

16

Abstract

17 **Background:** Depression is one of the most prevalent mental health conditions in the world.
18 However, the heterogeneity of depression has presented obstacles for research concerning
19 disease mechanisms, treatment indication, and personalization. So far, depression heterogeneity
20 research has mainly used latent variable modeling, assuming a latent cause, that overlooks the
21 possibility that symptoms might interact and reinforce each other. The current study used
22 network analysis to analyze and compare profiles of depressive symptoms present in community
23 samples, considering the relationship between symptoms.

24 **Methods:** Cross-sectional measures of depression using the Patient Health Questionnaire-9
25 (PHQ-9) were collected from community samples using data from participants scoring above a
26 clinical threshold of ≥ 10 points (N=2,023; 73.9% female; mean age 49.87, SD= 17.40). Data
27 analysis followed three steps. First, a profiling algorithm was implemented to identify all
28 possible symptom profiles by dichotomizing each PHQ-9 item. Second, the most prevalent
29 symptom profiles were identified in the sample. Third, network analysis for the most prevalent
30 symptom profiles was carried out to identify the centrality and covariance of symptoms.

31 **Results:** Of 382 theoretically possible depression profiles, only 167 were present in the sample.
32 Furthermore, 55.6% of the symptom profiles present in the sample were represented by only
33 eight profiles. Network analysis showed that the network and symptoms relationship varied
34 across the profiles.

35 **Conclusions:** Findings indicate that the vast number of theoretical possible ways to meet the
36 criteria for major depressive disorder is significantly reduced in empirical samples, and that the
37 most common profiles of symptoms have different networks and connectivity patterns. Scientific

38 and clinical consequences of these findings are discussed in the context of the limitations of this
39 study.

40 *Keywords:* Network Analysis, Depression Heterogeneity, Depression Profiles

41

Introduction42 *Why is depression a public health problem?*

43 Depression is the most prevalent mental health problem in the world affecting 4.7% of
44 the global population [1]. It has been classified as a public health problem due to its impact on
45 quality of life, work productivity, and mortality risk [2]. Despite global efforts to understand and
46 treat depression, its incidence has actually increased by 49% between 1990 and 2017 [3].
47 Currently, it is the third leading cause of disease burden and the single highest contributing
48 factor to global disability [3,4]. The impact of depression is not only felt by individuals, but also
49 by their families and communities, who suffer a direct cost related to treatment and an indirect
50 cost linked to an individual's reduced functional capacity [5,6]. Studies estimate that, when
51 diagnosed, depression could cost \$6,200 per person per year [7], while undiagnosed and
52 untreated depression contributes to an even more significant burden of illness, increasing
53 personal and societal costs [8]. Indeed, longer periods of undiagnosed and untreated depression
54 lead to negative outcomes including poorer treatment response and lower remission rates [9],
55 more severe cognitive impairment [10], and overall to poorer illness trajectories [11].

56

57 *What does heterogeneity in depression mean?*

58 Ever since the publication of DSM-III forty years ago, the field of healthcare has mainly
59 conceived depression as a homogeneous, distinct, and robust diagnostic category, outlined
60 broadly in the polythetic system of the DSM as Major Depressive Disorder (MDD) [12]. The
61 DSM's polythetic system masks a significant amount of syndromic heterogeneity, allowing for
62 multiple combinations of symptoms to exist under the same diagnostic label [13]. As a

63 consequence, the diagnostic criteria of MDD has led to the classification of people with only
64 some or even no symptoms in common into the same broad category, ignoring the specific
65 presentation of their symptoms and the interactions between specific symptoms [14–16].

66

67 *Consequences of heterogeneity in depression on treatment outcomes*

68 Failure to consider heterogeneity in depression has impacted the understanding of
69 etiological mechanisms and their physiological correlates [17–20] and it has also limited the
70 effectiveness of treatments [21]. Thus, it is highly important that research should consider the
71 heterogeneity of MDD in order to better address etiological processes and to implement smarter
72 and personalized treatment strategies [16].

73 It is estimated that nearly 85% of people who recover from MDD suffer a second episode
74 within 15 years, and that each additional MDD episode increases the risk of relapse by 18%
75 [22]. In addition, for 30% of patients diagnosed with MDD, symptoms do not remit despite
76 varied treatment attempts [2], with sleep problems and fatigue being the most prevalent residual
77 symptoms [23]. This highlights that patients will not respond similarly to different treatments
78 for MDD [24,25]. Even though clinicians typically adjust treatments to their specific patients,
79 often guidelines recommend treatments packages that are delivered to the ‘average depressed
80 patient’ and insufficient research has considered what are the specific modifications that should
81 be implemented to optimize a treatment for a particular subtype of depression. Thus, treatments
82 may yield sub-optimal effects, whereas parsing out heterogeneity in depression could enable the
83 design of evidence-based personalization strategies for treatments, thus leading to possible
84 improved patient outcomes.

85

86 *What are we missing by not looking at symptom-level heterogeneity?*

87 Research has typically approached depression as a common cause for diverse symptoms
88 and assumed that these symptoms are independent and have equal importance [14,26,27].
89 However, such a strategy has paid less attention to the interaction and mutual reinforcement
90 between symptoms [28]. This is a problem, considering that researchers have been attempting to
91 find associations between different symptoms of depression and distinct risk factors [29,30],
92 different gene polymorphisms [31], and different responses to treatment [32,33]. Moreover,
93 different symptoms have been associated with varying impacts on disability, with depressed
94 mood and concentration problems being the most disabling symptoms [34]. This is consistent
95 with research showing that patients who receive their optimal treatment (considering their
96 specific symptoms and personal characteristics) had clinically significant improvements in
97 depression [e.g., 21,35,36].

