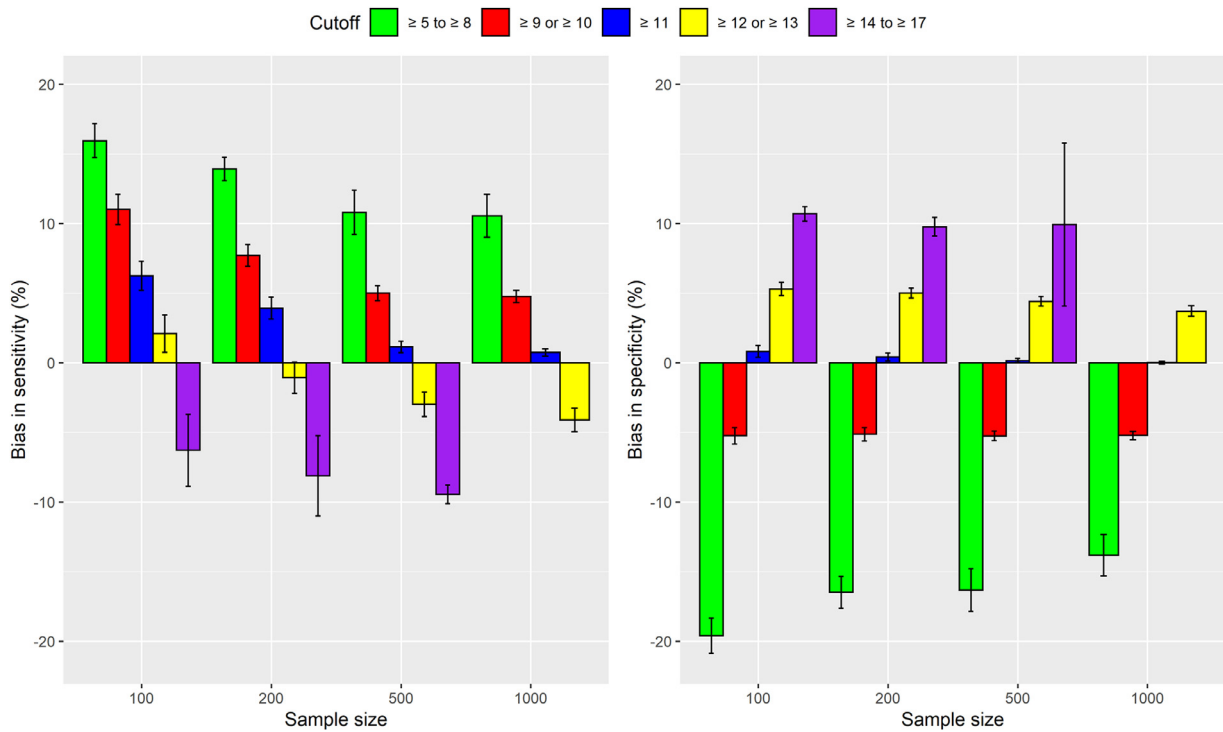


Fig. 3. Bias in accuracy estimates in simulated samples of sample size 100, 200, 500 and 1,000 stratified by sample optimal cutoffs. Optimal cutoff refers to the cutoff that maximized Youden’s J (sensitivity + specificity – 1) in the study sample. The error bars represent 95% confidence intervals of the bias in accuracy estimates.

18.3). For samples with sample-specific optimal cutoffs of ≥ 14 to ≥ 17 , sensitivity was underestimated by 6.3 pp (95% CI: -8.9, -3.7), and specificity was overestimated by 10.7 pp (95% CI: 10.2, 11.2).

As shown in [Figure 4](#), when sensitivity and specificity were calculated for cutoff ≥ 11 in each sample, the mean sensitivity and specificity were close to that of the population values. See also [Table 1](#).

