

A Taxometric Analysis of Type-D Personality

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Objective: To test the dimensionality of Type-D personality, using taxometric procedures, to assess if Type-D personality is taxonic or dimensional. Type-D personality is treated as a categorical variable and caseness has been shown to be a risk factor for poor prognosis in coronary heart disease. However, at present, there is no direct evidence to support the assumption that Type D is categorical and able to differentiate true cases from noncases. **Methods:** In total, 1012 healthy young adults from across the United Kingdom and Ireland completed the DS14, the standard index of Type D, and scores were submitted to two taxometric procedures MAMBAC and MAXCOV. **Results:** Graphical representations (comparing actual with simulated data) and fit indices indicated that Type D is more accurately represented as a dimensional rather than categorical construct. **Conclusion:** Type D is better represented as a dimensional construct. Implications for theory development and clinical practice with respect to Type D are examined as well as the wider use of taxometrics within psychosomatic medicine (e.g., to investigate if there are medically unexplained syndrome taxons, such as a Gulf War Syndrome taxon). **Key words:** Type D, taxometric analysis, taxon, dimensional, functional syndromes.

NA = negative affectivity; SI = social inhibition; MAMBAC = mean above minus below a cut; MAXCOV = maximum covariance; MAXEIG = maximum Eigenvalue; L-MODE = L-mode factor analysis; CCFI = curve comparison fit index.

INTRODUCTION

The Type D or “distressed” personality is defined as the “. . . tendency to experience negative emotions and to inhibit self-expression” (1 p. 970). Type D plays an important role in contemporary research on the relationship between personality characteristics and cardiovascular disease (1–5). Coronary heart disease patients classified as Type-D cases have a significantly poorer prognosis (1,4), leading Type D to be defined in the literature as a categorical risk factor in coronary heart disease. Whether or not initially intended, this implies a qualitative difference between Type-D cases and noncases. The researchers who originally developed the Type-D construct, Denollet and colleagues, have, in one paper, directly referred to it as a taxonomy (6), but in other papers they have referred to it as a “tendency.” This notion of “tendency” implies dimensionality and others have described Type D as consisting of “. . . negative affectivity (NA) and social inhibition (SI) personality dimensions” (7 p. 235). Whether Type D is dimensional (i.e., distributed as a continuous variable, with individuals varying quantitatively from each other) or taxonic (i.e., individuals are differentiated into nonarbitrary groups or categories) has important clinical and theoretical implications (8). It has been argued that, to explain dimensionality, the existence of multiple, additive causal fac-

tors that sum to produce quantitative variation on the trait is the most plausible model (8). This implies that clinicians and researchers should utilize the full range of scores rather than the use of arbitrary cutoffs (9) and the use of the full range of scores in research (10).¹ By contrast, for a categorical approach there is a need to explain the discontinuity between people. This suggests a greater number of possibilities including a single causal factor (e.g., genetic or threshold models) leading to a dichotomous outcome, or more complex interacting systems, such as environmental influences leading to developmental bifurcation (8). Although exact mechanisms may not be clear at present, the underlying principle is that the existence of dimensional and taxonic models requires very different theoretical accounts. Therefore, it is essential to test directly if Type D is taxonic or dimensional and this is the focus of this paper.

Evidence for the Dimensionality or Taxonicity of Type D

Recently, psychometric work, using item response theory, was interpreted as support for a categorical interpretation of Type D, using a cutoff score of 10 on each of the two subscales (NA and SI) of the DS14 (11). This interpretation should be reconsidered. First, it has been argued that, mathematically, item response theory methods produce unidimensional factors and so lack the specificity necessary to determine nonarbitrary cutoffs (12,13). Second, this study only considered a single cutoff of 10 on each subscale. There is no way to know if this is the optimal cutoff for achieving validity, or if a more reliable function may have been observed at other cutoffs. More importantly, it should be noted that the original decision to define these cutoffs for Type D was based on the combination of cluster analysis and the use of median splits rather than on theoretical considerations (4,14). Cluster analysis is limited when it comes to identifying taxons, as a) it always produces subgroups, and yet b) there is no way to establish the appropriate number of subgroups (15). Using median splits to identify Type-D cases amounts essentially to using arbitrary cutoffs, which others have argued against as a basis for suggesting taxonicity (9).

