Are the PHQ-9 and GAD-7 Suitable for Use in India? A Psychometric Analysis

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Background: Cross-cultural evidence on the factorial structure and invariance of the PHQ-9 and the GAD-7 is lacking for South Asia. Recommendations on the use of unit-weighted scores of these scales (the sum of items’ scores) are not well-founded. This study aims to address these contextual and methodological gaps using data from a rural Indian population.

Methods: The study surveyed 1,209 participants of the Kerala Diabetes Prevention Program aged 30–60 years (n at risk of diabetes = 1,007 and n with diabetes = 202). 1,007 participants were surveyed over 2 years using the PHQ-9 and the GAD-7. Bifactor-(S – 1) modeling and multigroup confirmatory factor analysis were used.

Results: Factor analysis supported the existence of a somatic and cognitive/affective subcomponent for both scales, but less explicitly for the GAD-7. Hierarchical omega values were 0.72 for the PHQ-9 and 0.76 for the GAD-7. Both scales showed full scalar invariance and full or partial residual invariance across age, gender, education, status of diabetes and over time. Effect sizes between categories measured by unit-weighted scores versus latent means followed a similar trend but were systematically higher for the latent means. For both disorders, female gender and lower education were associated with higher symptom severity scores, which corresponds with regional and global trends.

Conclusions: For both scales, psychometric properties were comparable to studies in western settings. Distinct clinical profiles (somatic-cognitive) were supported for depression, and to a lesser extent for anxiety. Unit-weighted scores of the full scales should be used with caution, while scoring subscales is not recommended. The
stability of these scales supports their use and allows for meaningful comparison across tested subgroups.

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http://www.anzctr.org.au/Trial/Registration/TrialReview.aspx?id=336603&isReview=true.

**Keywords:** Patient Health Questionnaire, India, measurement invariance, depression, generalized anxiety disorder assessment, sum score reliability, confirmatory bifactor modeling, omega hierarchical

### INTRODUCTION

Depressive and anxiety disorders are among the ten leading causes of global disability (Vos et al., 2015) and more than 80% of people who have mental disorders reside in low- and middle income countries (LMICs) (World Health Organization, 2008). In India, depressive and anxiety disorders have been shown to both have a crude prevalence of 3.3% and were responsible for 33.8% (29.5–38.5) and 19.0% (15.9–22.4) of disease-adjusted life years attributable to mental disorders (Sagar et al., 2020).

Self-reported measurement tools are crucial to estimate the burden of depressive and anxiety disorders at population level, to determine how this burden relates to subgroup characteristics (e.g., sociodemographic characteristics, other health conditions, etc.), and to measure the effect of public health interventions. At individual level, these tools can enhance the reliability of diagnoses and their ease of use makes them particularly useful in settings with poor mental health service provision and a lack of specialized staff (Mughal et al., 2020). However, most of the established self-reported measurement tools have been developed and evaluated in Europe and North America and may not perform in an equivalent way in different cultures or settings (Dere et al., 2015; Mughal et al., 2020).

The Patient Health Questionnaire (PHQ-9) and Generalized Anxiety Disorder (GAD-7) assessment can be used as screening tools as well as measures of symptom severity for depression (PHQ-9) and different types of anxiety (GAD-7) (Kroenke et al., 2001; Spitzer et al., 2006). Both tools are based on the Diagnostic and Statistical Manual of Mental Disorders criteria and have been found to be valid measures for detecting and monitoring depression or anxiety disorders in western countries (Kroenke et al., 2010). In South Asia, a region home to one quarter of the world’s population, assessment of these scales’ essential psychometric properties is lacking (Lamela et al., 2020). Below, we will discuss why assessment of such properties is crucial and their current level of evidence, focusing on: (1) the factorial structure; (2) the use of unit-weighted scores (i.e., the sum score of the item responses); and (3) invariance across subgroups.

The factorial structure of data from a specific population can provide empirical support for the potential existence of subdimensions of depression and anxiety disorders which can differ across cultures and settings (Leong and Tak, 2003). For instance, studies have revealed that somatic symptoms are more common in Indian people with depression compared with western populations (Grover et al., 2010). For the PHQ-9 and the GAD-7, which were initially intended to be used as unidimensional scales, a variety of measurement models have been proposed based on confirmatory factor analysis (CFA) investigations (Doi et al., 2018a,b; Moreno et al., 2019; Lamela et al., 2020). In these studies, mainly conducted in western settings, researchers have found both scales to fit unidimensional and multidimensional models (Lamela et al., 2020), a second-order model specifically for the GAD-7 (Doi et al., 2018a) and a bifactor model specifically for the PHQ-9 (Doi et al., 2018b). A recurrent finding is the identification of a factor consisting of items reflecting a somatic aspect and a factor consisting of items reflecting a cognitive/affective aspect (Beard and Björgvinsson, 2014; Lamela et al., 2020). For both disorders, these subdimensions correspond with clinical representations and pathophysiologic insights suggesting different subtypes of these conditions (Portman et al., 2011; Duivis et al., 2013; Lamela et al., 2020). For depression in particular, distinguishing between these subtypes has been shown important with regards to treatment and prognosis (Lamela et al., 2020). However, to our knowledge, no studies have assessed the factorial structure of these scales in India or anywhere else in South Asia. To examine the cross-cultural validity of these subdimensions, evidence on the factorial structure of these scales based on data from different settings is needed.

A key question when using self-reported measurement tools is to what extent the unit-weighted scores (i.e., the sum score of all or a subset of item responses) can be interpreted as a unidimensional representation of a specific construct. For instance, to what extent does the sum of all scale item scores represent depressive or anxiety symptom severity as an overarching construct. Items of these scales may belong to different subdimensions which may preclude the interpretation of the sum of their scores. In addition, clinicians or researchers may want to sum item responses belonging to one of these subdimensions and use this score as a reflection of that specific subdimension. Recent studies have defended scoring the total scale as well as its subdimensions for the PHQ-9 (Doi et al., 2018b; Lamela et al., 2020) and for the GAD-7 (Doi et al., 2018a) guided by the goodness of fit of CFA models. These recommendations are problematic for several reasons. First, Reise et al. (2013) argue that model selection based on CFA rarely informs researchers on the degree of multidimensionality such that it may justify the use of unit-weighted scores of subscales or total scales. Second, methodologists have called into question...
if specifying both, total and subdimension scores of the same scale can have an added value (Rodriguez et al., 2016a). Third, consensus is lacking on the choice of the most adequate CFA model for both, the PHQ-9 and GAD-7. To guide researchers on this matter, alternative analytic techniques have been proposed such as the Haberman (2008) procedure (Haberman, 2008) and the use of model-based reliability indices (Rodriguez et al., 2016b). To our knowledge, these techniques have not been applied yet to either the PHQ-9 or the GAD-7. Application of these techniques to data collected from different settings may redress current inconsistencies in recommendations on the use of unit-weighted scores.