98 An examination of symptom level heterogeneity in depression may also be crucial in
99 improving our understanding of differential developmental pathways towards psychopathology
100 from the perspective of equifinality and multifinality [37]. Indeed, by using a homogeneous
101 conceptualization of depression, different pathways to illness may be masked and thus, relevant
102 opportunities for prevention and personalization lost. Heterogeneity research in depression can
103 move the field towards a more person-centered approach that recognizes the relevance of
104 different developmental pathways to illness that may be related to or represented by different
105 profiles of depression [38].

106 Until now, depression has been studied through theoretical and empirical approaches that
107 have supplied evidence to its heterogeneity, identifying profiles of symptoms that may help map
108 out heterogeneity [2,16]. However, although empirical research has found profiles related to the
109 composition as well as severity of symptom profiles [e.g., 13, 39], it does not consider the
110 relation *between* symptoms within emerging profiles. Furthermore, studies show that not all
111 theoretically possible profiles of symptoms are actually present in clinical samples [13,39]. Still,
112 these studies have focused on the presence and prevalence of different profiles, leaving aside
113 how the symptoms are related to each other as an interrelated system. There are also studies that
114 are focused on seeing the interaction between depressive symptoms using network analysis and
115 other analytic strategies, but they usually analyze the depressive symptoms on total samples
116 without considering different profiles and interrelated networks between profiles [40].

117 As a result, it is not known which symptoms are present in each profile, how they are
118 related, or what the structure of the network of symptoms is like. In addition, we do not
119 understand how the empirical frequency of theoretical profiles differs when considering
120 community samples that include both help-seeking and non help-seeking individuals. The
121 current study is the first, to our knowledge, to examine the network structure and interactions
122 between symptoms on different symptom profiles of depression.

123

124

Methods

125 Participants

126 The study used secondary data derived from three community studies with nationally
127 representative samples: (1) the Chilean Longitudinal Social Survey (ELSOC); (2) the

128 Longitudinal Study of Intercultural Relations (ELRI); and (3) the Social Protection Survey
129 (EPS). These three studies were carried out between 2016 and 2020 and used multi-stage,
130 stratified, and probabilistic sampling. The inclusion criteria for the sampling of these studies
131 focused on female and male residents in urban areas, aged 18 to 99, and located in 13 different
132 (blinded for review). All three studies used the Patient Health Questionnaire (PHQ-9) to
133 measure depression symptoms. Only participants with clinically significant depressive
134 symptoms ($\text{PHQ-9} \geq 10$) were included in the present study. Of a total sample of 13,367
135 participants, 2,023 (15.13%) had a PHQ-9 score of 10 or above.

136 **Measures and data sources**

137 The Spanish-language version version of the PHQ-9 was used to measure depressive
138 symptoms [PHQ-9, 41]. It is a nine-item scale in which each item represents a DSM symptom
139 criterion. Participants are asked to report whether they have experienced the symptom in the last
140 two weeks on a Likert scale ranging from 0 to 3, where 0 is "not at all," and 3 is "almost every
141 day," resulting in a total score ranging from 0 to 27 points [42]. The PHQ-9 was designed to
142 screen for depression and has shown that scores ≥ 10 have a sensitivity of 88% and specificity of
143 88% for major depressive disorder compared to semi-structured interviews [43]. A diagnostic
144 cut-off of ≥ 10 is recommended for the detection of MDD, the criteria for classifying severity
145 levels of depression according the PHQ-9 are "moderate" for scores of 14 or below, "moderately
146 severe" for scores ranging between 15 and 19, and "severe depression" for scores of 20 or above.
147 [43].

148 **Statistical Analysis**

149 Descriptive statistics were generated for the total sample. To test for sex differences in
150 total PHQ-9 means, a t-test for independent samples was used.

151

152 All possible symptom profiles were identified for PHQ-9 scores equal or higher than 10
153 points (i.e., clinical sample) using an algorithm of combinatorial optimization. This was
154 calculated using the formula ${}^n C_r = \frac{n!}{r!(n-r)!}$ (for formula estimation see supplementary material),
155 that allows calculation of the number of ways of selecting r objects out of n different objects
156 [44]. The estimation resulted in 382 possible symptom combinations.

157

158 *Theoretical symptom profile analysis*

159 All possible symptom combinations were analyzed for the PHQ-9 using a profiling
160 algorithm developed by Banyard et al. [45]. In this algorithm, individual item responses to the
161 PHQ-9 were dichotomized, and coded as either "1" if a symptom was present (a score of 1-3) or
162 "0" if a symptom was absent (a score of 0). Using conditionals, each individual response was
163 matched to their corresponding profile (for details, see supplementary material). Different
164 theoretical profiles of depressive symptomatology could thus be constructed yielding a score of
165 10 or above 382 possible theoretical profiles, for details see supplementary material. Each
166 theoretical profile was assigned a number and its relative frequency was determined using
167 patient-level data, using a syntax that matches each participant's PHQ-9 responses to each of the
168 possible 382 theoretical symptom profile combinations. This is a method that prioritizes the
169 identification of *qualitatively distinctive* symptom profiles by emphasizing the absence-presence
170 of symptoms rather than emphasizing *quantitative differences* in their relative scores across each

171 Likert scale. This method was selected to maximize the probability of identifying qualitatively
172 different profiles, since prior research using continuous Likert-scale scores to identify latent
173 classes consistently show that such a method mainly parses cases into quantitatively distinctive
174 subgroups of cases with low-moderate-severe depression [e.g., see 27,46].

