Case Complexity as a Guide for Psychological Treatment Selection

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Objective: Some cases are thought to be more complex and difficult to treat, although there is little consensus on how to define complexity in psychological care. This study proposes an actuarial, data-driven method of identifying complex cases based on their individual characteristics.

Method: Clinical records for 1,512 patients accessing low- and high-intensity psychological treatments were partitioned in 2 random subsamples. Prognostic indices predicting post-treatment reliable and clinically significant improvement (RCSI) in depression (Patient Health Questionnaire-9; Kroenke, Spitzer, & Williams, 2001) and anxiety (Generalized Anxiety Disorder-7; Spitzer, Kroenke, Williams, & Löwe, 2006) symptoms were estimated in 1 subsample using penalized (Lasso) regressions with optimal scaling. A PI-based algorithm was used to classify patients as standard (St) or complex (Cx) cases in the second (cross-validation) subsample. RCSI rates were compared between Cx cases that accessed treatments of different intensities using logistic regression. Results: St cases had significantly higher RCSI rates compared to Cx cases (OR = 1.81 to 2.81). Cx cases tended to attain better depression outcomes if they were initially assigned to high-intensity (vs. low intensity) interventions (OR = 2.23); a similar pattern was observed for anxiety but the odds ratio (1.74) was not statistically significant. Conclusions: Complex cases could be detected early and matched to high-intensity interventions to improve outcomes.

What is the public health significance of this article?
Complex cases tend to have a poor prognosis after psychological treatment for depression and anxiety problems. An evidence-based model of defining complexity is proposed to guide therapists in matching patients to treatments of differing intensity. The findings indicate that this personalized method of treatment selection could lead to better outcomes for complex cases and could improve upon decisions that are informed by clinical judgment alone.

Keywords: stratified medicine, mental health, case complexity, psychotherapy
interventions offered in a stepped care model had comparable outcomes to patients whose treatments were selected and informed by clinical judgment (Van Straten, Tiemens, Hakkaart, Nolen, & Donker, 2006). Such evidence calls into question clinicians’ ability to match patients to treatments and supports current guidelines to apply a stepped care approach (National Institute for Health and Care Excellence [NICE], 2011). Overall, three key problems are apparent: a lack of conceptual clarity about complex cases, a gap between clinical judgment and research evidence, and limitations in clinicians’ ability to identify and select optimal treatments for complex cases.

Concerns regarding complexity are not exclusive to the practice of psychotherapy. The simultaneous growth and ageing of the general population have confronted many other areas of health care with the challenges of treating patients who present with multiple chronic conditions (Smith & O’Dowd, 2007), leading some to question the usefulness of evidence-based guidelines that are formulated for “prototypical” patients (Boyd et al., 2005; Tinetti, Bogardus, & Agostini, 2004). Consequently, theoretical models to account for case complexity in medicine have been proposed in the last decade. Some of these models conceptualize complexity as arising from a combination of clinical (e.g., diagnostic), biological, socioeconomic, cultural, environmental, and behavioral factors that are statistically associated with clinical prognosis (Safford, Allison, & Kiefe, 2007; Schaink et al., 2012). Individual patients may have protective or risk factors across these domains, and their overall complexity level results from the sum of risks. In an attempt to move beyond mere description, Shippee, Shah, May, Mair, and Montori (2012) proposed a cumulative complexity model that attempts to explain how risk factors accumulate and interact to influence health care outcomes. They proposed that (clinical, socioeconomic, cultural) risk factors complicate health care outcomes by disrupting the balance between patient workload (i.e., number and difficulty of daily life demands including self-care) and patient capacity (i.e., resources and limitations affecting ability to meet demands). From this perspective, effective health care for complex cases would not only require intensive treatment of acute symptoms and specific disease mechanisms, but also attending to wider biopsychosocial aspects that may redress the balance between demands and capacity. Common to these models are the focus on empirically supported prognostic factors, the consideration of factors across multiple domains, and the conceptual understanding of case complexity as resulting from the accumulation of risks and challenges to self-management.

Informed by these theoretical models emerging from the biomedical sciences, this study investigated the impact of case complexity in routine psychological care. Considering the problems outlined above, we sought to assess the merits of an actuarial, data-driven, cumulative model of defining case complexity. Specific objectives were (a) to identify prognostic variables associated with psychological treatment outcomes, (b) to develop an algorithm that could aid clinicians in identifying complex cases at risk of poor outcomes, (c) to determine whether or not complex cases respond differentially to treatments of differing levels of intensity, (d) to ascertain the extent to which patients are adequately matched to available stepped care interventions.

