

Affect Variability and Predictability: Using Recurrence Quantification Analysis to Better Understand How the Dynamics of Affect Relate to Health

Brooke N. Jenkins
Chapman University

John F. Hunter
University of California, Irvine

Michael J. Richardson
Macquarie University

Tamlin S. Conner
University of Otago

Sarah D. Pressman
University of California, Irvine

Changes in affect over time have been associated with health outcomes. However, previously utilized measurement methods focus on *variability* of affect (e.g., standard deviation, root mean squared successive difference) and ignore the more complex temporal patterns of affect over time. These patterns may be an important feature in understanding how the dynamics of affect relate to health. Recurrence quantification analysis (RQA) may help alleviate this problem by assessing temporal characteristics unassessed by past methods. RQA metrics, such as determinism and recurrence, can provide a measure of the *predictability* of affect over time, indexing how often patterns within affective experiences repeat. In Study 1, we first contrasted RQA metrics with commonly used measures of variability to demonstrate that RQA can further differentiate among patterns of affect. In Study 2, we analyzed the associations between these new metrics and health, namely, depressive and somatic symptoms. We found that RQA metrics predicted health above and beyond mean levels and variability of affect over time. The most desirable health outcomes were observed in people who had high mean positive affect, low mean negative affect, low affect variability, and high affect predictability. These studies are the first to demonstrate the utility of RQA for determining how temporal patterns in affective experiences are important for health outcomes.

Keywords: recurrence quantification analysis, affect variability, affect predictability, depressive symptoms, somatic symptoms

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Affective experiences are dynamic in nature. Feelings fluctuate from moment to moment and are intricately interconnected in a complex temporal system (Ebner-Priemer, Eid, Kleindienst, Stabnow, & Trull, 2009; Pressman, Jenkins, Kraft-Feil, Rasmussen, &

Scheier, 2017). Many studies have demonstrated that average affect is linked to a variety of psychological and physical health and health-relevant outcomes (e.g., Chida & Steptoe, 2008; Jenkins, Granger, et al., 2018; Pressman & Cohen, 2005; Pressman, Jenkins, & Moskowitz, 2018), but, increasingly, studies have revealed that the variation in affect over time may further predict health and health-relevant factors. For example, greater levels of affect variability, assessed with standard deviation (*SD*), have been associated with less favorable daily cortisol trajectories (Human et al., 2015) and worse immunocompetence (Jenkins, Hunter, Cross, Acevedo, & Pressman, 2018). These findings mirror patterns in mental health outcomes, which typically find greater affect variability in people with higher levels of depressive symptoms, greater anxiety, and/or poorer psychological well-being (Gruber, Kogan, Quidbach, & Mauss, 2013; Peeters, Berkhof, Delespaul, Rottenberg, & Nicolson, 2006). Therefore, it is critical that researchers be equipped with the proper tools to accurately assess the dynamics of affect. However, current measurements of affect

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Brooke N. Jenkins, Department of Psychology, Chapman University; John F. Hunter, Department of Psychological Science, University of California, Irvine; Michael J. Richardson, Department of Psychology, Macquarie University; Tamlin S. Conner, Department of Psychology, University of Otago; Sarah D. Pressman, Department of Psychological Science, University of California, Irvine.

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Correspondence concerning this article should be addressed to Brooke N. Jenkins, Department of Psychology, Chapman University, 110 Crean Hall, Orange, CA 92668. E-mail: bjenkins@chapman.edu

dynamics fail to capture important information about changes over time because of their limited ability to assess the temporal structure within complex patterns of affective change.

The most common metric used to assess affect variability is *SD* (Röcke, Li, & Smith, 2009). This approach captures affect variability as a single value by calculating the *SD* of an individual's scores on an affect measure over multiple time points (Eid & Diener, 1999; Gruber et al., 2013; Hardy & Segerstrom, 2017; Ram & Gerstorf, 2009; Röcke & Brose, 2013). An advantage of this approach is that *SD* is easily understood and the value can be used as a predictor or outcome variable in subsequent models (Röcke et al., 2009). Nevertheless, this single value assumes an independence of assessment such that an assessment at time, t , is not necessarily related to time, $t + 1$ (Ram & Gerstorf, 2009). Assessing only *SD* offers information about the magnitude of the change in affect but provides no information about the sequential dependence or deterministic *predictability* of affective experiences.