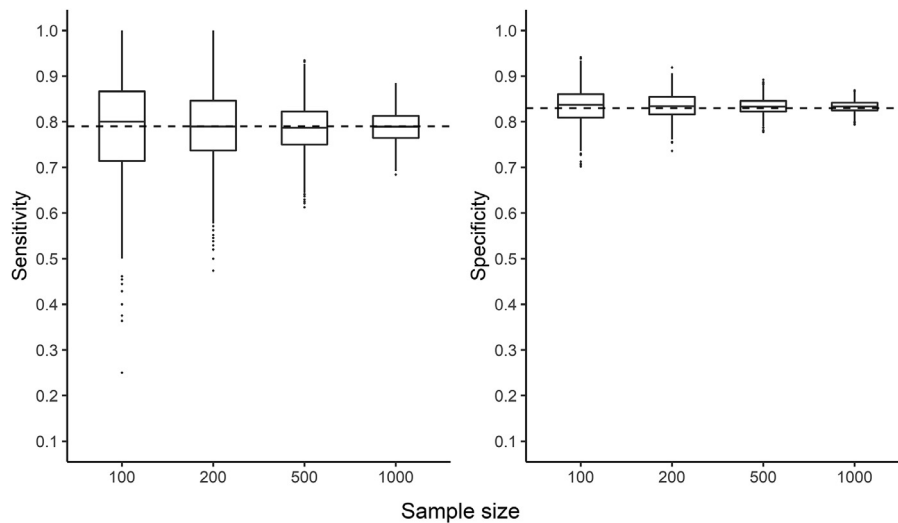


Fig. 4. Boxplots of accuracy estimates of the cutoff ≥ 11 in 1,000 simulated samples of sample size 100, 200, 500 and 1,000 compared to the accuracy estimates of cutoff ≥ 11 in the population.

Optimal cutoff refers to the cutoff that maximized Youden's J (sensitivity + specificity – 1) in the sample. Dotted horizontal line represents the accuracy of cutoff ≥ 11 in the population (full IPDMA dataset). Boxplots present quartiles (first quartile, median and third quartile) of accuracy estimates for the simulated samples.

4. Discussion

There were two main findings of this study. First, with very small sample sizes ($n = 100$), study-specific optimal cutoffs ranged from ≥ 5 to ≥ 17 (compared to the actual population optimal cutoff of ≥ 11). Even with samples of $n = 1,000$, optimal cutoffs ranged from ≥ 8 to ≥ 13 . Second, for samples of $n = 100$, mean overestimation of sensitivity was 6.5 pp, whereas mean underestimation of specificity was 1.3 pp. For larger samples ($n = 1,000$), sensitivity was overestimated, on average, by 1.4 pp and specificity underestimated by 1.0 pp. The degree and direction of bias from population-level estimates depended on the identified sample-specific optimal cutoff. For $n = 100$, for example, individual studies that identified optimal cutoffs from ≥ 5 to ≥ 8 overestimated sensitivity by an average of 16.0pp; studies that identified high optimal cutoffs (≥ 14 to ≥ 17), on the other hand, underestimated sensitivity by 6.3 pp.

The degree of variability identified in sample-specific optimal cutoffs, especially with smaller sample sizes, is concerning, because most diagnostic accuracy studies of depression screening tools are conducted in small samples. Among the 49 studies included in the present IPDMA database, 26 (53.1%) had sample size of < 200 , 19 (38.8%) had sample size of 200 to 500, 3 (6.1%) had sample size of 501 to 1,000 and only one (2.0%) had sample size $> 1,000$. A previous study examined sample sizes and the presence of sample size calculations in 89 studies of depression screening tool accuracy, not limited to the EPDS, and found that the median sample size was 224; 38 (42.7%) had sample size of < 200 , 33 (37.1%) had sam-

ple size of 200 to 500, 11 (12.3%) had sample size of 501 to 1,000 and 7 (7.9%) had sample size of $> 1,000$ [21]. Based on our findings, overall, many studies of depression screening tool accuracy likely overestimate sensitivity with only minor losses in specificity. A larger bias in sensitivity estimates as compared to bias in specificity estimates is intuitive, as most studies have much fewer participants with major depression (among whom sensitivity is estimated) than without (among whom specificity is estimated). Thus, optimal cutoff selection in some samples can result in substantial gains in sensitivity with relatively small compensation in specificity, particularly in small samples. As shown in the present study, however, mean differences do not capture what may occur in any given study, and depending on the specific sample, sensitivity may be overestimated or underestimated, sometimes substantially.

Surveys have shown that clinicians have difficulty understanding medical statistics, including conditional probabilities such as sensitivity, specificity, positive predictive value and negative predictive value [22–24]. Thus, clinicians may misinterpret EPDS cutoffs with inflated sensitivity estimates from data-driven procedures as being virtually diagnostic, and adopt such cutoffs for use in clinical practice, even when the actual positive predictive value may be much smaller [25].

Clinicians who use the EPDS in their practice should be wary of EPDS cutoff recommendations based on small individual studies that used data-driven methods to identify the optimal cutoff. Such cutoffs are likely to not truly be optimal for the population of interest, and accuracy estimates are likely to be overly optimistic compared to what would be obtained in actual clinical practice. Instead, clin-

icians should select EPDS cutoff thresholds from large, well-conducted meta-analyses or validated across multiple studies. In addition, clinicians may also opt to prioritize either sensitivity or specificity in different clinical settings, and select higher or lower thresholds, depending on their health and financial priorities.