175

176

177 *Network analysis*

178 Network analysis was used to examine the most prevalent profiles within the relationship
179 between symptoms, so that within a particular network, each *node* represents a PHQ-9 item (i.e.,
180 a depression symptom) and each *edge* represents the partial correlation between two symptoms.
181 Network estimation was conducted using Pairwise Markov Random Fields to calculate a
182 nondirected weighted network structure, and a Gaussian Graphical Model to estimate networks
183 with continuous data variables. By using the continuous item scores as inputs into the network
184 model, we were able to comprehensively identify *qualitatively distinctive* profiles (through the
185 prior step of analysis) while examining their *quantitative distinctive* network structures using the
186 full range of Likert scale responses.

187 The Fruchterman-Reingold algorithm was used to calculate the optimal layout of the
188 networks and to visualize more strongly connected nodes [47]. False-positive relations were
189 excluded by using the 'graphical least absolute shrinkage and selection operator' (GLASSO)
190 method, a statistical regularization technique, to increase the specificity of the network [48]. Due
191 to recent developments in network analysis discussing the use of regularized versus non-
192 regularized techniques for the estimation of psychopathology networks [49–51], both types of
193 analysis were conducted, and results are presented in supplementary materials. Finally, the

194 extended Bayesian information criterion (EBIC) was used to select the best-fitting model
195 (hyperparameters $\gamma = 0.5$ and $\lambda = 0.01$).

196 Strength centrality indexes were calculated for each network. This measure takes the
197 sum of all absolute edge weights to which a node is directly connected [52]. To estimate the
198 network stability, and considering the sample size for each depression symptom profile, a non-
199 parametric bootstrapping procedure was used with 1,000 sample simulations providing results
200 related to the edge-weight accuracy on each network [53,54]. A case-dropping subset bootstrap
201 was performed for the estimation of the centrality stability, which estimates a correlation
202 stability coefficient (CS-coefficient) representing the maximum proportion of the sample that
203 can be dropped and maintaining a 95% probability of a correlation between the original
204 centrality indices and the centrality metric equal or higher to 0.7. Thus, the centrality metric is
205 considered interpretable when the CS-coefficient is above 0.25 [53]. All the analyses were
206 performed using '*qgraph*' (55,56) and '*bootnet*' packages [53,54] on R studio version 4.0.0 [57].

207

208 **Results**

209 **Sample characteristics**

210 The total sample included PHQ-9 data from $N=2,023$ participants that had clinically
211 significant depression symptoms ($\text{PHQ-9} \geq 10$). Overall, 73.9% ($n = 1,495$) of participants were
212 female, the mean age of the total sample was 49.87 ($SD = 17.40$) years, and the mean PHQ-9
213 score was 14.7 ($SD = 4.36$). Approximately 57.7% of participants would be considered
214 moderately depressed ($n = 1,168$), 26.2% ($n = 531$) had moderately severe scores, and 16.0% (n

215 = 324) had severe depression. Supplementary Table 1 provides further sample characteristics
216 and details on item-level means and frequencies.

217

218 There were no statistically significant differences between female and male participants
219 regarding their mean depression severity scores ($t_{(2021)} = -1.51, p = .13$). In total, 35.5% of
220 participants reported having received a depression diagnosis, and 84.4% of participants who
221 received a diagnosis were women. Approximately 32% (649) reported having previously
222 received or currently were receiving treatment for depression at the time of the assessment.
223 Among participants with a past or current history of treatment, 83.6% (551) were female.

224

225 **Symptom profiles**

226 Of the 382 theoretically possible profiles of depressive symptoms operationalized by the
227 profiling algorithm for scores of 10 and above on the PHQ-9, 167 were actually present in the
228 sample. However, more than half of all cases present in the sample (55.6%) were accounted by
229 only 8 symptom profiles. Of all 167 profiles present in the sample, the most frequent was profile
230 1 ($n=510$), a “typical” depression profile that includes all 9 symptoms of depression measured
231 by the PHQ-9 (all 9 items are positive) and had a frequency of 25.2% of the sample. The mean
232 age for profile 1 was 49.87 ($SD= 17.40$) and the mean PHQ-9 score within this profile was 18.6
233 ($SD = 4.85$).

234 The second most frequent profile was profile 2 ($n=205$), a profile that includes all
235 symptoms, except for suicidal ideation (item 9), which accounted for 10.1% of cases present in
236 the sample. The mean age for profile 2 was 45.68 ($SD= 17.41$) and the mean PHQ-9 score was
237 14.88 ($SD = 4.09$).

259 The network of profile 1 is visualized in Figure 1 and shows a strong positive connection
260 between low mood and anhedonia ($pr = 0.30$) and also between sleep problems and low energy
261 ($pr = 0.26$). Also, there is a community of tightly interrelated symptoms including suicidal
262 ideation- concentration problems ($pr = 0.24$), suicidal ideation- changes in psychomotor
263 functioning (agitation or retardation) ($pr = 0.20$), concentration problems- changes in
264 psychomotor functioning ($pr = 0.20$).