Measurement invariance of a scale is an essential psychometric property when studying scale scores over time or between subgroups of a population (Vandenberg and Lance, 2000). Measurement invariance across subgroups corresponds to a latent construct being represented by the scale items in a similar way and suggests that this construct has a similar meaning to these groups. This implies that assessing invariance is crucial for the comparison of subgroups. When measurement invariance is violated across subgroups, prevalence or severity of a disorder may be under- or overestimated across these groups. Finally, analysis of invariance can provide insight into how the interpretation of a scale and the perception of an illness may differ across subgroups. This may have consequences with regards to the diagnosis and how people cope with their illness. Lack of invariance can lead to substantial bias when comparing different subgroups, especially when using unit-weighted scores (Steinmetz, 2013). Moreover, in addition to scalar invariance which is required to compare subgroups through a structural equation modeling (SEM) framework, the use of unit-weighted scores requires invariant indicator reliability (Vandenberg and Lance, 2000). Invariance of the PHQ-9 and GAD-7 scales has been supported by studies conducted in western settings for gender, ethnic and sociodemographic differences, but only few studies assessed invariance for age and over time (Hinz et al., 2017; Lamela et al., 2020). Moreover, other depression measures have shown non-invariance across different age groups (Estabrook et al., 2015). Only a handful of studies have tested invariance in LMICs and were focused on a specific population such as college students, pregnant women, etc. (Miranda and Scoppetta, 2018). To our knowledge, no studies have assessed any form of invariance of the PHQ-9 and GAD-7 in South Asian populations.

In sum, essential psychometric properties of the PHQ-9 and the GAD-7 remain underexplored among LMIC populations, in particular in South Asia. Furthermore, recommendations on the use of unit-weighted scores of these scales have been poorly supported.

The aim of this study was to simultaneously address these contextual and methodological gaps based on a state-of-the-art analytic approach and using data from a rural Indian population. Specifically, we aimed to: (1) assess existence of subdimensions in this population by assessing the factorial structure of data collected through these scales; (2) determine how precisely total and subscale scores reflect their intended constructs; (3) assess invariance across subgroups of age, gender, level of education, status of diabetes (at risk of versus with type 2 diabetes [T2D]) and different measurement occasions; and (4) if invariance could be established, compare the results of unit-weighted scores with latent means analysis in the assessment of differences in symptom severity across the same subgroups. With this last objective, we sought to assess: (1) the accuracy of unit-weighted scores when using latent mean levels as a standard; and (2) whether differences across subgroups correspond with regional trends that would support the validity of our data.

MATERIALS AND METHODS

Participants
The analysis was based on data collected from participants who took part in the baseline, cross-sectional, community-based survey of a diabetes prevention program in the state of Kerala in India: the Kerala Diabetes Prevention Program (K-DPP). A detailed description of the study design, participant screening and recruitment has been previously published (Sathish et al., 2013, 2017, 2019). Briefly, K-DPP was a cluster randomized controlled trial conducted in 60 randomly selected polling areas (electoral divisions) from a taluk (sub-district) in Trivandrum district of Kerala state. People aged 30–60 years were selected randomly from the electoral roll of the 60 polling areas and were approached at their households by trained data collectors. We screened 3,421 individuals for eligibility and those with a history of diabetes or other major chronic conditions, taking drugs influencing glucose tolerance (e.g., steroids), and who were illiterate in the local language were excluded (n = 835). Potentially eligible individuals (n = 2586) were screened with the Indian Diabetes Risk Score, and those with a score of ≥60 (n = 1529) were invited to undergo a 2-h oral glucose tolerance test (OGTT) at community-based clinics. Of these, 1,209 attended the clinics, of which 1,007 individuals were at high risk for developing diabetes and 202 were diagnosed with diabetes. Participant screening and recruitment were completed between January and October 2013. The 1,007 individuals at high risk were followed-up after 1 and 2 years of enrollment. These follow-up points were used for the analysis of invariance at different measurement occasions. Mean age of participants was 46.0 (SD: 7.5), 45.8% were female, and 95% were married. 25.3, 51.3, and 23.4% attended primary, secondary and higher education, respectively (Sathish et al., 2017).

Measures and Data Collection
Both the nine-item PHQ-9 and the seven-item GAD-7 use 4-point Likert-scaled items ranging from 0 (not at all) to 3 (nearly every day) (Kroenke et al., 2001; Spitzer et al., 2006). For the GAD-7, items 4, 5, and 6 have been found to reflect a somatic dimension (Rutter and Brown, 2017). For the PHQ-9, this was the case for items 3, 4, and 5 and in some studies items 7 and 8 (Lamela et al., 2020). The scales were translated to Malayalam and back-translated to English and were pilot tested. Interviews were administered by trained interviewers.
Data-Analysis

Confirmatory Factor Analysis

To evaluate whether our data supported a two dimensional model, a selection of models was tested with specifications based on theory and findings from previous studies. The aim of this analysis was to test if a model with one or two factors would be acceptable, rather than to select a specific model solely based on a better model fit. For the PHQ-9, we tested a correlated 2-factor model with items 3–4–5 loading onto one factor as was proposed by others (Lamela et al., 2020). For the GAD-7, we tested a 1-factor model with and without correlated residuals for items 4, 5, and 6 and a correlated 2-factor model with items 4, 5, and 6 loading onto one factor (Beard and Björgvinsson, 2014; Rutter and Brown, 2017; Doi et al., 2018a). Assessment of these models was based on their χ²-values, the item indicators’ loadings and the following sample-corrected for non-normal data goodness of fit indices (Brosseau-Liard et al., 2012) with target values as proposed by Hu and Bentler (1999): the comparative fit index (CFI) (∼0.95), the Tucker–Lewis index (TLI) (∼0.95), the root mean square error of approximation (RMSEA) (∼0.06), and the standardized root mean square residual (SRMR) (∼0.08). Since items’ distributions departed from normality, we used maximum likelihood estimation with robust (Huber–White) standard errors and a scaled test statistic that is (asymptotically) equal to the Yuan–Bentler test statistic.

Haberman Procedure

This procedure assesses whether the subscore provides a more accurate estimate of the construct it measures than the total score (Haberman, 2008). The proportional reduction of mean squared error (PRMSE) based on total scores was compared with the PRMSE based on subscale scores. In case the latter would be smaller, there is no psychometric justification to report the subscale scores. In addition, a hypothesis test was performed justifying the reporting of subscale scores if Olkin’s Z statistic was higher than 1.64 (Sinharay, 2019). However, this procedure does not test whether subscale scores provide meaningful information, while taking into account the total score (Reise et al., 2013). To assess this, we used the indices described in the next paragraph.