The Standards for Reporting of Diagnostic Accuracy Studies (STARD) reporting guideline recommends *a priori* sample size estimation for the desired precision level in accuracy estimates [26]. Results from our study show that setting sample size targets pre-study should also consider variability in the optimal cutoff that may be identified and not just variability in accuracy estimates. A previous study that examined sample sizes in 89 studies of depression screening tool accuracy found that only three reported *a priori* sample size calculations, and none specifically considered the issue of identifying an optimal cutoff and estimating accuracy in the same participant sample [21]. Authors of primary studies on depression screening tool accuracy could potentially use statistical methods to estimate confidence intervals for uncertainty around the optimal cutoff [27,28]. They could also employ internal validation methods such as cross-sampling, sample-splitting and bootstrapping to statistically adjust for the bias in accuracy estimates from data-driven optimal cutoff selection [29]. These methods, however, have not been demonstrated or tested in the context of mental health screening. Indeed, the most robust approach for identifying optimal cutoffs and generating accurate estimates of screening or diagnostic accuracy is through pooling large numbers of well-conducted primary studies and participants via meta-analysis, preferably IPDMA, which can ensure that all cutoffs are available for examination for all participants [30,31]. Researchers should report accuracy data from primary accuracy studies for all possible cutoffs in 2×2 form, at least in appendices, to facilitate subsequent synthesis and to avoid selective cutoff reporting bias [32].

As shown in our review of recent studies of the EPDS (Appendix-eMethods1), authors of diagnostic accuracy studies that identify study-specific optimal cutoffs that depart from standard cutoffs often conclude that this is evidence for the need to use different cutoffs in different populations. While this is possible, our full IPDMA of the screening accuracy of the EPDS did not find evidence for differential accuracy by subgroups. Since the same method was used in this study to identify the population optimal cutoff and the optimal cutoff in each simulated study sample, the reason why study sample optimal cutoffs diverge, particularly in very small samples is due to the sampling distribution of the mean in relatively small samples. Hence, authors of individual studies should avoid recommending specific cutoffs for specific populations unless the studies use very large samples, or the findings are replicated consistently across multiple studies.

Strengths of this study include the use of real-participant data instead of simulated data and hypothetical distribu-

tional assumptions and the large population from which we were able to draw samples. There are also limitations to consider. Our results on the bias in accuracy estimates is based on the analysis of a large dataset on depression screening accuracy of the EPDS, and the results may be different for a different test or study sample. Another possible limitation is that we only used Youden's J to select optimal cutoffs. It is the most commonly used method, by far, in depression screening accuracy studies and performs similarly to other indices, such as the Euclidean distance [14]. It is possible, however, that results might slightly differ for an alternative method for selecting optimal cutoffs.

5. Conclusions

We found that data-driven cutoff selection methods often result in optimal cutoffs that differ from the population optimal cutoff and in accuracy estimates that are overly optimistic. Researchers who conduct primary studies of diagnostic accuracy should calculate sample sizes *a priori* and describe related limitations; they should avoid recommending cutoffs for population subgroups when sample sizes are not sufficiently large; and should report results completely. Clinicians should be aware that cutoffs and accuracy from single studies may not reflect what will occur in practice and should select a cutoff from well-conducted meta-analyses or identified consistently across multiple studies.

The Depression Screening Data (DEPRESSD) EPDS Group

Ying Sun, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Chen He, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Danielle B. Rice, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Ankur Krishnan, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Yin Wu, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Marleine Azar, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Tatiana A. Sanchez, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Matthew J. Chiovitti, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Nazanin Saadat, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Kira E. Riehm, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Mahrukh Imran, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Zelalem Negeri, Lady Davis Institute for Medical Research, Jewish General Hospital, Montréal, Québec, Canada; Jill T. Boruff, Schulich Library of Physical Sciences, Life Sciences, and Engineer-