265 The nodes with the highest strength centrality in profile 1 were: low mood, low energy
266 and concentration problems. The least central nodes in terms of strength centrality were
267 anhedonia and sleep problems. Node strength centrality demonstrated an interpretable level of
268 stability ($CS (cor = 0.7) = 0.36$). Details of the centrality stability test are shown in
269 Supplementary Figures 1 and 2.

270

271 *Profile 2: Typical depression without suicidal ideation*

272 This profile included all typical depression symptoms except for suicidal ideation.

273

274 Figure 2. Network of symptoms and centrality plot for profile 2.

275

276 **INSERT-FIGURE-2**

277

278

279 Note: The centrality plot shows standardized strength indices.

280

281 The network of profile 2 is visualized in Figure 2 and shows a strong positive connection
282 between low mood and anhedonia ($pr = 0.27$), low mood and low energy ($pr = 0.26$), and low
283 mood and worthlessness ($pr = 0.18$). The nodes with the highest strength centrality were low
284 mood, low energy and worthlessness. The least central nodes in terms of strength centrality were
285 psychomotor functioning and anhedonia. These data must be interpreted with caution because
286 node strength centrality demonstrated low stability ($CS(\text{cor} = 0.7) = 0.12$). See details of the
287 centrality stability test in Supplementary Figures 4 and 5.

288

289 *Profile 3: All depressive symptoms except for psychomotor functioning and suicidal ideation*

290

291 This profile includes all PHQ-9 symptoms except for suicidal ideation and changes
292 related to psychomotor functioning.

293

294 Figure 3. Network of symptoms and centrality plot for profile 3.

295

296

INSERT-FIGURE-3

297

298

299 Note: The centrality plot shows standardized strength indices.

300 The network of profile 3 is visualized in Figure 3. Profile 3 was the third most frequent
301 in the sample. Due to the small sample size ($n = 81$) this network was estimated using a threshold
302 of ≤ 0.05 instead of applying the GLASSO method (which did not converge in this subgroup).

303 The network shows a strong positive connection between low mood and low energy ($pr = 0.37$),

304 low mood and anhedonia ($pr = 0.32$), sleep problems and anhedonia ($pr = 0.23$), sleep problems
305 and low energy ($pr = 0.22$), appetite and worthlessness ($pr = 0.23$).

306 The nodes with the highest strength centrality were low energy, low mood, and
307 worthlessness. In contrast, the nodes with the least strength centrality were sleep problems and
308 appetite changes. However, these data must be interpreted with caution because node strength
309 centrality demonstrated low stability related to the sample size ($CS(\text{cor} = 0.7) = 0.21$). For
310 details on the centrality stability test and accuracy (see Supplementary Figures 7 and 8).

311

312 Results indicated that node centrality varied across the most frequent profiles of
313 depression (see Figure 4 for a comparison). Consistently, the most central symptoms were low
314 mood and low energy, and the less central symptoms were anhedonia, change in the
315 psychomotor functioning, and appetite changes.

316

317 Figure 4. Strength centrality rankings indices for the three most prevalent profiles.

318

319 **INSERT-FIGURE-4**

320 Note: Numbers indicate Strength centrality rankings. Profile 1: all of the typical symptoms of
321 depression; Profile 2: typical depression without suicidal ideation; Profile 3: All depressive
322 symptoms except for psychomotor functioning and suicidal ideation. Figure adapted with
323 permission from Malgaroli et al. [40] and license provided by Elsevier.

324

325

Conclusions

326

327 The present study identified different depressive symptom profiles and examined their
328 network structure using PHQ-9 data from participants with clinically relevant depressive
329 symptoms drawn from three community samples. Results show that 167 of the 382 theoretically
330 possible symptom combinations were present in the sample, and 55.6% of all profiles were
331 accounted for by only 8 profiles. The most frequent symptom profile included all typical
332 symptoms of depression measured by the PHQ-9 (25.2%). These results are consistent with
333 studies that applied a similar approach using patient-level data, which show that many cases
334 display similar symptom profiles. For example, Zimmerman et al. [13] similarly found in a
335 community sample that out of the 227 symptom combinations calculated using semi-structured
336 interviews (SCID-I), just 170 were empirically observed and concluded that nine combination
337 profiles accounted for the depression symptoms of 40% of patients. These findings align with
338 those of Park et al. [39], who identified in a clinical sample 119 symptom combinations within
339 their sample. Both studies, using a different approach from the one used in the present study,
340 concluded that combinatorial patterns with all nine symptoms of depression were the most
341 prevalent in samples from the USA and South Korea [13,39]. Overall, the extent of diagnostic
342 heterogeneity observed empirically within clinical and community-based samples is lower than
343 has been previously suggested based on theoretical arguments [e.g., 16].

344

345 The three most prevalent profiles showed similar mean levels of overall symptom
346 severity on the PHQ-9 total score. However, there were evident differences in their centrality
347 indices and in the interrelations between symptoms. This is clinically relevant, considering that
348 one of these profiles shows suicidal ideation and is rated with the same severity as the other

349 profiles, supporting the idea that looking at total scores in scales omits important qualitative
350 differences between symptoms concerning their hierarchy and clinical relevance [28,58].

351

352 Regarding the relationship between symptoms, there are several differences between the
353 profiles related to the centrality indices and the connection between them. The most common
354 profile, namely profile 1, showed a strong relationship between concentration problems, suicidal
355 ideation, and psychomotor functioning, with concentration problems constituting a rather strong
356 node within this profile. This is different to those profiles that do not include suicidal ideation.
357 These results are in line with those reported in two meta-analyses that found an association
358 between the attentional process and suicidal spectrum behaviors [59,60] . Thus, this profile could
359 be relevant in identify vulnerable people in the population because it has been highlighted that
360 sad mood and concentration problems, the two most central symptoms for this profile, are the
361 most disabling symptoms of depression [34].