Model-Based Psychometric Indices

Rodriguez et al. (2016a,b) proposed the following indices to assess the degree to which total and subscale scores reflect their intended constructs. Total omega (omegaT) estimates the proportion of variance in the unit-weighted total score due to all common factors including the general and group factors (Zinbarg et al., 2005). Hierarchical omega (omegaH) estimates the proportion variance due to a general factor. Omega hierarchical subscale (omegaHS) estimates the variance due to a specific group factor while controlling for the general factor. Recommended minimum values were described for OmegaH (0.70) and for OmegaHS (0.50) (Reise et al., 2013; Rodriguez et al., 2016a).

The indices were calculated based on a confirmatory bifactor modeling approach (Reise et al., 2013) using the semTools package in R (Jorgensen et al., 2020). A bifactor model was deemed appropriate to calculate these indices as it is a less restrictive model (than, e.g., a hierarchical model) and as the structure of the response data was assumed to be consistent with a bifactor structure: i.e., a single general trait reflecting the target construct and the presence of subdomain constructs due to clusters of similar items (Rodriguez et al., 2016a). Bifactor models can be specified by all items loading onto a general factor as well as on group factors representing the subdomains. In addition, group factors are assumed to be uncorrelated with other group factors as well as with the general factor. However, since only two group factors were present, these bifactor models will be unidentified, which implies that an infinite number of hierarchical factor models can be found for the same covariance matrix (Zinbarg et al., 2007). To address this problem, we used a modified version of the traditional bifactor model: the bifactor-(S − 1) model described by Eid et al. (2016). The name of this model refers to the number of specific group factors being less than the actual number of the scale’s subdimensions being considered. This modification makes the discarded group factor a reference group for the general factor and solves the identification problem (Eid et al., 2016).

Invariance

Invariant indicator reliabilities of unit-weighted scores across groups requires residual or strict invariance which was tested using multigroup confirmatory factor analysis (MGCFA) (Vandenberg and Lance, 2000). The following levels of invariance were assessed: equal form (i.e., configural invariance), equality of factor loadings (i.e., metric invariance), equality of indicator intercepts (scalar invariance), and equality of residuals (i.e., residual invariance) (Vandenberg and Lance, 2000). However, for residual invariance to be analogous to indicator reliability invariance, the last step requires invariance of factor variances which was tested first (Vandenberg and Lance, 2000). Criteria of invariance between nested models included a difference in CFI < −0.01 combined with difference in RMSEA < 0.015 or a non-significant scaled χ²-square difference test (Putnick and Bornstein, 2016).

Group Differences

For the unit-weighted scores, standardized effect sizes between different subgroups were calculated using robust regression of the unit-weighted scores based on MM-estimation (i.e., an extension of the maximum likelihood estimate method). Standardized effect sizes of unit-weighted scores were compared with the standardized effect sizes of the difference in latent mean levels estimated through multigroup structural equation modeling. Data were analyzed using R software with the packages “lavaan” (Rosseel, 2012) and “semTools” (Jorgensen et al., 2020).

Missing Data

Missing data occurred in 0.4% of the GAD-7 data, in 3.4% of the PHQ-9 data and in 0.0% of the demographic variables (sex, education, and age). It was deemed implausible that the probability of missing data would significantly differ in specific groups or cases, assuming they were missing completely at random (MCAR). This was supported by Little’s test hypothesis not being rejected for a subset of the GAD-7 and demographic variables (p = 0.30) and the PHQ-9 and
demographic variables ($p = 0.07$). For these reasons, complete case analysis was preferred.

**Ethical Approval**

The study was approved by the Institutional Ethics Committee of the Sree Chitra Tirunal Institute for Medical Sciences and Technology, Trivandrum, Kerala, and by the Human Research Ethics Committees of Monash University, Australia and the University of Melbourne, Australia. The study was also approved by the Health Ministry Screening Committee of the Government of India.

**RESULTS**

**Factor Structure**

As mentioned previously, the aim of this analysis was to assess the existence of a somatic and a cognitive subdimension in the response data. For this purpose, we assessed whether model fit criteria of 1- and 2-factor models were acceptable (see Figures 1 and 2 and Table 1). CFA of the models proposed for the PHQ-9 revealed an acceptable fit for a 2-factor model, but not for a 1-factor model (see Table 1). The factors of the 2-factor model were highly correlated ($r = 0.77$). Factor loading estimates revealed that the indicators were strongly related to their purported factors (range $\lambda = 0.45$–0.73) with $p$-values below 0.001 (see Figure 1).

CFA of the models proposed for the GAD-7 revealed an acceptable fit for a 1-factor model with correlated residuals between items 3, 5, and 6, and a 2-factor model (see Table 1). The correlation between the cognitive and somatic factor of the 2-factor model was high ($r = 0.84$). Factor loading estimates revealed that the indicators were strongly related to their purported factors (range $\lambda = 0.58$–0.88), except for item 6 ($\lambda = 0.27$) (see Figure 2). For all parameters, $p$-values were below 0.001 except for the correlation between the residuals of items 5 and 6 ($p$-value $= 0.05$).

**Haberman Procedure**

PRMSE based on subscale scores was smaller than the PRMSE of the total score for the somatic dimension of the PHQ-9 (0.57 vs. 0.63) and the GAD-7 (0.63 vs. 0.81) and for the cognitive dimension of the GAD-7 (0.80 vs. 0.83). This was confirmed by a formal hypothesis test with the Olkin’s $Z$ statistic lower than 1.64. These results imply that there is no added value in reporting these subscale scores as total scores would be a relatively more precise indicator of subscale true scores than the actual subscale scores. For the cognitive dimension of the PHQ-9, the PRMSE based on the subscale scores was higher than the PRMSE based on the total score (0.69 vs. 0.62), which indicates that this subscale score is a more precise indicator of its true score.

**Model-Based Psychometric Indices**

As mentioned earlier, a bifactor-($S - 1$) model was fitted to calculate these indices (see Figure 3). For both scales, the somatic factor was discarded making it the reference group for the general factor. Model fit of these models was acceptable for the PHQ-9 and excellent for the GAD-7 (see Table 2). For both scales, estimates of omegaH above the recommended minimum value and the difference between omegaT and omegaH was relatively small (i.e., 0.08 for both scales), which indicates that the general factor is the major determinant of the variance underlying the unit-weighted total scale scores. Estimates of the omegaHS were small for both scales, reflecting little unique variance due to any specific group factor.