¹Although this should be generally true, a single cause is always a possibility.

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Received for publication December 23, 2008; revision received July 29, 2009.

DOI: 10.1097/PSY.0b013e3181bd888b

The other type of evidence that could be offered to suggest that Type D is taxonic is based on the following argument: Using these cutoffs, Type D demonstrates good prognostic outcomes. There is a large body of evidence showing that Type-D cases differ from noncases in terms of mortality (4), morbidity (1), biological markers (16–18), physiological stress responses (19), and psychosocial factors (5,20). Although this evidence seems compelling, it does not demonstrate that Type D is taxonic, just that there are between-group differences based on median splits. It is possible to show that median splits for known dimensional traits (e.g., neuroticism, health anxiety) exhibit differences on numerous outcomes (memory effects, reaction times) (8,10,21). This does not mean that these traits are taxonic. The above evidence that Type D is categorical is, therefore, weak.

Furthermore, even if there is a Type-D taxon, but the current cutoff is incorrectly positioned and the Type-D non-case category includes cases, then key psychobiological differences may be obscured, at-risk cases may be missed, and prognostic power may be reduced. Conversely, if the Type-D case category includes noncases, biological differences may be obscured and prognostic power would be reduced, although Type-D cases would not be missed.

The idea that Type D is a dimensional construct comes from the finding that a number of individual differences theoretically and empirically related to the subcomponents of Type D—worry, depression, anxiety (14,22,23)—are dimensional (8). This implies that the latent structure of the subcomponents of Type D, NA and SI, should be dimensional. Thus, although indirect, this evidence is based on taxometric evidence and it is reasonable to hypothesize that Type D may be dimensional.

The Present Study

At present, there is no direct evidence concerning whether or not Type D is dimensional or taxonic. However, even in the absence of direct evidence that Type D is taxonic, the “established cutoffs” are still used to split samples into cases and noncases. Given the very different implications for theory, research, and clinical practice afforded by dimensional and taxonic conceptualizations, a direct test of the dimensionality of Type D is urgently required (8). This paper tests the dimensionality of Type-D, using taxometric procedures (13,24–26), in an unselected, healthy, young adult sample of university students. We deliberately avoided recruiting a sample containing subpopulations with particular disorders (in this case, cardiovascular disease) alongside groups of healthy persons, as such a combination of heterogeneous groups could lead to the erroneous identification of a taxon (i.e., a pseudotaxon) (8,15,26–28). In the case of Type D, the evidence shows that the prevalence of Type D is significantly higher in clinical groups with cardiovascular disease compared with healthy controls (14). This increased prevalence of Type-D cases in clinical samples may lead to the identification of a pseudotaxon if combined with a healthy sample where the prevalence rate is significantly lower. Furthermore, a sample of healthy participants is likely to cover the full range of

scores on Type D and be normally distributed, limiting skew. Positive skew is known to lead to the identification of pseudotaxa with low base rates, whereas negative skew can misidentify a high base-rate taxon (29). Furthermore, the initial analysis of the DS14 (14) by Denollet confirmed that the same factor structure was present in both clinical cardiovascular and nonclinical samples. As such, the use of a non-clinical sample in this study is appropriate and consistent with other psychometric analyses (14).

METHODS

Sample and Sampling

In total, 1012 healthy young adults took part in this study ($n = 787$ females, 225 males; mean age = 20.5 years, standard deviation (SD) = 4.84; age range = 17–61 years). These subjects were recruited via convenience sampling from eight universities across England, Scotland, Northern Ireland and Republic of Ireland. The samples from England, Scotland, Northern Ireland and Republic of Ireland did not differ significantly in terms of age ($F(5,1006) = 0.740, p = .59$), or gender ($\chi^2(6, n = 1012) = 6.59, p = .29$). The study was conducted over 12 months across 2006 to 2007. The study was approved by the relevant local ethics committees.

Measures

Type D Personality

Type D was assessed, using the DS14 (14). It is a 14-item measure answered on a 5-point Likert type scale ranging from 0 (false) to 4 (true), consisting of two subscales assessing the NA and SI components of Type D. Participants who score highly on both NA and SI, using a cutoff point of ≥ 10 on both scales, are classified as having a Type-D personality (14). Both subscales were internally consistent in the current study ($\alpha = 0.85$ and 0.82 for NA and SI, respectively).