362

363 Another difference between the profiles is present in profile 2, which shows a strong
364 connection between low mood, low energy, and self-perception. In this profile, with all of the
365 symptoms except for suicidal ideation, worthlessness takes a key role in comparison to profile 1
366 which includes all of the symptoms. On the other hand, in profile 3 worthlessness is strongly
367 related to changes in appetite, which is unique to this profile. Profile 3 is characterized by the
368 centrality of low energy which is different from profiles 1 and 2, where low mood is the most
369 central symptom.

370

371 In the most frequent profiles of depressive symptoms, results show that low mood and
372 low energy are consistently among the three most central symptoms; this is similar to the results
373 reported in a systematic review that considered the results of 58 cross-sectional depression
374 networks. Interestingly, anhedonia does not appear as a central node on these profile networks,
375 even though it has a strong positive connection with low mood ($pr = 0.30, 0.27, 0.21$), showing a
376 consistent relationship on the three profiles analyzed. This is also consistent with previous
377 studies that found the connection between low mood and anhedonia was the networks' most
378 frequent and robust edge [40]. This is theoretically interesting, considering that anhedonia has
379 been conceived as a main symptom according to the DSM diagnostic criteria for major
380 depressive disorder.

381
382 In terms of methodology, there are limitations related to the sample sizes for each profile
383 subsample that must be considered when interpreting these results. The estimation method could
384 impact the visualization of the networks for small sample sizes, generating networks that overfit
385 to data and impacting the stability of the centrality indexes [53]. Another limitation of this study
386 is the use of cross-sectional data, which provides only a static vision of the profile symptoms that
387 could change over time. Future studies should consider these limitations and explore the
388 relationship between symptoms over time using longitudinal designs with repeated measures.
389 Also, it could be relevant to understand the possible directional influence between the symptoms
390 considering time-series data. An additional limitation of this study, as well as depression
391 heterogeneity research, that has been shown in previous studies [61], is that different instruments
392 can assess different symptoms of depression. Therefore, this study captures the heterogeneity of
393 the specific screening instrument that was used, and other instruments that capture additional

394 symptoms or that phrase the same symptoms differently may yield different heterogeneity
395 profiles. Consequently, no claims can be made about substantive heterogeneity as it occurs in
396 nature (i.e., carving nature at its joints) but rather as it emerges from the use of the PHQ-9, a
397 widely used screening measure and recommended as a preferred measure for the screening of
398 depression [62,63]. It is important to acknowledge that finding common ground for the screening
399 of depression by utilizing one instrument also may have a negative impact on the efforts to map
400 out heterogeneity; it is easier to aggregate findings from different studies but all researchers are
401 looking through the same lens, that could narrow the comprehension of depression [64,65].

402

403 Even with these limitations, this study is the first to our knowledge that combined the
404 identification of qualitatively distinctive symptom profiles and examined the network structure
405 of such profiles using network analyses. Previous network analysis research has shown the
406 relevance of investigating the interrelations between symptoms of depression [40]. While the
407 present results support this approach, they also expand previous research about network analysis
408 and depression and provide empirical support regarding the relevance of looking at the different
409 profiles and the different relations between symptoms for each one. Also, these results could be
410 relevant considering treatment personalization.

411

412 **Author Statement Contributors**

413 Conceptualization: C.N., A.B; Data curation: C.N., A.B; Formal analysis: C.N. with the support
414 of A.B.; Data visualization: C.N, A.B , Writing—original draft: C.N., A.B., J.D., M.B. ;
415 Writing—review & editing: C.N., A.B., J.D., M.B. All authors gave the final approval.

416

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430

431 **Conflicts of Interest**

432 The authors declare that they have no known competing interests that could have appeared to
433 influence the work reported in this paper.

434

435

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628 **Table 1**
 629 Frequency and composition of symptom profiles sample (n=2,023)
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 631

Symptom Profile	Anhedonia	Low mood	Difficulty with sleep	Energy levels	Appetite	Worthlessness	Ability to concentrate	Psychomotor functioning	Suicidal ideation	%	CF%	N
1										25.2%	25.2%	510
2										10.1%	35.3%	205
3										4.0%	39.3%	81
4										3.9%	43.2%	79
5										3.6%	46.8%	74
6										3.2%	50%	65
7										2.9%	52.9%	60
8										2.7%	55.6%	55

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637 **Table 2**
 638 Descriptive statistics of the 8 most frequent theoretical symptom profiles
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	Profile 1	Profile 2	Profile 3	Profile 4	Profile 5	Profile 6	Profile 7	Profile 8
	(N = 510)	(N =205)	(N =81)	(N =79)	(N =74)	(N= 65)	(N=60)	(N=55)
Demographics								
Mean Age (<i>SD</i>)	49.87 (17.40)	45.68 (17.41)	44.47 (15.65)	46.35 (17.37)	50.36 (17.49)	51.23 (16.18)	48.23 (18.02)	44.60 (15.24)
Female (%)	75%	74%	81%	76%	72%	77%	72%	76%
Mean PHQ-9 (<i>SD</i>)	18.60 (4.85)	14.88 (4.09)	13.74 (3.00)	15.18 (3.96)	13.22 (2.58)	15.37 (3.88)	13.17 (2.64)	13.67 (3.12)
Moderate depression rating	23%	57%	69%	51%	70%	42%	67%	71%
Moderately severe depression rating	34%	26%	25%	32%	27%	40%	33%	20%
Severe depression rating	42%	18%	6%	18%	3%	18%	0%	9%
Diagnostic	40%	40%	27%	43%	30%	43%	35%	24%
Treatment	37%	32%	31%	44%	34%	42%	22%	27%