**Invariance**

**PHQ-9**

Applying CFA to the different subgroups of gender, age, education, T2D status, and different measurement occasions, resulted in good model fit, except for the age group 30–40 and the group with higher education (see Table 3). Scalar invariance could be established across all different subgroups (see Table 3). Scalar invariance could be established across all different subgroups, except for education. Full or partial residual invariance could be established across all different subgroups. Partial residual invariance across all three measurement occasions required free estimation of residuals of at least six items. Per two measurement occasions, partial invariance could be obtained by freeing two ($T0$–$T1$) to five residuals ($T1$–$T2$).

**GAD-7**

Applying CFA resulted in excellent model fit for all different subgroups using the one-factor model with correlated residuals (see Table 3). Scalar invariance could be established across all different subgroups (see Table 4). Invariance of latent factor variance could be established for all subgroups, except for education. Full or partial (for gender and measurement occasions) residual invariance could be established across all subgroups.

**Group Differences**

For both scales, group differences in symptom severity calculated from unit-weighted scores versus latent mean scores show similar results, although the effect size based on latent mean scores was systematically higher (see Table 5). This difference was more pronounced for the GAD-7. Symptom severity scores were higher among women and groups with lower educational attainment. There was no difference between people with or at risk of T2D. Age groups did not differ, except for higher PHQ-9 scores among the eldest cohort. Severity scores decreased over different measurement occasions.

**DISCUSSION**

This is the first study assessing the factorial structure, unit-weighted score reliability, and invariance of the PHQ-9 and the GAD-7 in population in South Asia. For both scales, our findings support the presence of a dominant general factor, but also suggest a somatic and a cognitive/affective subcomponent. This was less explicit for the GAD-7. However, this multidimensionality was not sufficient to justify the scoring
of subscales. Our findings only supported the use of unit-weighted scores of the full scales. Scales also showed full scalar invariance and partial to full residual invariance across gender, age, education, measurement occasions and T2D status.

For the PHQ-9, only a 2-factor model showed an acceptable fit in our population in support of a somatic and cognitive/affective subcomponent. While initially, previous research proposed a one-factor model for the PHQ-9, this has been attributed to
TABLE 1 | Model fit of the PHQ-9 (n = 1207) and the GAD-7 (n = 1205).

| Scale | Model | \( \chi^2 \) | df | p-value | CFI  | TLI  | RMSEA | 90% CI | SRMR |
|-------|-------|-------------|----|---------|------|------|-------|--------|------|
| PHQ-9 | 1-factor | 148.37 | 27 | < 0.001 | 0.907 | 0.876 | 0.083 | 0.070–0.096 | 0.050 |
| PHQ-9 | 2-factor | 96.34 | 26 | < 0.001 | 0.947 | 0.926 | 0.064 | 0.051–0.078 | 0.038 |
| GAD-7 | 1-factor | 133.13 | 14 | < 0.001 | 0.935 | 0.903 | 0.114 | 0.097–0.132 | 0.039 |
| GAD-7 | 1-factor + | 33.97 | 14 | 0.001 | 0.988 | 0.979 | 0.053 | 0.033–0.075 | 0.024 |
| GAD-7 | 2-factor | 46.61 | 21 | < 0.001 | 0.982 | 0.971 | 0.062 | 0.044–0.082 | 0.032 |

\( \chi^2 \), degrees of freedom; CFI, comparative fit index; TLI, Tucker–Lewis index; RMSEA, root mean square error of approximation; 90% CI, 90% confidence interval for RMSEA; SRMR, standardized root mean square residual.

practical reasons, rather than being supported by a conceptual model (Lamela et al., 2020). More recently, and in line with our findings, studies have provided empirical support for a 2-factor model (Lamela et al., 2020) with a somatic and a cognitive/affective factor. This model corresponds with the occurrence of two clinically distinct profiles of symptoms: one with a combination of high levels of cognitive/affective symptoms and low levels of somatic symptoms, and one with a combination of high levels of both subtypes of symptoms and has been supported by neurobiological findings (Lamela et al., 2020). Distinction of these subtypes has been shown to have clinical relevance in terms of prognosis and treatment (Lamela et al., 2020).

For the GAD-7, models with the somatic items as a different group or with their residuals correlated showed an adequate fit, while a unidimensional model did not. This provides empirical support for an overarching construct reflecting symptom severity, but also supports the eventual distinction of a somatic and a cognitive/affective subcomponent. Based on several samples of American psychiatric patients, researchers have argued for a one-factor model with the same correlated residuals, reflecting a method effect (referring to the nature of the question) rather than a somatic subtype (Kertz et al., 2012; Rutter and Brown, 2017; Johnson et al., 2019). Others have argued for a two-factor model, with the same items as one factor (Beard and Björgvinsson, 2014; Moreno et al., 2019). In a sample of Japanese psychiatric patients, a higher order 2-factor model was proposed (Doi et al., 2018a). Model selection in those studies was typically based on their goodness of fit. We argue that beyond these statistical criteria, conclusions should be informed by evidence from other domains. The existence of a worry subtype and a somatic subtype has been suggested based on pathophysiologic findings and clinical manifestations (Portman et al., 2011), however, more evidence is needed regarding their clinical relevance.

Factor loadings of the GAD-7 indicators ranged from acceptable to high, except for item 6. Lower factor loadings of this item have been reported in several other studies (Zhong et al., 2015; Rutter and Brown, 2017; Johnson et al., 2019) albeit not as low as in our study. Despite this low factor loading, reliability indices did not change substantially when the item was excluded. This finding in combination with the scale’s established validation in a variety of settings supports the use of the scale.
including item 6. However, we recommend further investigation of the item’s wordings and understanding when being used in similar settings.

Our findings indicate that for both scales, although they are not fully unidimensional, the general factor is the major determinant of the variance underlying the unit-weighted total scale scores. This supports unit-weighted scores of the full scales as being a reliable reflection of depression or anxiety symptom severity. However, the Haberman procedure showed that reporting unit-weighted subscale scores does not have an added value as they are no more precise than the total scores in estimating the corresponding subdimension, except for the cognitive dimension of the PHQ-9. In addition, low OmegaHS indices suggested that unit-weighted subscale scores do not provide reliable measures of constructs independently of the general construct. In other words, the subscale scores were redundant with the total score. This is in contrast with recommendations made in recent studies (Doi et al., 2018a,b). Our study expands on their findings which were based on the goodness of fit of a confirmatory factor model, an approach that has been argued insufficient for this purpose (Reise et al., 2013).