ing, McGill University, Montréal, Québec, Canada; Pim Cuijpers, Department of Clinical, Neuro and Developmental Psychology, Amsterdam Public Health Research Institute, Vrije Universiteit, Amsterdam, the Netherlands; Simon Gilbody, Hull York Medical School and the Department of Health Sciences, University of York, Heslington, York, UK; John P.A. Ioannidis, Department of Medicine, Department of Health Research and Policy, Department of Biomedical Data Science, Department of Statistics, Stanford University, Stanford, California, USA; Lorie A. Kloda, Library, Concordia University, Montréal, Québec, Canada; Roy C. Ziegelstein, Department of Medicine, Johns Hopkins University School of Medicine, Baltimore, Maryland, USA; Liane Comeau, International Union for Health Promotion and Health Education, École de santé publique de l'Université de Montréal, Montréal, Québec, Canada; Nicholas D. Mitchell, Department of Psychiatry, University of Alberta, Edmonton, Alberta, Canada; Marcello Tonelli, Department of Medicine, University of Calgary, Calgary, Alberta, Canada; Simone N. Vigod, Women's College Hospital and Research Institute, University of Toronto, Toronto, Ontario, Canada; Franca Aceti, Department of Neurology and Psychiatry, Sapienza University of Rome, Rome, Italy; Rubén Alvarado, School of Public Health, Faculty of Medicine, Universidad de Chile, Santiago, Chile; Cosme Alvarado-Esquivel, Laboratorio de Investigación Biomédica, Facultad de Medicina y Nutrición, Avenida Universidad, Dgo, Mexico; Muideen O. Bakare, Child and Adolescent Unit, Federal Neuropsychiatric Hospital, Enugu, Nigeria; Jacqueline Barnes, Department of Psychological Sciences, Birkbeck, University of London, UK; Amar D. Bavle, Department of Psychiatry, Rajarajeswari Medical College and Hospital, Bengaluru, Karnataka, India; Cheryl Tatano Beck, University of Connecticut School of Nursing, Mansfield, Connecticut, USA; Carola Bindt, Department of Child and Adolescent Psychiatry, University Medical Center Hamburg-Eppendorf, Germany; Philip M. Boyce, Discipline of Psychiatry, Westmead Clinical School, Sydney Medical School, University of Sydney, Sydney, Australia; Adomas Bunevicius, Neuroscience Institute, Lithuanian University of Health Sciences, Kaunas, Lithuania; Tiago Castro e Couto, Federal University of Uberlândia, Brazil; Linda H. Chaudron, University of Rochester School of Medicine and Dentistry, Rochester, New York, USA; Humberto Correa, Medicine Faculty - Universidade Federal de Minas Gerais. Belo Horizonte, MG, Brazil; Felipe Pinheiro de Figueiredo, Department of Neurosciences and Behavior, Ribeirão Preto Medical School, Brazil; Valsamma Eapen, University of New South Wales and Ingham Institute South West Sydney LHD, Australia; Nicolas Favez, Faculty of Psychology and Educational Sciences, University of Geneva, Geneva, Switzerland; Ethel Felice, Department of Psychiatry, Mount Carmel Hospital, Attard, Malta; Michelle Fernandes, Faculty of Medicine, Department of Paediatrics, University of Southampton, Southampton and Nuffield De-

partment of Women's & Reproductive Health, University of Oxford, Oxford, UK; Barbara Figueiredo, School of Psychology, University of Minho, Portugal; Jane R. W. Fisher, School of Public Health and Preventive Medicine, Monash University, Melbourne, Australia; Lluïsa Garcia-Esteve, Perinatal Mental Health Unit CLINIC-BCN, Institut Clínic de Neurociències, Hospital Clínic, Barcelona, Spain; Lisa Giardinelli, Psychiatry Unit, Department of Health Sciences, University of Florence, Firenze, Italy; Nadine Helle, Department of Child and Adolescent Psychiatry, University Medical Center Hamburg-Eppendorf, Germany; Louise M. Howard, Institute of Psychiatry, Psychology & Neuroscience, King's College London, London, UK; Dina Sami Khalifa, Ahfad University for Women, Omdurman, Sudan; Jane Kohlhoff, University of New South Wales, Kensington, Australia; Zoltán Kozinszky, Department of Obstetrics and Gynaecology, Danderyd Hospital, Stockholm, Sweden; Laima Kusminskas, Private Practice, Hamburg, Germany; Lorenzo Lelli, Psychiatry Unit, Department of Health Sciences, University of Florence, Firenze, Italy; Angeliki A. Leonardou, First Department of Psychiatry, Women's Mental Health Clinic, Athens University Medical School, Athens, Greece; Michael Maes, Department of Psychiatry, Faculty of Medicine, Chulalongkorn University, Bangkok, Thailand; Valentina Meuti, Department of Neurology and Psychiatry, Sapienza University of Rome, Rome, Italy; Sandra Nakić Radoš, Department of Psychology, Catholic University of Croatia, Zagreb, Croatia; Purificación Navarro García, Perinatal Mental Health Unit CLINIC-BCN. Institut Clínic de Neurociències, Hospital Clínic, Barcelona, Spain; Daisuke Nishi, Department of Mental Health, Graduate School of Medicine, The University of Tokyo, Japan; Daniel Okitundu Luwa E-Andjafono, Unité de Neuropsychologie, Département de Neurologie, Centre Neuro-psycho-pathologique, Faculté de Médecine, Université de Kinshasa, République Démocratique du Congo; Susan J. Pawlby, Institute of Psychiatry, Psychology & Neuroscience, King's College London, London, UK; Chantal Quispel, Department of Obstetrics and Gynaecology, Albert Schweitzer Ziekenhuis, Dordrecht, the Netherlands; Emma Robertson-Blackmore, Halifax Health, Graduate Medical Education, Daytona Beach, FL. USA; Tamsen J. Roach, MRC/Developmental Pathways to Health Research Unit, School of Clinical Medicine, University of Witwatersrand, South Africa; Heather J. Rowe, School of Public Health and Preventive Medicine, Monash University, Melbourne, Australia; Deborah J. Sharp, Centre for Academic Primary Care, Bristol Medical School, University of Bristol, UK; Bonnie W. M. Siu, Department of Psychiatry, Castle Peak Hospital, Hong Kong SAR, China; Alkistis Skalkidou, Department of Women's and Children's Health, Uppsala University, Uppsala, Sweden; Alan Stein, University of Oxford, Oxford, UK; Robert C. Stewart, Department of Mental Health, College of Medicine, University of Malawi, Malawi; Kuan-Pin Su, College of Medicine, China Medical