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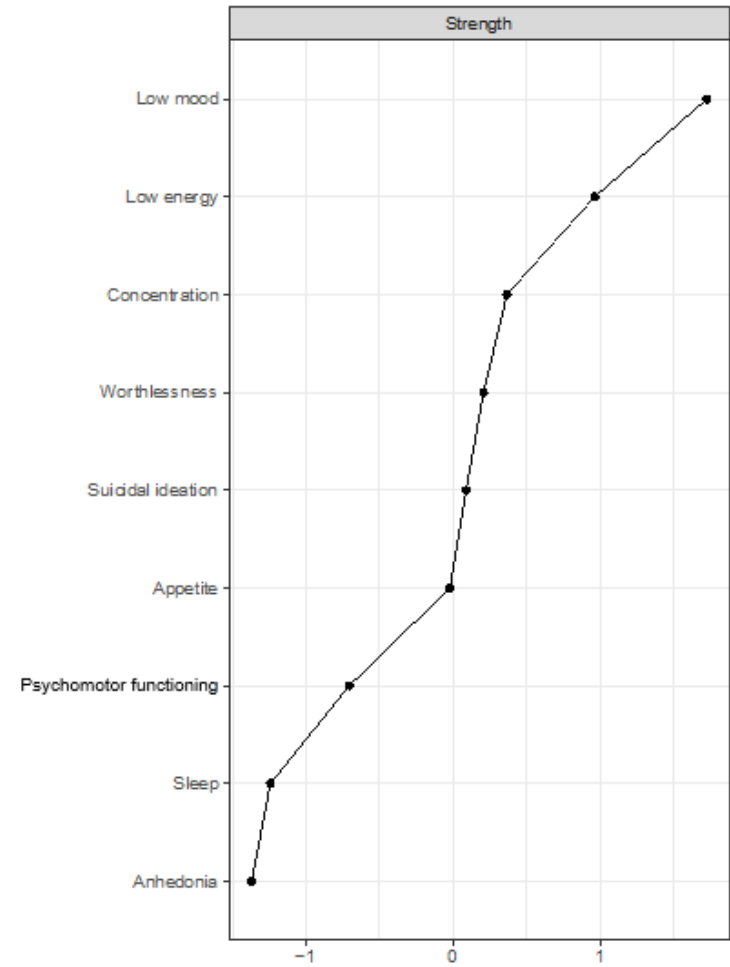
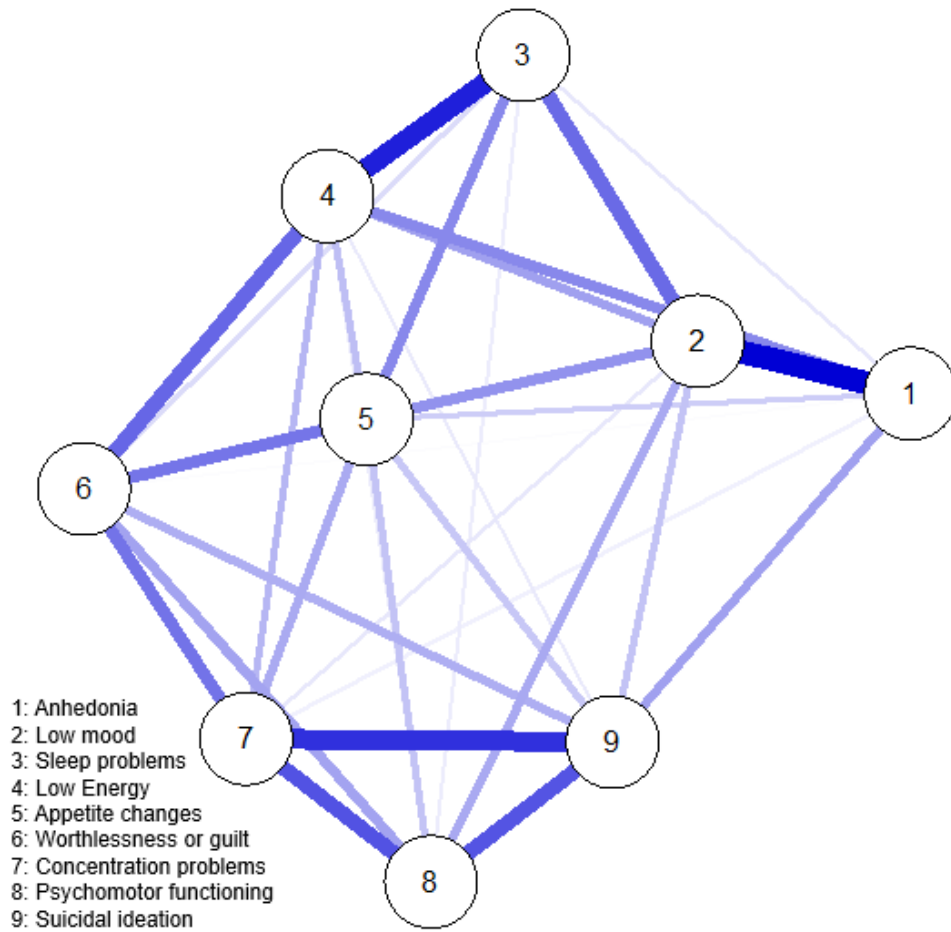
642 Note: 'Diagnostic' is used to identify participants who self-reported having received a diagnosis of depression. On the other hand,

643 'Treatment' is assigned to participants currently undergoing depression treatment.