Scalar invariance was established for all studied subgroups, which suggests a similar response and interpretation of scale items across these groups (Vandenberg and Lance, 2000). Scalar invariance also justifies comparison using SEM. The use of unit-weighted scores requires invariant indicator reliability that implies invariant residuals after having established invariance of latent factor variances (Vandenberg and Lance, 2000). The latter could not be established for education (PHQ-9 only) and gender (GAD-7 only) which does not rule out indicator reliability, however, precludes the interpretation of residual invariance as indicator reliability. Full or partial residual invariance was found for the remaining subgroups. The effect of residual invariance being partial rather than full has been poorly studied and

### TABLE 2

| Indices of fit | PHQ-9 | GAD-7 |
|----------------|-------|-------|
| $\chi^2$      | 133.530 | 27.987 |
| df            | 21     | 21    |
| $p$-value     | <0.001 | <0.001 |
| CFI           | 0.948  | 0.991 |
| TLI           | 0.910  | 0.982 |
| RMSEA         | 0.071  | 0.049 |
| 90% CI        | 0.059-0.082 | 0.028-0.071 |

Reliability indices

| OmegaT       | 0.80  | 0.84  |
| OmegaH       | 0.72  | 0.76  |
| OmegaHS somatic | 0.27  | 0.27  |
| OmegaHS cognitive | 0.22  | 0.23  |

### TABLE 3

| N   | PHQ-9 | GAD-7 |
|-----|-------|-------|
|     | $\chi^2$ | df | CFI | TLI | RMSEA | SRMR | $\chi^2$ | df | CFI | TLI | RMSEA | SRMR |
| Gender |       |     |     |     |       |     |       |     |     |     |       |     |
| Female  | 553  | 65.737 | 26 | 0.000 | 0.917 | 0.067 | 0.044 | 554  | 33.022 | 12 | 0.001 | 0.978 | 0.067 |
| Male    | 654  | 57.140 | 26 | 0.000 | 0.949 | 0.059 | 0.041 | 651  | 14.268 | 12 | 0.284 | 0.965 | 0.073 |
| Age     |       |       |     |     |       |     |       |     |     |     |       |     |
| 30–40   | 326  | 66.215 | 26 | 0.000 | 0.949 | 0.068 | 0.059 | 324  | 17.777 | 12 | 0.013 | 0.967 | 0.097 |
| 41–50   | 493  | 49.860 | 26 | 0.000 | 0.968 | 0.059 | 0.043 | 495  | 26.334 | 12 | 0.074 | 0.967 | 0.097 |
| 51–60   | 388  | 61.913 | 26 | 0.000 | 0.940 | 0.066 | 0.059 | 386  | 19.846 | 12 | 0.074 | 0.967 | 0.097 |
| Education |     |       |     |     |       |     |       |     |     |     |       |     |
| Primary | 305  | 57.584 | 26 | 0.000 | 0.940 | 0.066 | 0.059 | 305  | 25.109 | 12 | 0.014 | 0.986 | 0.097 |
| Secondary| 619  | 56.576 | 26 | 0.000 | 0.940 | 0.066 | 0.059 | 617  | 15.282 | 12 | 0.226 | 0.996 | 0.097 |
| Higher  | 283  | 39.294 | 26 | 0.000 | 0.940 | 0.066 | 0.059 | 283  | 26.647 | 12 | 0.074 | 0.967 | 0.097 |
| T2D-Status |     |       |     |     |       |     |       |     |     |     |       |     |
| At risk | 1006 | 85.341 | 26 | 0.000 | 0.940 | 0.066 | 0.059 | 1003 | 25.562 | 12 | 0.012 | 0.986 | 0.097 |
| T2D     | 201  | 40.088 | 26 | 0.000 | 0.940 | 0.066 | 0.059 | 202  | 14.962 | 12 | 0.244 | 0.996 | 0.097 |
| Time    |     |       |     |     |       |     |       |     |     |     |       |     |
| T0      | 982  | 39.152 | 26 | 0.046 | 0.930 | 0.049 | 0.042 | 981  | 21.062 | 12 | 0.049 | 0.930 | 0.049 |
| T1      | 983  | 39.152 | 26 | 0.046 | 0.930 | 0.049 | 0.042 | 983  | 39.152 | 12 | 0.049 | 0.930 | 0.049 |

df, degrees of freedom; CFI, comparative fit index; TLI, Tucker–Lewis index; RMSEA, root-mean-square error of approximation; SRMR, standardized root mean square residual; T2D, type 2 diabetes.

df, degrees of freedom; CFI, comparative fit index; TLI, Tucker–Lewis index; RMSEA, root-mean-square error of approximation; 90% CI, 90% confidence interval for RMSEA; SRMR, standardized root mean square residual; OmegaT, total omega; OmegaH, hierarchical omega; and omegaHS, omega hierarchical subscale.
TABLE 4 | Assessment of measurement invariance of the PHQ-9 and GAD-7 across demographic subgroups, over time and status of diabetes.