University, Taichung, Taiwan; Inger Sundström-Poromaa, Department of Women's and Children's Health, Uppsala University, Uppsala, Sweden; Meri Tadinac, Department of Psychology, Faculty of Humanities and Social Sciences, University of Zagreb, Croatia; S. Darius Tandon, Northwestern University Feinberg School of Medicine, Chicago, IL, USA; Iva Tendais, School of Psychology, University of Minho, Portugal; Pavaani Thiagayson, Institute of Mental Health, Singapore; Annamária Tőreki, Department of Emergency, University of Szeged, Hungary; Anna Torres-Giménez, Perinatal Mental Health Unit CLINIC-BCN. Institut Clínic de Neurociències, Hospital Clínic, Barcelona, Spain; Thach D. Tran, School of Public Health and Preventive Medicine, Monash University, Melbourne, Australia; Kylee Trevillion, Institute of Psychiatry, Psychology & Neuroscience, King's College London, London, UK; Katherine Turner, Epilepsy Center-Child Neuropsychiatry Unit, ASST Santi Paolo Carlo, San Paolo Hospital, Milan, Italy; Johann M. Vega-Dienstmaier, Facultad de Medicina Alberto Hurtado, Universidad Peruana Cayetano Heredia, Lima, Perú; Karen Wynter, School of Nursing & Midwifery, Deakin University, Melbourne, Australia; and Kimberly A. Yonkers, Department of Psychiatry, Yale School of Medicine, New Haven, Connecticut, USA.

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Author Contributions

PMBhandari, BL, DNeupane, JTB, PC, SG, JPAI, LAK, SBP, IS, RCZ, LC, NDM, MTonelli, SNV, BDT and ABenedetti were responsible for the study conception and design. JTB and LAK designed and conducted database searches to identify eligible studies. FA, RA, CAE, MOB, JB, ADB, CTB, CB, PMBoyce, ABunevicius, TCeC, LHC, HC, FPF, VE, NF, EF, MF, BF, JRWF, LGE, LG, NH, LMH, DSK, JK, ZK, LK, LL, AAL, MM, VM, SNR, PNG, DNishi, DOLEA, SJP, CQ, ERB, TJR, HJR, DJS, BWMS, ASkalkidou, AStein, RCS, KPS, ISP, MTadinac, SDT, IT, PT, AT, ATG, TDT, KTrevillion, KTurner, JMVD,

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Ethics Statement

As this study involved only analysis of previously collected de-identified data and because all included studies were required to have obtained ethics approval and informed consent, the Research Ethics Committee of the Jewish General Hospital determined that ethics approval was not required.

Data Sharing

Requests to access data should be made to the corresponding authors.

Supplementary material

Supplementary material associated with this article can be found, in the online version, at doi:[10.1016/j.jclinepi.2021.03.031](https://doi.org/10.1016/j.jclinepi.2021.03.031).

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