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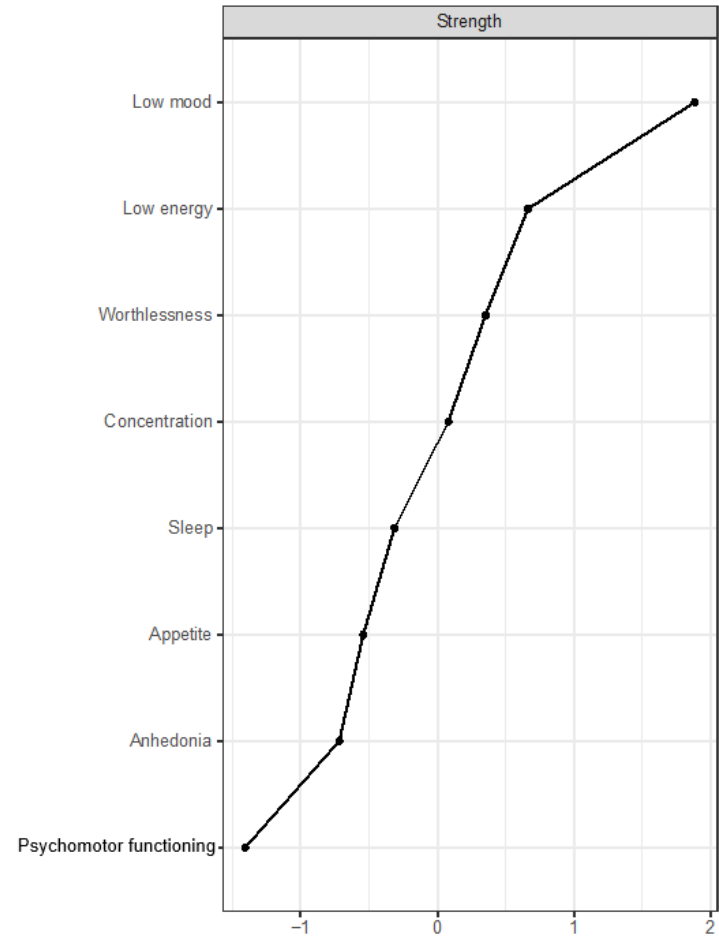
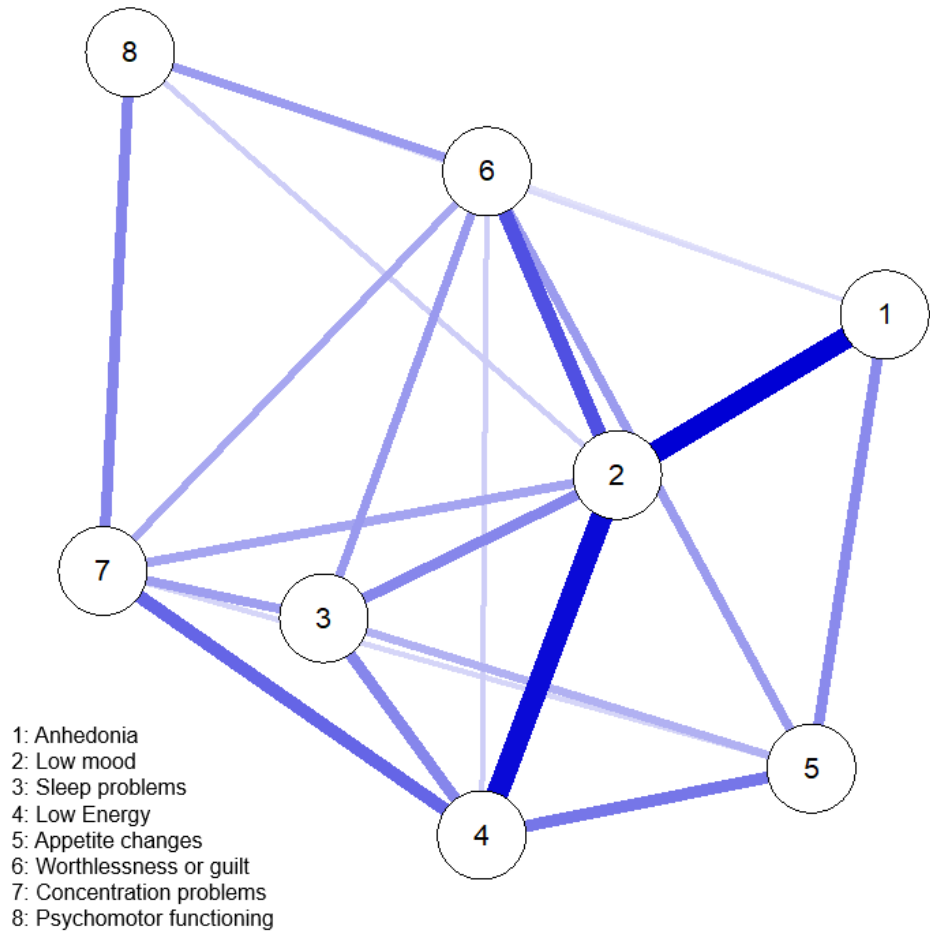
646 Figure 1