| PHQ-9                        | Items constrained | $\chi^2$ | df | $p$  | CFI   | RMSEA | SRMR | $\Delta \chi^2$ | $\Delta p$ | $\Delta CFI$ | $\Delta RMSEA$ | $\Delta SRMR$ |
|-----------------------------|-------------------|---------|----|------|-------|-------|------|----------------|------------|------------|----------------|---------------|
| Gender                      | Configural        | 122.124 | 52 | <0.001 | 0.945 | 0.063 | 0.039 | –              | –          | –          | –               | –             |
| Metric                      | 121.308           | 59      | <0.001 | 0.948 | 0.057 | 0.043 | 3.666 | 0.817          | 0.004      | –0.006     | 0.004           | 0.004         |
| Scalar                      | 131.714           | 66      | <0.001 | 0.948 | 0.054 | 0.044 | 8.101 | 0.324          | –0.001     | –0.003     | 0.001           | 0.001         |
| Factor variance             | 137.501           | 68      | <0.001 | 0.944 | 0.056 | 0.069 | 5.047 | 0.080          | –0.004     | 0.001      | 0.025           | 0.025         |
| Residual*                   | 191.250           | 75      | <0.001 | 0.894 | 0.072 | 0.069 | 39.371 | <0.001        | –0.054     | 0.018      | 0.018           | 0.025         |
| Partial residual* 2,6,7     | 142.568           | 72      | <0.001 | 0.941 | 0.055 | 0.052 | 11.250 | 0.081          | –0.007     | 0.001      | 0.007           | 0.007         |
| Age                         | Configural        | 176.513 | 78  | <0.001 | 0.932 | 0.072 | 0.044 | –              | –          | –          | –               | –             |
| Metric                      | 177.109           | 92      | <0.001 | 0.935 | 0.065 | 0.055 | 114.809 | 0.648          | 0.003      | –0.007     | 0.011           | 0.011         |
| Scalar                      | 198.213           | 106     | <0.001 | 0.934 | 0.061 | 0.057 | 172.130 | 0.245          | –0.001     | –0.004     | 0.002           | 0.002         |
| Factor variance             | 209.552           | 110     | <0.001 | 0.927 | 0.063 | 0.084 | 98.314  | 0.043          | –0.007     | 0.002      | 0.027           | 0.027         |
| Residual*                   | 208.579           | 124     | <0.001 | 0.928 | 0.059 | 0.069 | 21.647  | 0.248          | –0.006     | –0.002     | 0.012           | 0.012         |
| Education                   | Configural        | 156.348 | 78  | <0.001 | 0.941 | 0.064 | 0.043 | –              | –          | –          | –               | –             |
| Metric                      | 186.644           | 92      | <0.001 | 0.926 | 0.066 | 0.063 | 29.826  | 0.008          | –0.016     | 0.002      | 0.020           | 0.020         |
| Partial metric 7            | 167.316           | 90      | <0.001 | 0.940 | 0.060 | 0.052 | 13.392  | 0.341          | –0.001     | –0.004     | 0.009           | 0.009         |
| Scalar                      | 185.049           | 104     | <0.001 | 0.941 | 0.055 | 0.053 | 12.461  | 0.569          | 0.001      | –0.004     | 0.004           | 0.004         |
| Factor variance             | 218.204           | 108     | <0.001 | 0.916 | 0.065 | 0.110 | 19.521  | <0.001        | –0.025     | 0.009      | 0.057           | 0.057         |
| Residual*                   | 274.559           | 122     | <0.001 | 0.859 | 0.079 | 0.109 | 58.147  | <0.001        | –0.082     | 0.023      | 0.056           | 0.056         |
| Partial residual* 6,7       | 200.944           | 118     | <0.001 | 0.932 | 0.056 | 0.064 | 20.026  | 0.129          | –0.009     | 0.000      | 0.012           | 0.012         |
| T2D status                   | Configural        | 131.678 | 52  | <0.001 | 0.946 | 0.064 | 0.038 | –              | –          | –          | –               | –             |
| Metric                      | 130.505           | 59      | <0.001 | 0.948 | 0.060 | 0.044 | 58.489  | 0.558          | 0.001      | –0.006     | 0.006           | 0.006         |
| Scalar                      | 139.259           | 66      | <0.001 | 0.949 | 0.056 | 0.044 | 43.797  | 0.735          | 0.001      | –0.004     | 0.000           | 0.000         |
| Factor variance             | 136.175           | 68      | <0.001 | 0.951 | 0.054 | 0.049 | 0.8415  | 0.657          | 0.002      | –0.002     | 0.005           | 0.005         |
| Residual*                   | 144.056           | 75      | <0.001 | 0.945 | 0.054 | 0.050 | 11.655  | 0.233          | –0.004     | –0.001     | 0.006           | 0.006         |
| Time                        | Configural        | 173.821 | 78  | <0.001 | 0.968 | 0.050 | 0.033 | –              | –          | –          | –               | –             |
| Metric                      | 197.636           | 92      | <0.001 | 0.963 | 0.049 | 0.046 | 24.934  | 0.035          | –0.005     | –0.001     | 0.013           | 0.013         |
| Scalar                      | 247.699           | 106     | <0.001 | 0.954 | 0.051 | 0.049 | 67.865  | <0.001        | –0.009     | 0.002      | 0.003           | 0.003         |
| Factor variance             | 255.133           | 110     | <0.001 | 0.952 | 0.051 | 0.063 | 7.928   | 0.094          | –0.002     | 0.000      | 0.014           | 0.014         |
| Residual*                   | 489.852           | 124     | <0.001 | 0.845 | 0.086 | 0.102 | 127.070 | <0.001        | –0.109     | 0.035      | 0.054           | 0.054         |
| Partial residual* 1,2,5,6,7,8| 144.056           | 112     | <0.001 | 0.945 | 0.054 | 0.050 | 15.264  | 0.018          | 0.009      | –0.003     | –0.003          | –0.003        |

(Continued)
| GAD-7 Items constrained | \( \chi^2 \) | df | \( \rho \) | CFI | RMSEA | SRMR | \( \Delta \chi^2 \) | \( \Delta \rho \) | \( \Delta \) CFI | \( \Delta \) RMSEA | \( \Delta \) SRMR |
|------------------------|--------|-----|------|-----|------|------|-----------|--------|----------|----------|-----------|
| **Gender**             |        |     |      |     |      |      |           |         |          |          |           |
| Configural             | 45.472 | 24  | 0.005| 0.987| 0.053| 0.025|          | -       | -        | -        | -        |
| Metric                 | 46.633 | 30  | 0.027| 0.990| 0.043| 0.033| 27.551   | 0.839   | 0.002    | -0.010   | 0.008     |
| Scalar                 | 64.176 | 36  | 0.003| 0.984| 0.049| 0.040| 230.732  | < 0.001 | -0.006   | 0.006    | 0.008     |
| Factor variance        | 88.158 | 37  | <0.001| 0.970| 0.065| 0.136| 162.348  | < 0.001 | -0.014   | 0.017    | 0.095     |
| Residual*              | 131.510| 43  | <0.001| 0.941| 0.085| 0.073| 45.674   | < 0.001 | -0.002   | 0.036    | 0.032     |
| Partial residual*      | 66.057 | 41  | 0.013| 0.986| 0.042| 0.039| 15.748   | 0.008   | -0.010   | 0.010    | 0.009     |
| **Age**                |        |     |      |     |      |      |           |         |          |          |           |
| Configural             | 63.391 | 36  | 0.003| 0.986| 0.057| 0.027|          | -       | -        | -        | -        |
| Metric                 | 75.737 | 48  | 0.007| 0.985| 0.051| 0.049| 134.609  | 0.336   | -0.001   | -0.006   | 0.022     |
| Scalar                 | 95.195 | 60  | 0.003| 0.983| 0.049| 0.052| 198.170  | 0.071   | -0.002   | -0.002   | 0.003     |
| Factor variance        | 95.908 | 62  | 0.004| 0.983| 0.048| 0.064| 18.565   | 0.395   | 0.000    | -0.001   | 0.011     |
| Residual*              | 101.703| 74  | 0.018| 0.985| 0.042| 0.053| 11.880   | 0.616   | 0.002    | -0.007   | 0.001     |
| **Education**          |        |     |      |     |      |      |           |         |          |          |           |
| Configural             | 66.657 | 36  | 0.001| 0.984| 0.060| 0.026|          | -       | -        | -        | -        |
| Metric                 | 83.912 | 48  | 0.001| 0.980| 0.059| 0.050| 183.622  | 0.105   | -0.005   | -0.001   | 0.024     |
| Scalar                 | 110.261| 60  | <0.001| 0.974| 0.060| 0.053| 301.806  | 0.000   | -0.005   | 0.000    | 0.003     |
| Factor variance        | 120.791| 62  | <0.001| 0.969| 0.064| 0.100| 75.149   | 0.023   | -0.005   | 0.005    | 0.016     |
| Residual*              | 123.384| 74  | <0.001| 0.971| 0.057| 0.065| 17.481   | 0.231   | -0.003   | -0.003   | 0.011     |
| **T2D status**         |        |     |      |     |      |      |           |         |          |          |           |
| Configural             | 40.073 | 24  | 0.021| 0.991| 0.046| 0.023|          | -       | -        | -        | -        |
| Metric                 | 42.627 | 30  | 0.063| 0.992| 0.038| 0.029| 42.037   | 0.649   | 0.001    | -0.008   | 0.006     |
| Scalar                 | 50.677 | 36  | 0.053| 0.992| 0.036| 0.030| 76.128   | 0.268   | 0.000    | -0.002   | 0.001     |
| Factor variance        | 50.969 | 37  | 0.063| 0.992| 0.035| 0.038| 0.693    | 0.406   | 0.000    | -0.001   | 0.008     |
| Residual*              | 66.057 | 43  | 0.013| 0.986| 0.042| 0.039| 14.072   | 0.050   | -0.006   | 0.007    | 0.008     |
| **Time**               |        |     |      |     |      |      |           |         |          |          |           |
| Configural             | 84.234 | 36  | <0.001| 0.988| 0.051| 0.022|          | -       | -        | -        | -        |
| Metric                 | 94.655 | 48  | <0.001| 0.988| 0.044| 0.032| 11.560   | 0.482   | 0.000    | -0.007   | 0.010     |
| Scalar                 | 135.387| 60  | <0.001| 0.982| 0.048| 0.038| 54.319   | < 0.001 | -0.006   | 0.003    | 0.005     |
| Factor variance        | 149.413| 62  | <0.001| 0.979| 0.051| 0.074| 9.799    | 0.007   | -0.003   | 0.004    | 0.006     |
| Residual*              | 210.355| 74  | <0.001| 0.964| 0.061| 0.058| 61.287   | < 0.001 | -0.018   | 0.013    | 0.021     |
| Partial residual*      | 4,5    | 66.057| 70  | 0.013| 0.986| 0.042| 0.039    | 14.077  | 0.170    | -0.001   | -0.002    | 0.003     |