Taxometric Analyses

The analysis used taxometric procedures developed by Meehl and colleagues (13,24,25,27) and implemented following recommendations made by Ruscio and colleagues (8,26,28–32). These procedures run over three main steps in terms of a) identifying valid construct indicators; b) applying the appropriate taxometric techniques; and c) interpretation (30–32).

Identifying Valid Indicators

Indicators are retained if they showed good indicator validity, distinguishing Type-D cases (termed taxons) from noncases (termed complements). Meehl (27) has suggested that valid indicators should have a mean separation expressed in standard units in terms of a Cohen's d of 1.25. Second, of the valid indicators, those with high item-total correlations are retained as they represent the most valid indicators of the construct (26). Finally, evidence for nuisance covariance in the retained indicators is explored. For nuisance covariance to be tolerable, this should be ≤ 0.30 (27).

Taxometric Techniques

The basic principle of Meehl's taxometric technique is termed “coherent cut kinetics” where indicator variables are split into input and output variables. For successive divisions along the input variable, either mean differences either side of a cut (mean above minus below a cut: MAMBAC), or covariances (maximum covariance: MAXCOV) or Eigenvalues (maximum Eigenvalues: MAXEIG) within a cut (or subsample) are computed for the remaining indicators (called output variables) (25,30–32). Based on the general covariance mixture theorem (25,30), the mean difference, covariances or Eigenvalues for valid indicators will be at a maximum when the sample contains equal proportions of both taxons and complements and at a minimum when either complement or taxons are present alone. With divisions represented on the x-axis and mean differences, covariance or Eigenvalues on the y-axis, characteristic curves represent either a taxonic or dimensional solution. If the latent

TABLE 1. Indicators Statistics

Items	Domain	Item-Total	Cohen's <i>d</i>	Skew
1. I make contact easily when I meet people	SI	0.49	1.05	0.55
2. I often make a fuss about unimportant things	NA	0.33	0.63	-0.19
3. I often talk to strangers	SI	0.20	0.56	-0.08
4. I often feel unhappy	NA	0.60	1.05	0.29
5. I am often irritated	NA	0.51	0.93	0.02
6. I often feel inhibited in social interactions	SI	0.63	1.37	0.13
7. I take a gloomy view of things	NA	0.64	1.21	0.75
8. I find it hard to start a conversation	SI	0.57	1.25	0.47
9. I am often in a bad mood	NA	0.60	1.06	0.68
10. I am a closed kind of person	SI	0.52	1.26	0.38
11. I would rather keep other people at a distance	SI	0.50	1.15	0.72
12. I often find myself worrying about something	NA	0.45	0.85	-0.39
13. I am often down in the dumps	NA	0.68	1.23	0.62
14. When socializing, I don't find the right things to talk about	SI	0.56	1.22	0.64
Two subcomponents				
Negative affect ($\alpha = 0.85$)	NA		1.48	0.24
Social inhibition ($\alpha = 0.82$)	SI	0.41 ^a	1.91	0.31
Three indicators				
Negative affect ($\alpha = 0.85$)	NA	0.41	1.48	0.24
Reticence ($\alpha = 0.78$)	SI	0.61	1.85	0.30
Social discomfort ($\alpha = 0.70$)	SI	0.43	1.25	0.24

^a Interscale correlation.

SI = social inhibition; NA = negative affectivity.

structure is taxonic (the sample is a mixture of taxons and complements), the curve will be characteristically peaked for MAMBAC (identifying the division that maximally separates complement from taxon) or either peaked or cusped for MAXCOV/MAXEIG.² If the structure is dimensional (i.e., there is not a specific taxon), then at different cuts/subpopulations the values should remain virtually unchanged and the graph is either flat or concave.

Which taxometric technique to use depends, in part, on the number of indicators (30). MAMBAC requires a minimum of two indicators, one is designated the input and one is designated the output. The mean difference on the output variable is calculated above and below successive cuts on the input variable. MAXCOV is used with at least three indicators. One indicator acts as an input variable and is used to divide the sample into a succession of overlapping subsamples or windows. The covariance between the other output indicators is calculated at successive windows. All possible combinations of input-output triplets are used. MAXEIG is used when there are >3 indicators, this is an extension of MAXCOV, where the 1st Eigenvalue is calculated across successive windows on the remaining output variables. An alternative approach to "coherent cut kinetic" is to base taxometric analyses on factor analytic procedures (25). For example, L-Mode factor analysis requires multiple indicators and uses factor analytic procedures to calculate scores on the 1st principal component and the distribution of these is plotted. A unimodal distribution indicates a dimensional solution and a bimodal distribution indicates a taxonic solution.