### Method

#### Primary outcome measures.

Patients accessing IAPT services self-complete standardized outcome measures on a session-to-session basis to monitor response to treatment. The Patient Health Questionnaire-9 (PHQ-9; Kroenke et al., 2001) is a nine-item screening tool for major depression, where each item is rated on a 0 to 3 Likert scale, yielding a total depression severity score between 0 and 27. A cut-off ≥10 has been recommended to detect clinically significant depression symptoms (Kroenke et al., 2001), and a difference of ≥6 points between assessments is indicative of reliable change (Richards & Borglin, 2011).

The Generalized Anxiety Disorder-7 (GAD-7; Spitzer, Kroenke, Williams, & Löwe, 2006) is a seven-item measure developed to screen for anxiety disorders. It is also rated using Likert scales, yielding a total anxiety severity score between 0 and 21. A cut-off score ≥8 is recommended to identify the likely presence of a diagnosable anxiety disorder (Kroenke, Spitzer, Williams, Monahan, & Löwe, 2007), and a difference of ≥5 points is indicative of reliable change (Richards & Borglin, 2011). Pre-treatment and last observed PHQ-9 and GAD-7 scores were available for analysis.

#### Other measures.

The Work and Social Adjustment Scale (WSAS; Mundt, Marks, Shear, & Greist, 2002) is a measure of functioning across five domains: work, home management, social leisure activities, private leisure activities, family and close relationships. Each item is rated on a scale of 0 (no impairment) to 8 (very severe impairment), rendering a total functional impairment score between 0 and 40.
The Standardized Assessment of Personality–Abbreviated Scale (SAPAS) is an eight-item questionnaire developed to screen for the likely presence of a personality disorder (Moran et al., 2003). Each question prompts respondents to endorse specific personality traits (yes/no), yielding a total score between 0 and 8 where a cut-off >3 is indicative of cases with a high probability of diagnosable personality disorders. The WSAS and SAPAS were gathered at the time of initial assessments.

De-identified treatment and demographic data were also available, including information on referral sources, the intensity and sequence of treatments received (LIT and/or HIT along the stepped care pathway), age, gender, ethnicity and employment status. Formal diagnostic assessments were not carried out in routine care, but primary presenting problems noted in clinical records were available in summary form as group-level percentages.

Sample Characteristics

The study included case records for a total of 2,202 patients who had been discharged from the service at the time of data collection. Complete data (described earlier) were available for 1,512 (68.7%) cases. More than half were females (63.9%), with a mean age of 41.99 (SD = 14.54; range = 16 – 87) and of white British ethnic background (88.2%). A quarter (24.9%) of all cases were unemployed and/or in receipt of incapacity benefits; Approximately 59.9% were referred to treatment by general medical practitioners; the remainder self-referred (24.3%) or were referred by other social and health care providers (15.8%). The presenting problems noted in clinical records were depression (21.0%), recurrent depression (6.6%), obsessive–compulsive disorder (4.4%), adjustment disorders (5.7%), somatoform disorders (0.4%), eating disorders (5.7%), other anxiety disorders (42.4%), and unspecified mental health problems (13.4%). Mean baseline severity scores for the whole cohort were PHQ-9 = 14.86 (SD = 6.33), GAD-7 = 13.27 (SD = 5.07), WSAS = 18.39 (SD = 9.46), SAPAS = 3.82 (SD = 1.89; cases with SAPAS >3 = 54.2%). Many patients had comorbid presentations, where 71.4% of cases had case-level symptoms in both PHQ-9 and GAD-7. Approximately 76.6% of patients were initially assigned to LIT; 28.5% of those initially assigned to LIT; 28.5% of those initially assigned to HIT.

Statistical Analysis

Consistent with the objectives of the study, data analyses were performed in four stages aiming to develop, validate and assess the clinical utility of a cumulative complexity model. The primary analyses were carried out in the dataset of cases with complete data (N = 1,512). Following a cross-validation approach, we partitioned this dataset into two random halves which were treated as estimation (N = 755) and validation (N = 757) samples. In order to assess the potential influence of missing data, a single imputed estimation sample (N = 1,108) was derived using an expectation-maximization method (Schafer & Olsen, 1998) and was used for sensitivity analyses described below.