*df*, degrees of freedom; \( \Delta \chi^2 \), change in \( \chi^2 \); CFI, comparative fit index; RMSEA, root-mean-square error of approximation; SRMR, standardized root mean square residual; \( \Delta \) CFI, change in CFI; \( \Delta \) RMSEA, change in RMSEA; \( \Delta \) SRMR, change in SRMR; T2D, type two diabetes. *Compared to the model with constrained loadings and intercepts (i.e., the scalar invariant model).
### TABLE 5 | Descriptive statistics of the PHQ-9 and GAD-7 per category and comparison of effect sizes between unit-weighted and latent means.

| Category | Subgroup   | N   | Unit-weighted scores | Unit-weighted score differences<sup>a</sup> | Latent mean differences<sup>b</sup> |
|----------|------------|-----|----------------------|-------------------------------------------|-------------------------------------|
|          |            |     | Mean     | SD       | SES     | P-value | Mean     | SD       | SES     | P-value |
| PHQ-9    | Total      | 1207 | 3.71     | 3.72    |         |         | 0.417    | <0.001   | −0.429  | <0.001  |
|          | Gender     |      |          |         |         |         |          |          |         |         |
|          | Female     | 553  | 4.66     | 3.93    | −0.417  | <0.001  | −0.429  | <0.001  |         |         |
|          | Male       | 654  | 2.91     | 3.34    | 0.076   | 0.203   | 0.094   | 0.228   |         |         |
|          | Age        |      |          |         |         |         |          |          |         |         |
|          | 30–40      | 326  | 3.24     | 3.27    | 0.152   | 0.016   | 0.194   | 0.001   |         |         |
|          | 41–50      | 493  | 3.66     | 3.78    |         |         |         |         |         |         |
|          | 51–60      | 388  | 4.16     | 3.97    |         |         |         |         |         |         |
|          | Education  |      |          |         |         |         |          |          |         |         |
|          | Primary    | 305  | 4.83     | 4.34    |         |         |         |         |         |         |
|          | Secondary  | 619  | 3.72     | 3.67    | −0.195  | 0.001   | −0.259  | <0.001  |         |         |
|          | Higher     | 283  | 2.45     | 2.56    | −0.446  | <0.001  | −0.602  | <0.001  |         |         |
|          | Status     |      |          |         |         |         |          |          |         |         |
|          | At risk    | 1006 | 3.77     | 3.77    |         |         |         |         |         |         |
|          | T2D        | 201  | 3.4      | 3.47    | −0.046  | 0.477   | −0.085  | 0.508   |         |         |
|          | Time       |      |          |         |         |         |          |          |         |         |
|          | T0         | 1006 | 3.77     | 3.77    |         |         |         |         |         |         |
|          | T1         | 982  | 3.09     | 3.49    | −0.168  | <0.001  | −0.151  | <0.001  |         |         |
|          | T2         | 963  | 2.68     | 3.19    | −0.255  | <0.001  | −0.262  | <0.001  |         |         |
| GAD-7    | Total      | 1205 | 3.16     | 3.33    |         |         |         |         |         |         |
|          | Gender     |      |          |         |         |         |          |          |         |         |
|          | Female     | 554  | 3.9      | 3.72    | −0.299  | <0.001  | −0.603  | <0.001  |         |         |
|          | Male       | 651  | 2.52     | 2.8     | 0.002   | 0.973   | 0.130   | 0.101   |         |         |
|          | Age        |      |          |         |         |         |          |          |         |         |
|          | 30–40      | 324  | 3.09     | 3.19    | −0.039  | 0.471   | 0.042   | 0.583   |         |         |
|          | 41–50      | 495  | 3.09     | 3.26    |         |         |         |         |         |         |
|          | 51–60      | 386  | 3.3      | 3.53    |         |         |         |         |         |         |
|          | Education  |      |          |         |         |         |          |          |         |         |
|          | Primary    | 305  | 3.91     | 3.81    |         |         |         |         |         |         |
|          | Secondary  | 617  | 3.1      | 3.2     | −0.144  | 0.007   | −0.292  | <0.001  |         |         |
|          | Higher     | 283  | 2.47     | 2.86    | −0.328  | <0.001  | −0.541  | <0.001  |         |         |
|          | Status     |      |          |         |         |         |          |          |         |         |
|          | At risk    | 1003 | 3.21     | 3.37    | −0.065  | 0.265   | −0.138  | 0.181   |         |         |
|          | T2D        | 202  | 2.88     | 3.11    |         |         |         |         |         |         |
|          | Time       |      |          |         |         |         |          |          |         |         |
|          | T0         | 1003 | 3.21     | 3.37    |         |         |         |         |         |         |
|          | T1         | 981  | 2.62     | 2.94    | −0.132  | <0.001  | −0.180  | <0.001  |         |         |
|          | T2         | 963  | 2.42     | 2.81    | −0.172  | <0.001  | −0.256  | <0.001  |         |         |

<sup>a</sup>Based on robust regression with MM-estimation.