More recently, researchers have begun using root mean squared successive difference (RMSSD; Ebner-Priemer et al., 2009). This metric captures the degree of change in affect from one time point to the next by squaring all successive differences, averaging these squared differences, and then taking the square root of that average. RMSSD takes into account temporal ordering by reflecting the relative instability of affective change from one time point to the next. Nevertheless, RMSSD does not capture patterning of change over time beyond average change among two time points. A similar limitation occurs when measuring emotional inertia, which is often operationalized as the degree of autocorrelation between consecutive time points (Kuppens, Allen, & Sheeber, 2010).

Information about the dynamics of affect over large time spans can provide more detailed insights over and above current affect variability measures. For example, an individual who has more structured or “predictable” negative affect (NA) values that denote a more recurrent pattern of affective experiences over time (i.e., exhibits a less stochastic and more stationary or periodic structure of affective change over time) may have much different outcomes compared with an individual who has “less predictable” values of NA that create a more inconsistent (i.e., stochastic) or less recurrent pattern of affect (see Figure 1). The predictability of fluctuations in affective experiences over time (i.e., regularity and consistency) may matter more than (or in addition to) simply assessing the magnitude of those fluctuations. Broadening the quantification

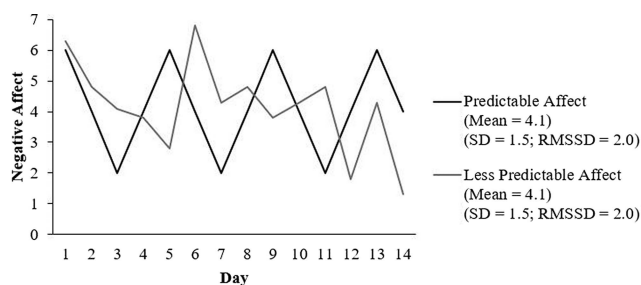


Figure 1. Two individuals with different predictability of affect but with identical means and variability. *SD* = standard deviation; RMSSD = root mean squared successive difference.

of change to include metrics of predictability in addition to variability may allow researchers to better understand the full dynamics of affective experience over time.

RQA may be one such method for assessing patterns of affect predictability. RQA is a nonlinear event- or time-series analysis method that assesses the dynamics of temporal sequences of change over time without researchers having to make any a priori assumptions about the nature of the dynamics that define a given behavioral event- or time-series recording. Although RQA has not been previously used to examine intraindividual changes in affect over time, it has been successfully used to investigate a wide range of other dynamic human behaviors (e.g., eye gaze, Anderson, Bischof, Laidlaw, Risko, & Kingstone [2013]; posture changes, Riley & Clark [2003]; and affect synchrony between mother–child dyads Main, Paxton, & Dale [2016]). Because of the temporal nature of affect, RQA lends itself nicely to studying dynamics of affect and, in particular, the degree of affect predictability (Richardson, Dale, & Marsh, 2014). RQA provides multiple metrics of predictability, two of the most common being the percentage of deterministic structure (%DET) within a measured series and recurrence rate (i.e., percent recurrence, %REC). %DET measures the degree to which recurrent states exhibit regular or structured patterns of change over time (i.e., repeated sequences of recurrent states). With regard to how affect changes over time, %DET captures the degree to which the same (or similar) sequences of affective change occur over time, such that more structured or predictable patterns of affective change over time should result in high levels of %DET. %REC is a measure of state regularity that, in terms of affect, reflects how often a person experiences the same (or similar) level of affect over time (i.e., the degree to which the same state of affect reoccurs over time). Although %DET and %REC may provide different information for longer time-series data (e.g., over 50 time points), they are often correlated for shorter time-series (e.g., 15 time points).