Stage I involved the development of a prognostic index and classification method to identify complex cases in routine care. The dependent variable in all models was a binary indicator of post-treatment reliable and clinically significant improvement (RCSI), with separate models for depression (PHQ-9) and anxiety (GAD-7) measures. RCSI was determined using the criteria proposed by Jacobson and Truax (1991), based on combining reliable change indices for PHQ-9 (≥6) and GAD-7 (≥5) described by Richards and Borglin (2011) and diagnostic cut-offs for each measure (PHQ-9 < 10; GAD-7 < 8). The dependent variable was coded as follows: 0 = RCSI; 1 = no RCSI, such that the prognostic models would be constructed to identify (more complex) cases with increased probability of poor outcomes.

As an initial variable screening step, we used univariate logistic regressions to examine the goodness-of-fit (based on −2 log likelihood test and magnitude of AIC and BIC statistics) of linear and nonlinear trends for continuous variables, as well as alternative ways to model the SAPAS Questionnaire (as a total score, dichotomized based on a cut-off >3, or entered as a series of eight binary items). Entering all eight SAPAS binary items yielded the best fitting models in preliminary tests (i.e., lowest AIC and BIC, significant −2 log likelihood tests) and confirmed that only five items were significant (p < .05) predictors of outcome. Furthermore, baseline severity (PHQ-9, GAD-7), impairment (WSAS) and age variables were optimally modeled using nonlinear trends. Age was rescaled to ordinal decade groups (e.g., teens, twenties, thirties, etc.) and reverse scored (oldest group coded 0, youngest group coded 6) on the basis of the observed trend of correlations between age and RCSI.

Informed by these preliminary tests, we applied penalized categorical regressions with optimal scaling (CATREG-Lasso) in the main analysis. CATREG applies classical linear regression to predictor variables that are transformed to categorical quantifications which are optimally suited to explore nonlinear relations in the data (Gifi, 1990). Continuous variables were thus transformed using a monotonic spline scaling level to examine nonlinear associations with the dependent variable. Variable selection and regularization were performed combining the Lasso procedure (Tibshirani, 1996) and the .632 bootstrap resampling method (Efron & Tibshirani, 1997). The Lasso imposes a penalty term that shrinks coefficients toward zero, penalizing the sum of the squared regression coefficients. This yields more generalizable prediction equations compared with conventional regression models which are prone to overfitting and are less reliable in the presence of multicollinearity. Because using different penalty terms results in different shrunken coefficients, resampling techniques are often used to determine an optimal penalty. The .632 bootstrap resampling method is a smoothed version of the leave-one-out cross-validation strategy, which permits the estimation of a model’s expected prediction error. This resampling method was applied 1,000 times to each Lasso model, iteratively increasing the penalty term in 0.01 units, until all coefficients were shrunk to zero. The one-standard-error rule was applied to select the most parsimonious Lasso model within one standard error of the model with minimum expected prediction error. The predictors entered into CATREG-Lasso models included clinical (baseline PHQ-9, GAD-7, WSAS), personality (SAPAS items 1, 2, 3, 5, 7) and demographic variables (age groups, gender, ethnicity, employment status). Shrunken coefficients from the optimal models were used to calculate a prognostic index (PI) for each patient, where a higher PI denotes poorer...
prognosis. PIs were retained in the CATREG quantifications scale, with signed and continuous scores centered at zero.

The above procedure was conducted in the estimation samples with complete and imputed data, allowing us to compare the area under the curve (AUC) for the PIs derived from each dataset as an indicator of predictive accuracy. PIs derived using complete and imputed samples had comparable AUC statistics albeit with some shrinkage observed in the imputed dataset (PHQ-9: 0.67 ± 0.04 vs. 0.63 ± 0.05; GAD-7: 0.74 ± 0.04 vs. 0.66 ± 0.04). Therefore, subsequent analyses were applied in the dataset with complete data.