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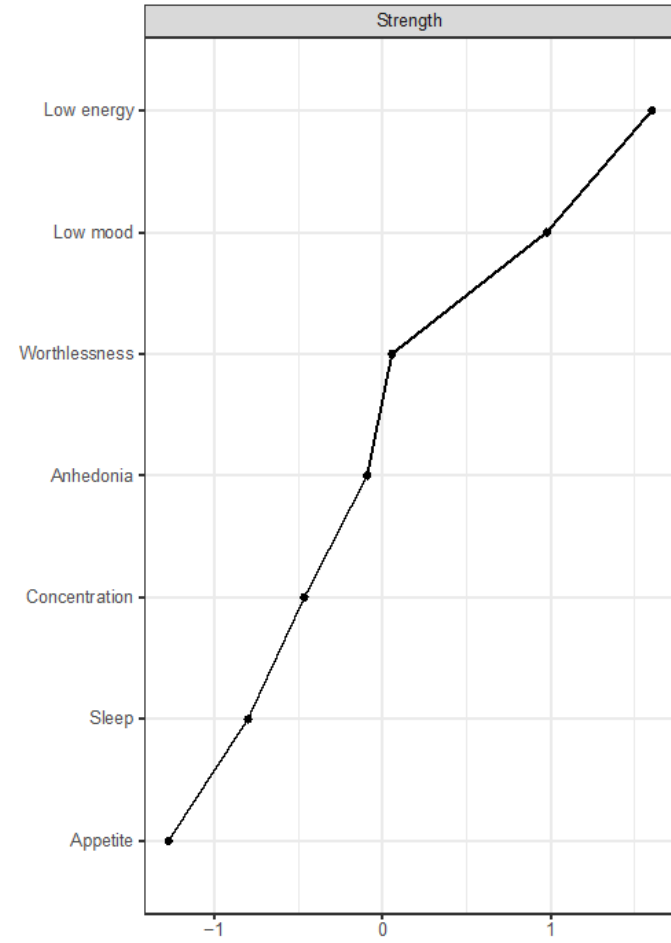
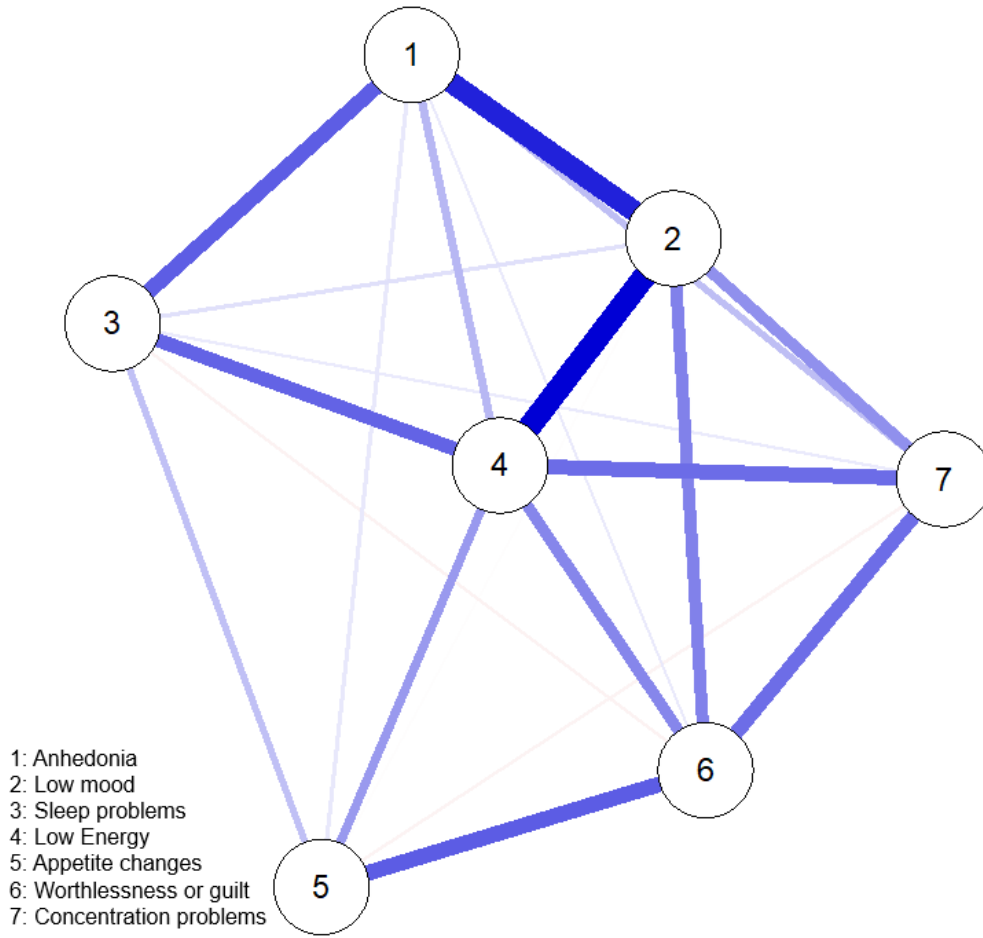
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649 Figure 2



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652 Figure 3



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655 Figure 4



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