Interpretation

Interpretation of taxometric analysis involves inspecting the characteristic shape of the curves. As visual inspection can be open to judgment errors and skewed indicators can lead to the erroneous identification of taxa (29–32), Ruscio and colleagues have developed a set of interpretative techniques based on comparisons to simulated taxonic and dimensional curves and the development of the curve comparison fit index (CCFI) (30–32). This approach is based on generating a series of simulated taxonic and dimensional curves (based on ten replications) derived from the actual data characteristics (30).

²A MAXCOV/MAXEIG curve is cusped if there are too few number of either taxon or complement subgroups either side of the most heterogeneous subsample (8,31).

The actual data are plotted relative to the simulated curves plotted ± 1 SD. This allows a contextualized interpretation. To further aid interpretation the CCFI, a relative fit index, is used (8,30–32). The CCFI varies between 0 and 1 and is symmetric around 0.5. Values of >0.5 indicate a taxonic solution and values of <0.5 indicate a dimensional solution. Extensive Monte-Carlo studies support the precision of these methods relative to all other procedures (30,32). Finally, the results need to show consistency. This means using a number of taxometric procedures to show consistency across methods and showing the same results at the same time successively increasing the number of cuts/windows (30,32).

RESULTS

Prevalence of Type D

Using the recommended cutoff point of ≥ 10 on both NA and SI to define Type D (14), 39.6% of females and 34.7% of males were categorized as Type D. Type-D classification did not vary as a function of either sex ($\chi^2(1) = 1.83, p = .41$) or sample region ($\chi^2(6) = 16.32, p = .33$). As Type D is hypothesized to be an independent risk factor for cardiovascular disease, this prevalence rate seems high and highlights how the simple taxonic model can be problematic as a prognostic risk factor in a healthy population.

Selection of Indicators

The indicator validities (expressed in standard units as Cohen's *d*), item-total correlations, and skew for each indicator are presented in Table 1.

Initially, these statistics were examined for the 14 items of the DS14. All indicators showed a degree of skew that was generally within the ranges reported for other taxometric studies (29). However, only three items, all SI items, had indicator validities ≥ 1.25 . Thus, at the item level, there was not sufficient taxon-complement separation to conduct taxometric analyses.

TABLE 2. Rotated Factor Matrix for the Social Inhibition Items

Item	F1	F2	Construct
10. I am a closed kind of person	0.79	0.15	Reticence
11. I would rather keep other people at a distance	0.79	0.00	Reticence
14. When socializing, I don't find the right things to talk about	0.69	0.27	Social discomfort
6. I often feel inhibited in social interactions	0.68	0.31	Social discomfort
3. I often talk to strangers (<i>r</i>)	-0.02	0.87	Social poise
1. I make contact easily when I meet people (<i>r</i>)	0.44	0.68	Social poise
8. I find it hard to start a conversation	0.56	0.57	Social discomfort
α	0.78	0.70	

F1 = reticence; F2 = social discomfort.

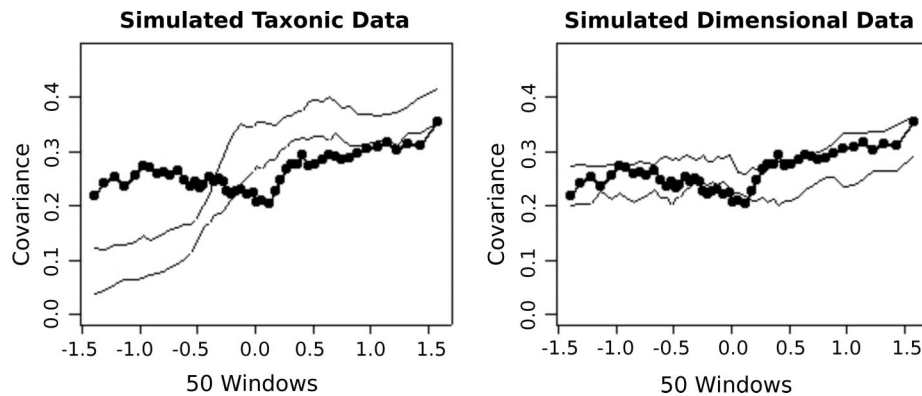


Figure 1. Maximum covariance (MAXCOV) curves based on 50 windows. Smooth lines mark the simulated curves +1 and -1 SD from the mean simulated data. Dark lines represent the actual data. SD = standard deviation.