In Stage II, we applied receiver operating characteristic (ROC) curve analysis (Altman & Bland, 1994) in the estimation sample to determine empirical cut-offs that optimally balanced sensitivity and specificity on each PI. Consistent with our assumptions about clinical complexity, cases where both (PHQ-9 and GAD-7) PIs were above empirical cut-offs were classed as complex (Cx), and others (including all those with subclinical symptoms) were classed as standard (St) cases. The agreement of both PIs was taken as a conservative means of minimizing “false positive” classifications, and limiting the Cx classification to cases with the poorest prognoses across both outcome domains. We then tested our assumptions about prognosis and cumulative complexity in the validation subsample, with cases whose symptoms were above diagnostic cut-offs for each outcome measure (PHQ-9: N = 675; GAD-7: N = 755). ROC curve analyses were used to assess how well the PIs (using Lasso-based shrunk coefficients from the estimation sample) performed out-of-sample (in a statistically independent validation sample). In addition, separate logistic regression models were applied for each outcome (PHQ-9, GAD-7), where the dependent variable was post-treatment RCSI status (0 = no RCSI; 1 = RCSI) and the predictors included case complexity (0 = Cx, 1 = St) controlling for baseline severity of symptoms (PHQ-9 or GAD-7, respectively).

Stage III analyses were also conducted in the validation sample. A logistic regression model predicting (HIT vs. LIT) group membership based on clinical and demographic characteristics was performed to estimate propensity scores, denoting the predicted probability of completing a treatment episode at HIT. Propensity scores were entered as a covariate in subsequent analyses to control for confounding by indication. Next, logistic regression models were applied with RCSI status as a dependent variable, entering baseline severity (PHQ-9 or GAD-7, respectively), propensity scores, and treatment pathway (LIT or HIT only vs. LIT + HIT) as predictors. The models were performed separately in the subgroups of Cx (N = 269) and St (N = 425) cases (with available data to estimate propensity scores), to minimize multicollinearity between propensity scores and case complexity dummy variables in the same model.

In Stage IV, we assessed the extent to which initial treatment assignment (LIT or HIT) determined by clinical judgment was consistent with the assignment that would be indicated by the prognostic method described previously. A prognostic treatment assignment was coded for all patients, where starting at HIT was recommended for Cx cases and starting at LIT was recommended for all other cases. Next, agreement codes were noted for each case in the full sample, where “1” indicated agreement between clinical judgment and prognosis, and “0” indicated disagreement. Agreement codes were aggregated across the entire sample to estimate a “hit rate,” denoting the percentage of cases where clinicians’ decisions converged with a prognostic strategy for treatment assignment. Next, we applied Cohen’s kappa across agreement codes to derive a Treatment Matching precision (TMaP) score, which takes into account the probability that “hit rates” may be due to chance. The TMaP score is therefore a robust measure of convergence between clinical and empirical decision-making strategies, ranging between 1 (perfect agreement) and −1 (complete disagreement), where 0 is indicative of agreement by chance. TMaP scores were estimated for the full sample and for individual clinicians that undertook initial assessments and made decisions about treatment assignment for at least 20 patients (to eschew extreme scores in caseloads with small base rates).

Results

Estimation of Prognostic Equations

Using the CATREG-Lasso procedure in the estimation sample, we arrived at prognostic models that explained between 9% (PHQ-9: optimal scaling adjusted $R^2 = 0.09$) and 15% (GAD-7: adjusted $R^2 = 0.15$) of variance in posttreatment RCSI. Regression and ROC curve model estimates for each outcome measure are presented in Table 1 (with detailed outputs in Appendix A). Several predictors were selected into optimal Lasso models, including demographic (age, ethnicity, employment), personality (SAPAS items: $2 = $interpersonally avoidant$, 3 = suspicious$, $5 = impulsive$, $7 = dependent$), and clinical features (baseline PHQ-9, GAD-7, WSAS).

The $R^2$ share statistic reflects the relative contribution of each predictor to the model’s overall adjusted $R^2$, after partialing out the specific and combined effects of the other variables. In the depression model, demographics had relatively greater explanatory influence (22.5%) relative to personality (14.7%) and clinical features (15%), although the remaining $R^2$ variance was large (47.9%) and reflected the combined influence of all variables in the model. In the anxiety model, clinical features (55.9%) had two to three times greater explanatory power relative to personality (23.9%) and demographic features (15.2%), leaving only 5% of the remaining $R^2$ variance to combined effects. The F tests for specific variables in both models suggested that the removal of clinical factors (particularly PHQ-9) significantly deteriorated the predictive power of regression models. AUC statistics for the depression ($0.67, SE = 0.02$) and anxiety ($0.74, SE = 0.02$) prognostic indices applied to predict RCSI in the estimation sample were both statistically significant ($p < .001$; ROC curves are shown in Appendix B).