Of particular relevance to the current study is the fact that the RQA metrics of %DET and %REC should provide more information about the structural dynamics of affective change than standard (linear) variability statistics. In other words, the benefit of these RQA metrics with regard to understanding the complex dynamics of affect is that they can quantify the *predictability* of affect over time. Although previous research has demonstrated that greater affect variability (e.g., higher *SD*) has negative implications for health (e.g., Gruber et al., 2013), no previous research has examined whether higher affect predictability may be beneficial for health. Specifically, the regularity or predictability of how affect changes over time may allow individuals to better prepare and then cope with affective experiences. For example, an individual who knows that NA is always high Monday mornings may be better prepared to cope with such NA. Thus, it is important to test how mean levels of affect, affect variability, and affect predictability interact, as these factors may operate together with health implications.

The purpose of the current research was to demonstrate the validity of the RQA method for quantifying the structure or predictability of affective time-series using simulated data (Study 1) and then apply RQA to a large, real data set (Study 2). In Study 1, the RQA metrics of %DET and %REC were compared with common variability metrics (*SD*, RMSSD) using simulated data. This simulation compared predictability and variability metrics

and determined whether each could add independent information about the patterns of affect across time (i.e., days). Specifically, we hypothesized that the same values of *SD* or *RMSSD* would be associated with, but not differentiate between, patterns that are more or less predictable (i.e., stochastic and/or periodic), but that patterns could be quantifiably differentiated using the RQA metrics of %DET and %REC. In Study 2, the RQA metrics of %DET and %REC were then used, along with mean levels and variability, to predict health outcomes. We hypothesized that, as in previous literature, more variability would be associated with worse psychological and physical health outcomes (i.e., more depressive and somatic symptoms). In contrast, we also hypothesized that predictability would be associated with better psychological and physical health outcomes (i.e., fewer depressive and somatic symptoms).

Study 1

Method

Data simulation. Affect data were simulated by creating 14 instances for 900 cases, thus, emulating typical daily dairy data collected once a day for 2 weeks with 900 participants. Within these 900 cases, nine distinct groups of 100 cases each were created to alter variability and predictability (see Table 1). Data were first simulated using the Stata 15 (StataCorp, 2017) generate function to randomly draw values from normal distributions to alter the variability. Groups 1 through 3 had integer values generated from a normal distribution with a mean of 3.00 and a *SD* of 0.25 (see Table 1 column 3). Groups 4 through 6 had integer values generated from a normal distribution with a mean of 3.00 and a *SD* of 0.50. Groups 7 through 9 had integer values generated from a normal distribution with a mean of 3.00 and a *SD* of 1.00. Generating the integer valued time-series in this way ensured that Groups 1 through 3 would have small *SDs* (i.e., low variability), Groups 4 through 6 would have medium *SDs* (i.e., medium variability), and Groups 7 through 9 would have large *SDs* (i.e., high variability).

In addition to altering the variability of integer values, some cases had value sequences repeated (see Table 1 column 4) to alter predictability of the data, such that the data series contained levels of periodic structure. Repeating the values ensured that there would be greater predictability within these groups of data series, with greater levels of repeated (periodic) structure corresponding to higher predictability. One third of the groups had no values repeated (i.e., low predictability). One third of the groups had

instances 1 through 7 repeated once (i.e., medium predictability). One third of the groups had instances 1 and 2 repeated 7 times (i.e., high predictability). It is important to note that this method of generating the data was not specific to a particular affect measure per se, but was used to simply represent different amounts of variability and predictability that might be associated with changes in affect over time when rated on continuous scales (e.g., changes in positive affect [PA] or NA across a 14 day daily diary period).

Measures.

Mean. Means were calculated within individuals by summing the values for each of the 14 days and then dividing by 14. Each participant's mean, therefore, represents their average score over the entire daily diary period.