<sup>b</sup>Based on multigroup structural equation modeling using a second order model for the PHQ-9 to allow for comparison of the general factor latent mean. SD, standard deviation; SES, standardized effect size; T2D, type two diabetes. Interpretation of the SES corresponds to Cohen’s d (conventionally: d = 0.20, 0.50, and 0.80 for small, medium, and large effects, respectively). Empty rows indicate the reference category.

Currently, no agreement exists on the required number of items to obtain an acceptable level of invariance (Putnick and Bornstein, 2016). Some experts argue that two items is sufficient (Putnick and Bornstein, 2016).

The differences in effect size between subgroups turned out to be systematically higher using latent means compared to unit-weighted scores. This is expected since random error is ignored when comparing latent means. However, the difference in effect size was much smaller for the PHQ-9. This can be explained by the use of a second-order model for this scale representing the common effect of the subfactors. Unit-weighted scores and the unidimensional model for the GAD-7 did not distinguish between general and subfactor effects, resulting in relatively higher effect sizes. We conclude that our findings support the use of unit-weighted scores to detect differences of a reasonable effect size across different subgroups. However, we do recommend the use of SEM whenever possible and certainly if a precise estimate is at stake.

Our findings regarding invariance of the PHQ-9 are in line with previous studies conducted in western settings assessing residual invariance for gender and age (Lamela et al., 2020). Those studies also showed residual invariance for education. Full or partial (after freeing 1 item) residual invariance over different measurement occasions was found among primary care patients (González-Blanch et al., 2018) and COPD patients (Schuler et al., 2018). However, time intervals were smaller (3 months) in those studies. Our findings on invariance of the GAD-7 are in line with previous studies conducted in western settings reporting scalar invariance across gender and age (Hinz et al., 2017; Rutter and Brown, 2017); residual invariance was not tested in these studies. A study using a digitalized version of the GAD-7 found residual invariance across gender, age, education and different measurement occasions (Moreno et al., 2019). For both scales,
our study is the first to establish invariance between people at risk of T2D and with T2D.

Higher severity scores of depression and anxiety in women and people with lower education are in line with national and global trends (Bjelland et al., 2008; Grover et al., 2010; Hinz et al., 2017; Sagar et al., 2020). Prevalence of both disorders followed the same trends. Higher severity scores of depression in the older subgroup and the status quo of anxiety severity across age groups were compatible with national trends (Sagar et al., 2020).

Our findings are in support of the use of both scales to measure symptom severity among individuals. However, health workers need to take into account that part of the scores does not reflect symptom severity, but random error, and this part differs across individuals. Indicator reliability across certain categories (e.g., gender and education) may also slightly vary since full residual invariance could not be established. Moreover, the lack of full residual invariance over time suggests that scores' reliability may vary over time (i.e., years) which should be taken into account when using the scales for patient follow-up. We conclude that, while scores of an individual may give an idea of symptom severity, further examination is essential for an accurate diagnosis.

Strengths and Limitations
A major strength of our study was the use of a large sample size which was generally representative of Kerala's general population in terms of age structure, education, occupation, marital status, and household size at the time of study enrollment (Office of the Registrar General and Census Commissioner of India, 2011). However, the gender ratio was lower than the state's average and the study population was restricted to people aged 30–60 years, with no history of chronic conditions yet at risk of diabetes. Generalizing of our findings to other states of India also warrants caution since Kerala's literacy rate and health indicators are better than most Indian states. However, since the scales showed invariance for age, gender, education and status of diabetes, we assume that differences in these characteristics may not have a substantial influence. We conducted an extensive assessment of reliability based on bifactor modeling and compared differences between unit-weighted scores and latent means. To our knowledge, these techniques have not been applied previously to these scales. A weakness of the study is that we did not examine criterion validity nor did we evaluate invariance for ethnicity and socioeconomic status. Another limitation was the use of a maximum likelihood estimator for the parameter estimation of the factor analyses, which likely contributed to a negative bias in parameter estimates because of the non-normal data (Li, 2016). Weighted least square estimation would normally be preferable as it has been shown more accurate if the normality assumption is violated (Li, 2016). However, the use of tetrachoric or polychoric correlations has been shown to overestimate reliability (Revelle and Condon, 2018). A model-based reliability index for categorical data has been proposed, but evidence about its performance is lacking (Green and Yang, 2009). Since assessment of reliability was the primary objective of this study, we therefore opted for a maximum likelihood estimator. As expected, sensitivity analysis with diagonally weighted least squares resulted in higher factor loadings and improved model fit. Despite this, the difference was not large enough to alter our conclusions. Using the formula proposed by Green and Yang resulted in similar estimates of reliability: 0.73 for PHQ-9 and 0.79 for GAD-7. Finally, it would be interesting for future studies to study invariance of response patterns across global regions.

In conclusion, our findings support the existence of a somatic and a cognitive subtype of depression and to a lesser extent for anxiety in a rural Indian population. Unit-weighted scores of the full scales can be used at individual and population level, reflecting a single construct of symptom severity. However, one needs to take into account that part of the score corresponds to error. Scoring of the subscales is redundant. Both scales showed to be stable across demographic subgroups and over time, which supports their use and allows for meaningful comparison across tested subgroups in the Indian context. For both scales, psychometric properties are comparable to studies in western settings.

DATA AVAILABILITY STATEMENT
The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT
The studies involving human participants were reviewed and approved by the Institutional Ethics Committee of the Sree Chitra Tirunal Institute for Medical Sciences and Technology, Trivandrum, Kerala, and by the Human Research Ethics Committees of Monash University, Australia and the University of Melbourne, Australia. The study was also approved by the Health Ministry Screening Committee of the Government of India. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS
PA, TS, BO, JD, and EW played a major role in the conception of the study. PA, TS, KT, BO, and EW contributed to the design of the study. JD drafted the manuscript and analyzed and interpreted the data. PA, Lj, AD, TS, TH, KT, BO, and EW critically revised the manuscript for important intellectual content, and read and approved the final version. All authors contributed to the article and approved the submitted version.

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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