Validation of Case Complexity Model

PIs using the shrunk coefficients derived from the estimation sample were applied in the validation sample, yielding stable and statistically significant ($p < .001$) AUC estimates for depression ($0.64, SE = 0.02$) and anxiety ($0.70, SE = 0.02$) measures (see ROC curves in Appendix B). Overall, 28.6% of all patients were classified as Cx by the prognostic classification rule derived using ROC curve analyses. The proportion of Cx cases was lower in the subsample of patients who only accessed LIT (15.9%) by compar-
**Main Findings**

This study set out to contribute to the understanding of case complexity in psychological care, in view of the limited conceptual clarity and evidence base surrounding this topic. Our findings demonstrate that (a) several patient characteristics have a cumulative effect on treatment outcomes, (b) it is possible to make reasonably accurate prognoses using this information, and (c) prognostic models can help us to operationalize case complexity in a way that is clinically useful. Cases classed as Cx (28.6%) on the basis of prognostic data tended to have significantly poorer outcomes after psychological treatment. Furthermore, Cx cases were two times (OR = 2.23) more likely to attain RCSI in depression symptoms if they were initially assigned to a high-intensity intervention instead of usual stepped care. A similar trend was observed for anxiety symptoms, although this did not reach statistical significance.

**Discussion**

The aggregated hit rate in the full sample indicated that clinicians’ treatment assignment decisions agreed with the prognostic strategy in 65.6% of cases. The TMaP score for the full sample, however, was low (κ = 0.09, SE = 0.02, p < .001). A closer examination of individual therapists’ treatment assignment decisions (N = 1,247 nested within 26 therapists) revealed considerable variability in their hit rates (range = 36.5% to 84.7%; M = 62.9, SD = 14.3) and TMaP scores (range = −0.27 to 0.44; M = 0.05, SD = 0.20). As shown in Figure 3, hit rates and TMaP scores were moderately correlated (r = .67, p < .001), and approximately 48% of therapists had TMaP scores <0.

**Table 1**

<table>
<thead>
<tr>
<th>Predictor</th>
<th>B</th>
<th>SE</th>
<th>R² share</th>
<th>Removal impact</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gender (female = 0, male = 1)</td>
<td>.000</td>
<td>.002</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Ethnicity (White British = 0, minority ethnic group = 1)*</td>
<td>.000</td>
<td>.009</td>
<td></td>
<td>.000</td>
</tr>
<tr>
<td>Employment (employed = 0, unemployed = 1)*</td>
<td>.065</td>
<td>.037</td>
<td>13.7%</td>
<td>.067</td>
</tr>
</tbody>
</table>

**Depression (PHQ-9) model parameters**

F(591) = 4.13, p < .001

Adjusted R² = .09

AUC = .67, SE = .02 (.64, .71), p < .001

**Anxiety (GAD-7) model parameters**

F(643) = 6.85, p < .001

Adjusted R² = .15

AUC = .74, SE = .02 (.70, .78), p < .001

**Lasso-based coefficients**

<table>
<thead>
<tr>
<th>Predictor</th>
<th>B</th>
<th>SE</th>
<th>R² share</th>
<th>Removal impact</th>
</tr>
</thead>
<tbody>
<tr>
<td>SAPAS item (not endorsed = 0, endorsed = 1)</td>
<td>.000</td>
<td>.026</td>
<td></td>
<td>.000</td>
</tr>
</tbody>
</table>

**Note.** Dependent variables in both models are categorical markers for post-treatment reliable and clinically significant improvement (0 = Reliable and clinically significant improvement (RCSI), 1 = no Reliable and clinically significant improvement (RCSI)). Beta coefficients are expressed in a Categorical Quantification Scale; AUC = area under the curve statistic; R² share = squared partial correlation between predictor and outcome/adjusted R²; Removal impact = F test probability of model deterioration if the predictor is removed; SE = standard errors aggregated over 1,000 bootstrap samples. SAPAS = Standardized Assessment of Personality–Abbreviated Scale; PHQ-9 = Patient Health Questionnaire-9; GAD-7 = Generalized Anxiety Disorder-7; WSAS = Work and Social Adjustment Scale.

* Predictors selected into optimal Lasso model. A “no” answer to Item 3 is reverse scored = 1.