Variability. Variability was assessed by *SD* and *RMSSD*. *SDs* were calculated within individuals by summing the squared distances for each day from the overall mean and then averaging those squared distances. *RMSSD* was calculated by squaring all successive differences, averaging them together, and then taking the square root of that average. The following formulas were used:

$$SD = \sqrt{\frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n}}$$

$$RMSSD = \sqrt{\frac{\sum_{i=2}^n (x_i - x_{i-1})^2}{n-1}}$$

Predictability. Predictability was measured using RQA. The RQA measures of percent determinism (%DET) and percent recurrence (%REC) were calculated using the RQA software developed by Richardson, Riley, Shockley, and Dale (2015; see supplemental material A for a detailed description of how to compute %DET and %REC). Given that integer value time-series data were investigated here and in Study 2, a form of RQA known as Categorical-RQA was used. As illustrated in Figure 2, this method of RQA first involves identifying reoccurring (recurrent) values within a discrete time-series by plotting them on a two-dimensional recurrence plot (see Figure 2). Essentially, a data time-series is represented on both the *x* and *y*-axis of a two-dimensional grid, with recurrent points indicating when the same value within the data series reoccurs. For example, a row of simulated data is presented in Figure 2. Recurrent states (i.e., "points") within the recurrence plot correspond to when the same value reoccurs indicating that the same level of affect that was previously experienced has reoccurred. Note that given that each value within the data series is recurrent with itself, the main

Table 1
Data Simulation Parameters and Predicted Results

Group	Group name	Normal distribution (mean, <i>SD</i>)	Number of days repeated	Predictions	
				Variability	Predictability
1	Low variability–Low predictability	(3, .25)	None	Low	Low
2	Low variability–Medium predictability	(3, .25)	7 (first 7 are repeated a second time)	Low	Medium
3	Low variability–High predictability	(3, .25)	12 (first 2 are repeated 7 more times)	Low	High
4	Medium variability–Low predictability	(3, .5)	None	Medium	Low
5	Medium variability–Medium predictability	(3, .5)	7 (first 7 are repeated a second time)	Medium	Medium
6	Medium variability–High predictability	(3, .5)	12 (first 2 are repeated 7 more times)	Medium	High
7	High variability–Low predictability	(3, 1)	None	High	Low
8	High variability–Medium predictability	(3, 1)	7 (first 7 are repeated a second time)	High	Medium
9	High variability–High predictability	(3, 1)	12 (first 2 are repeated 7 more times)	High	High

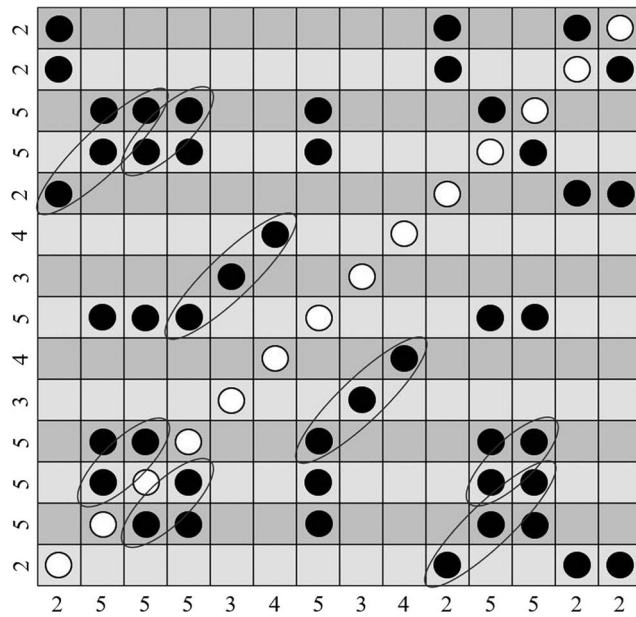


Figure 2. Data analysis for recurrence quantification analysis (RQA). The same data time-series is placed on the x - and y -axis. Recurrent states or “points” are placed where same values intersect. For example, in the lower left corner a “point” is placed because there is a 2 on the x -axis and a 2 on the y -axis. Points that are circled are those that lie on a diagonal line. Only points that form diagonal lines (excluding the line of identity; the white points) that contain two or more recurrent points are used in the determinism calculation.

diagonal of an (auto-) recurrence plot (i.e., single time-series recurrence plot), also known as the line of identity, is ignored. That is, %DET and %REC quantifications exclude recurrent states along the main diagonal.