Examining the two main subcomponents of Type D (i.e., NA and SI) indicated that both showed good indicator validity, had acceptable skew, and were correlated with each other ($0.42, p < .001$). There is some evidence that NA and SI may themselves be multidimensional factors (11). Principal components analysis was applied separately to the NA and SI items. The results showed that the NA items loaded on a single factor (Eigenvalue = 3.819, % variance = 54.5%). The SI items loaded onto two separate factors (Eigenvalues = 3.466 and 1.004; % variance = 63.8%). The varimax rotated solution for the SI items are presented in Table 2. The first factor represents being closed and keeping others at a distance and we refer to this primarily as a “reticence factor.” The second factor represents finding social contact inhibiting and uncomfortable and we refer to this as a “social discomfort factor.” These three factors (NA, reticence, and social discomfort) all showed good indicator validity, item-total correlations, and acceptable skew. Based on these analyses, MAMBAC was applied to NA and SI and MAXCOV to the NA, reticence, and social discomfort. With a maximum of three indicators, there are not a sufficient number to justify MAXEIG or L-Mode factor analysis. MAMBAC analyses were performed with 50, 150, and 200 cuts and MAXCOV with 50, 150 and 200 windows with 90% overlap.

Within the putative taxon and complement (using the recommended cutoff point of ≥ 10 on both NA and SI), the nuisance covariance for NA and SI for the taxon was 0.30, and for the complement, it was -0.09 . For the three indicators, the mean nuisance covariance for the taxon was 0.22 and 0.09 for

the complement. Furthermore, nuisance covariance for the two indicators in the MAMBAC analysis for the taxon and complement at 50 cuts were -0.14 and -0.06 , respectively; at 150 cuts, they were -0.15 and -0.06 , respectively, and for 200, they were -0.15 and -0.06 , respectively. Similarly, mean nuisance covariance for the three indicators in the MAXCOV analyses for the taxon and complement at 50 windows were 0.23 and 0.26, respectively; at 150 windows, they were 0.23 and 0.20, and for 200 windows, they were 0.11 and 0.24, respectively. As such, there were no problems with nuisance covariance.

Taxometric Analyses

For the MAMBAC analyses, the CCFIs were 0.32, 0.32, and 0.35 for 50, 150, and 200 cuts, respectively; for the MAXCOV, the CCFIs were 0.21, 0.35, and 0.36 for 50, 150, and 200 windows, respectively. All the CCFI values are < 0.50 , suggesting a dimensional interpretation of the latent dimensional structure of Type D is a more accurate representation than a taxonic one. Confirming this, the taxometric curves for the MAMBAC and MAXCOV analyses all indicated that the actual data were more similar to the simulated dimensional rather than the simulated taxonic data curves. Figure 1 provides the MAXCOV taxometric curves at 50 windows as an example.³

³The remaining MAMBAC and MAXCOV curves are available from the first author on request.

DISCUSSION

The pattern of results reported here indicates clearly that Type D is better represented as a dimensional rather than a categorical construct. This finding has important theoretical and clinical implications for Type D. Theoretical models should focus on additive multicausal agents or risk factors (8), as research into Type D indicates that there are a wide variety of mechanisms associated with Type D (4,5,18,20). However, more research is now needed to explore the additive nature of these different mechanisms. Thus, although it is implicitly used as a taxonic construct, the research agenda supporting Type D conforms more to one defined for a dimensional construct.

Type D as a Dimensional Construct in Research and Practice

Based on the above, theoretical models and clinical interventions should be examined using regression approaches with large unselected samples (8), which also have the advantage of increasing statistical power (33). One of the main assumptions of Type D is that it is defined synergistically with ill effects only seen for high NA in combination with high SI; when either trait is low, there are no effects on health. Given the dimensional nature of Type D, these potential synergistic effects of Type D should be tested, using both additive and multiplicative regression terms (18) (Williams L, O'Connor RC, Grubb, N, O'Carroll RE. Type D personality predicts medication non-adherence in myocardial infarction patients. Submitted for publication, 2009). If the same prognostic power for Type D is not seen when treated as the interaction between these continuous traits, then its construct validity could be questioned.