**Case Complexity and Treatment Selection**

Logistic regression models presented in Table 3 indicated that Cx cases had a significantly greater probability of RCSI in depression symptoms if they directly accessed HIT, by comparison to a standard stepped care pathway LIT + HIT (OR = 2.23, p = .01). There was also a trend indicating the same advantage of HIT for Cx cases in the anxiety model, although this did not reach statistical significance (OR = 1.74, p = .08). No significant differences were found between treatment pathways in the regression models applied to St cases. These analyses controlled for baseline symptom severity and propensity scores (derived from logistic regression model in Appendix C). The results for the depression outcomes are illustrated in Figure 2; where Cx cases that were initially assigned to HIT (optimal prognostic treatment assignment) had a 16.3% increased probability of RCSI by comparison to Cx cases that were assigned to a conventional stepped care pathway (LIT + HIT).

**Clinical Judgment Versus Prognostic Models**

The aggregated hit rate in the full sample indicated that clinicians’ treatment assignment decisions agreed with the prognostic
A Conceptual Bridge Between Prognosis and Case Complexity

These results lend support to the clinical notion that some cases are more difficult to treat due to various complicating factors (Ruscio & Holohan, 2006), although clinicians’ intuitions and treatment planning are often inconsistent with the evidence base (Garb, 2005). We found that treatment assignment decisions guided by clinical judgment were consistent with prognostic models in 65.6% of cases. This rate of agreement could be achieved by chance, or simply by mechanically following stepped care guidelines and assigning all cases initially to LIT, because the base rate of Cx cases is relatively low (under 30%). This was evidenced more clearly by examining the aggregated TMaP score (0.09) which was close to zero. Overall, the findings indicate that depression improvement (RCSI) rates for Cx cases could be significantly increased (by approximately 16.3%) if clinical judgment was supported by prognostic treatment selection models.

This gap between practice and science is perhaps accentuated by an unwieldy literature on the topic of prognosis in psychological care. Previous authors have attempted to synthesize findings across multiple studies to elucidate predictors of depression and anxiety outcomes (e.g., Driessen & Hollon, 2010; Haby, Donnelly, Corry, & Vos, 2006; Hamilton & Dobson, 2002; Keeley et al., 2008; Kessler et al., 2017; Licht-Strunk et al., 2007; Nilsen, Eisemann, & Kvernmo, 2013). Although some convergent findings are evident, meta-analytic reviews that privilege data from clinical trials are limited by typically small samples with sparse and heterogeneous prognostic variables, often gathered in highly selected participants (i.e., those with specific disorders) that may not be representative of complex cases seen in routine care (Chambless & Ollendick, 2001). Naturalistic cohort studies can offer informative evidence to complement findings from controlled trials, especially where multiple variables are measured systematically across large health care populations, as exemplified in this study. Several such studies are

Table 2
Validation of Prognostic Indices Applied Out-of-Sample-Using Logistic Regression

<table>
<thead>
<tr>
<th>Predictor</th>
<th>Depression (PHQ-9) model parameters</th>
<th>Anxiety (GAD-7) model parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>B</td>
<td>SE</td>
</tr>
<tr>
<td>Baseline severity a</td>
<td>-.06**</td>
<td>.02</td>
</tr>
<tr>
<td>Case complexity (Cx = 0, St = 1)</td>
<td>.59**</td>
<td>.20</td>
</tr>
<tr>
<td>Constant</td>
<td>.57</td>
<td>.46</td>
</tr>
</tbody>
</table>

*Note.* Nagelkerke $R^2 = .08$. Dependent variables in both models are categorical markers for post-treatment reliable and clinically significant improvement (0 = no Reliable and clinically significant improvement (RCSI), 1 = Reliable and clinically significant improvement (RCSI)). PHQ-9 = Patient Health Questionnaire-9; GAD-7 = Generalized Anxiety Disorder-7; Cx = complex cases; St = standard cases.

a The baseline severity measure (either PHQ-9 or GAD-7) entered in each model matched the relevant outcome variable.

** $p < .01$. ***$p < .001$. 

Figure 1. Reliable and clinically significant improvement (RCSI) in cases classified as standard (St) and complex (Cx).
yielding replicated findings (e.g., Beard et al., 2016; Delgadillo, Moreea, & Lutz, 2016; Delgadillo, Dawson, Gilbody, & Böhnke, 2017; Firth, Barkham, Kellett, & Saxon, 2015; Goddard, Wingrove, & Moran, 2015; Licht-Strunk et al., 2009).

Overall, the emerging literature on outcome prediction points to factors clustered around clinical (i.e., baseline symptom severity, diagnosis, comorbidity, functioning and disability, physical illnesses), demographic (i.e., age, ethnicity, employment, socioeco-

Table 3

Logistic Regression Models Assessing Case Complexity and Treatment Selection

<table>
<thead>
<tr>
<th>Predictor</th>
<th>Depression (PHQ-9) model parameters</th>
<th>Anxiety (GAD-7) model parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>B</td>
<td>SE</td>
</tr>
<tr>
<td>Subsample of Cx casesa</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Baseline severityb</td>
<td>-.10**</td>
<td>.03</td>
</tr>
<tr>
<td>Propensity score</td>
<td>-3.28*</td>
<td>1.27</td>
</tr>
<tr>
<td>Treatment = LIT + HIT (ref)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Treatment = LIT</td>
<td>-.02</td>
<td>.34</td>
</tr>
<tr>
<td>Treatment = HIT</td>
<td>.80*</td>
<td>.31</td>
</tr>
<tr>
<td>Constant</td>
<td>3.67**</td>
<td>1.23</td>
</tr>
</tbody>
</table>

| Subsample of St casesc |         |          |        |        |          |          |        |        |
| Baseline severityb  | -.03     | .03      | .97    | .91, 1.03| .05      | .03      | 1.05  | .98, 1.11|
| Propensity score    | -.51     | .98      | .60    | .09, 4.06| -3.69*** | .92      | .03   | .004, .15|
| Treatment = LIT + HIT (ref) |          |          |        |        |          |          |        |        |
| Treatment = LIT    | -.14     | .25      | .87    | .53, 1.43| -.18     | .24      | .83   | .52, 1.34|
| Treatment = HIT    | .32      | .34      | 1.38   | .70, 2.70| .00      | .32      | 1.00  | .54, 1.86|
| Constant           | 1.03     | .71      | 2.79   |         | 1.81**   | .66      | 6.13  |         |

Note. Dependent variables in both models are categorical markers for post-treatment reliable and clinically significant improvement (0 = no Reliable and clinically significant improvement (RCSI), 1 = Reliable and clinically significant improvement (RCSI)); PHQ-9 = Patient Health Questionnaire-9; GAD-7 = Generalized Anxiety Disorder-7; Cx = complex cases; LIT = low intensity treatment; HIT = high intensity treatment; ref = reference category; St = standard cases.

a For depression model parameters, Nagelkerke $R^2 = .11$; for anxiety model parameters, Nagelkerke $R^2 = .10$. b The baseline severity measure (either PHQ-9 or GAD-7) entered in each model matched the relevant outcome variable. c For depression model parameters, Nagelkerke $R^2 = .01$; for anxiety model parameters, Nagelkerke $R^2 = .06$.

$p < .05$. ** p < .01. *** p < .001.

Overall, the emerging literature on outcome prediction points to factors clustered around clinical (i.e., baseline symptom severity, diagnosis, comorbidity, functioning and disability, physical illnesses), demographic (i.e., age, ethnicity, employment, socioeco-

nomic depravation, marital status), characterological (i.e., personality disorder diagnoses or traits, interpersonal problems and style, trait anxiety and neuroticism), and dispositional domains (i.e., readiness to change, expectancy). Informed by advances in the biomedical literature (Safford et al., 2007; Shippee et al., 2012), we propose that complex cases in psychological care are characterized by the presence of measurable factors that map onto multiple domains (clinical, demographic, characterological and dispositional), which are statistically associated with clinical prognosis and have a cumulative—detrimental—effect on treatment outcomes. The concept of case complexity is, therefore, dimensional.

Figure 2. Reliable and clinically significant improvement (RCSI) in cases classified as standard (St) and complex (Cx) according to treatment pathway.

Figure 3. Distribution of hit rates and treatment matching precision (TMaP) scores across 26 therapists.
(i.e., degrees of complexity on a continuum), and complex cases can be distinguished from others using empirically derived population norms and classification rules.