How should Type D be used in clinical practice as a dimensional rather than a taxonic construct? It has been argued that it is appropriate to draw distinctions within a latent dimensional construct as long as these are systematic and empirically justifiable (8). This approach is similar to the use of continuous diagnostic signs in physical medicine (e.g., blood pressure, heart rate, temperature) that are integrated to reach a final differential diagnosis. One way to achieve this with a dimensional construct is to identify "inflection points" (34). An inflection point expresses the association between the continuous dimensional construct and other relevant clinical data (e.g., biomarkers, prognostic clinical outcomes, or treatment responses). The point of inflection marks the position on the continuum where these show dramatic accelerated changes (34). For example, with Type D, this may indicate the point where there is an increase in cardiovascular mortality. However, the issue for Type D concerns its definition along two constructs. It may be, therefore, that inflection points need to be either a) identified for both separately and then combined or b) combine the two scales and identify a single inflection point or c) use interaction terms specified with different sliding cuts to identify the point at which the interaction best predicts future prognostic outcomes. With respect to diagnosis, it should also be noted that Type D is a risk factor and not

a clinical diagnosis in itself and should be used in conjunction with other information. Therefore, the issue of cutoffs is perhaps less crucial rather than defining people with respect to the "normal" range of scores.

Taxometrics and Psychosomatic Medicine: Functional Syndromes and Biomarkers

It has been strongly argued that taxometric approaches have implications for many fields beyond their current application mainly to investigating psychopathology (e.g., depression, personality disorder) (15). Within the field of psychosomatic medicine, these may be applied initially to the dimensionality of traits directly relevant to psychosomatic medicine. For example, health anxiety, alexithymia, anxiety, and posttraumatic stress disorder have all already been shown to be dimensional (8,21,35), whereas Type A and self-monitoring have been identified as taxonic (8). Future work could examine the dimensionality of functional syndromes. This would allow key questions to be answered, such as whether or not medically unexplained/functional syndromes, such as irritable bowel syndrome, chronic fatigue syndrome, and Gulf War syndrome are taxonic representing unique syndromes (36). Taxometric analyses could, and should, be applied to existing symptom databases to address these fundamental and important questions. Taxometric procedures could also prove useful with other data types relevant to psychosomatic medicine, such as biological markers (37) and social interactions (38). For example, is there a particular taxon for cortisol under- or overreactors? Are there people who are able to cope with social stress in a qualitatively different way? Once evidence on the dimensionality (boundaries within a disorder: taxon versus complement) has been identified, issues pertaining to the boundaries between syndromes (if chronic fatigue syndrome and irritable bowel syndrome are identified as taxonic, are they distinct or do they overlap?) can be examined (28).

Caveats

The use of a convenience sample of healthy young adults may lead to questions about the generalizability of the findings. However, the question addressed herein is about the latent-dimensional structure of Type D: If the construct is a general personality trait, then its latent structure should be stable across populations. Denollett (14) used a healthy non-clinical sample as part of the development of the DS14. Furthermore, using a sample that is selected to combine clinical cases (with cardiovascular disease) with healthy nonclinical cases may lead to the identification of a pseudotaxon (8). That is, evidence shows that the prevalence of Type-D cases is significantly higher in cardiovascular clinical groups compared with healthy controls (14). This increased prevalence of Type-D cases in clinical samples may lead to the identification of a pseudotaxon if combined with a healthy sample, where the prevalence rate is significantly lower. However, given that the majority of research in the area of Type D has been conducted on clinical cardiac samples, it is important that the

results of these taxometric analyses are replicated in a) an appropriately sampled clinical sample of cardiovascular disease respondents, and b) a larger healthy community sample without cardiovascular disease, but not in samples that combine clinical and nonclinical cardiovascular samples.

CONCLUSIONS

The present taxometric analysis indicates that Type D is better represented as a dimensional construct. Future theorizing and research examining the links between Type D and cardiovascular disease should consider dimensional approaches to move this area of inquiry forward.

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