Case complexity may challenge psychological improvement through several mechanisms. One possibility is that an accumulation of disadvantages (e.g., poverty, interpersonal difficulties, functional impairment, outgroup derogation due to minority ethnic status) could disrupt the balance between life stressors and coping resources (Shippee et al., 2012). Complexity could also interfere with adequate engagement with therapy, for example, by undermining expectancy, which is a well-established predictor of treatment outcomes (Constandino et al., 2011). Baseline severity is an important contributor to complexity, so another possibility is that high baseline severity does not completely block improvement but may dampen the effect of treatment (i.e., cases with high severity can attain reliable improvement even if their symptoms do not reach subclinical levels). Furthermore, our findings suggest that specific features (i.e., demographic, clinical, characterological) influence specific clinical outcomes (remission of depression, anxiety) differentially. For example, demographic factors (e.g., young age, unemployment) had a considerably larger influence over depression outcomes relative to clinical and characterological factors. Future research could focus on exploring the relative contribution of different prognostic domains to multiple outcome domains (symptoms, quality of life, functioning) and the mechanisms through which these cumulative disadvantages may complicate or undermine treatment.

Limitations

Some limitations should be considered when interpreting the results of this study. As is common in naturalistic data sets, we encountered several cases with missing data (>30%). To deal with this, we applied multiple imputation and sensitivity analyses that yielded similar prognostic models, albeit with some shrinkage observed in the imputed dataset. On this basis, it was appropriate to perform further validation analyses using cases with complete data, to simulate how prognostic assessments would be applied in routine care, where data imputation of missing values is unfeasible.

Another limitation concerning the data used in this study was that we only had access to pre-post outcome measures for the entire treatment pathway, and it was not possible to disaggregate the outcomes for LIT and HIT for cases that accessed both steps. However, we were able to determine that Cx that only accessed HIT tended to have better outcomes compared to those who accessed LIT + HIT (a lengthier and costly treatment pathway). This suggests that there are no benefits of having LIT sessions preceding HIT, and hence the advantage of being initially assigned to HIT may not be solely due to having a lengthier treatment episode. Previous research using more granular outcomes data for each treatment step suggested that cases with poor prognostic features had a higher probability of dropout and lower probability of improvement at the LIT step by comparison to HIT (Delgadillo et al., 2016). These emerging findings suggest that assigning complex cases directly to HIT seems justified, although future randomized controlled trials of this strategy are necessary to determine if it is indeed more cost-effective.

Other limitations include the lack of formal diagnostic assessments and the analysis of a limited number of prognostic variables. It is known, for example, that specific diagnoses such as post-traumatic stress disorder, eating disorders and obsessive–compulsive disorder are associated with poorer outcomes in stepped care services (Delgadillo et al., 2017), and it is plausible that such diagnoses could interact with other prognostic features. Notwithstanding these limitations, it is remarkable that this narrow range of variables yielded an accurate and clinically useful prognostic model. Other studies using routine practice data have shown that similar variables can be used to identify subgroups of cases with depression and anxiety problems that attain similar outcomes (Delgadillo et al., 2016; Lutz, Lowry, Kopta, Einstein, & Howard, 2001; Lutz et al., 2005; Saunders, Cape, Fearon, & Pilling, 2016).

Clinical Implications

In line with recent findings in stepped-care psychological treatment settings (Delgadillo et al., 2016; Lorenzo-Luaces, DeRubeis, van Straten, & Tiemens, 2017), the present study provides further evidence that applying prognostic indices to guide personalized treatment recommendations is likely to improve treatment outcomes. Low-intensity guided self-help interventions are recommended as first-line treatments for several common mental disorders (NICE, 2011) and are becoming widely available in routine stepped care services (Clark, 2011). The application of evidence-based treatment selection algorithms like the one demonstrated in this study could help to maximize the cost-effectiveness of LIT by selectively offering it to those who are most likely to derive benefits. Equally, prognostic models could be used to fast-track complex cases to HIT in a timely way.

References

Appendix A

Penalized Categorical Regression Models With Optimal Scaling

Table A1

<table>
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<th>Sum of squares</th>
<th>df</th>
<th>M square</th>
<th>F</th>
<th>p</th>
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<tr>
<td>Regression</td>
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<td>4.130</td>
<td>.000</td>
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<td>Residual</td>
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<td>579</td>
<td>.942</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>592.000</td>
<td>591</td>
<td></td>
<td></td>
<td></td>
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</tbody>
</table>

Note. N = 592; adjusted $R^2 = 0.09$; dependent variable = PHQ9_No_RCSI. ANOVA = analysis of variance; PHQ-9 = Patient Health Questionnaire-9.