In short, %DET equals the percentage of recurrent points that form diagonal lines (in this case 20; excluding the line of identity) of the total number of points (in this case 46) within a recurrence plot, where a diagonal line corresponds to two or more consecutive recurrent points. For instance, in **Figure 2**, $\%DET = 20/46 = .43 \rightarrow 43\%$. With regard to an affect data series, %DET represented the percentage of time an individual experiences the same pattern of change in affect over time and, therefore, the degree of predictability or deterministic structure within an affect data series. %REC is calculated by dividing the number recurrent points that do not fall along the main diagonal (in this case 46) by the number of spaces (in this case 182). In **Figure 2**, $\%REC = 46/182 = .25 \rightarrow 25\%$. %REC indexes the proportional degree to which an individual experiences the same measured states of affect over time. Note that although %REC and %DET often covary, %REC does not provide a measure of predictability on its own, but only in relation to %DET. For instance, randomly shuffling the example time-series in **Figure 2** would not alter %REC, but would significantly affect %DET. That is, %DET has the potential to provide the best picture of the degree to which the states of a system or data series repeated the same sequential dependent sequences over time (i.e., are governed by a deterministic or predictable dynamic process).

Statistical approach. Analysis of variance (ANOVA) and post hoc pairwise comparisons with a Bonferroni correction for

familywise error were used to assess differences in the metrics (i.e., mean, SD , RMSSD, %DET, and %REC) among the nine groups; 95% confidence intervals (CIs) are presented for mean values. Effect sizes of η^2 and their corresponding 90% CIs¹ are presented (Steiger, 2004). A power analysis conducted using G*Power (Faul, Erdfelder, Buchner, & Lang, 2009) revealed that a sample size of 100 per group would be sufficient to detect a medium effect size with statistical power to evaluate this hypothesis at the 0.95 level.

Results

Figure 3 presents a visual depiction of the mean, variability, and predictability measures for each of the nine groups. ANOVA results indicated that there were no overall mean differences among the nine groups $F(8, 891) = 0.45, p = .891, \eta^2 < 0.01$, 90% CI [0.00, 0.00] (see **Table 2**). As can be seen in **Figure 3a**, the mean was relatively stable over the nine groups.

ANOVA results indicated that there were overall differences among the nine groups for the variability measures (SD : $F(8, 891) = 163.77, p < .001, \eta^2 = 0.60$, 90% CI [0.56, 0.62]; RMSSD: $F(8, 891) = 106.21, p < .001, \eta^2 = 0.49$, 90% CI [0.45, 0.52]). Variability metrics tended to be in line with predictions such that Groups 1 through 3 had the lowest values while Groups 7 through 9 had the highest values (see **Figure 3a** and **Table 2**).

ANOVA results also indicated that there were overall differences among the nine groups for the predictability measures (%DET: $F(8, 891) = 392.96, p < .001, \eta^2 = 0.78$, 90% CI [0.76, 0.79]; %REC: $F(8, 891) = 230.63, p < .001, \eta^2 = 0.67$, 90% CI [0.65, 0.69]). The %DET and %REC measures followed similar patterns as the hypothesized results (see **Figure 3b**). Looking within low (Groups 1 through 3), medium (Groups 4 through 6), and high (Groups 7 through 9) variability groups, %DET and %REC tended to become larger as the data were repeated more often (i.e., more predictable). For example, %DET grew from 64.57 to 71.30 to 99.44 across Groups 4 through 6, respectively (see **Figure 3b** and **Table 2**). Although the SD was held constant for these groups, more instances of repeated data (i.e., predictability) occurred for Group 6 compared with Group 5 and more repeated data occurred for Group 5 compared with Group 4. However, when the SD was smaller (i.e., Groups 1 through 3), there was less discrepancy between %REC and %DET values among the groups. For example, the pairwise comparisons between Groups 1 and 2 and Groups 2 and 3 were no longer significantly different, $ps > .05$ (see **Table 2**). Nevertheless, the same pattern of results occurred throughout the data whereby as predictability increased, so did %DET and %REC values.

Discussion

The results demonstrate that the RQA metrics of %DET and %REC can be used to index the dynamics of variables like affect and provide different and new pieces of information about how affect changes over time compared with traditional measures of variability. RQA further differentiates simulated cases based on

¹ Because of the one-tailed probability test of ANOVA, 90% confidence intervals should be used to ensure that confidence intervals do not include 0 when p values are less than .05 (Steiger, 2004).

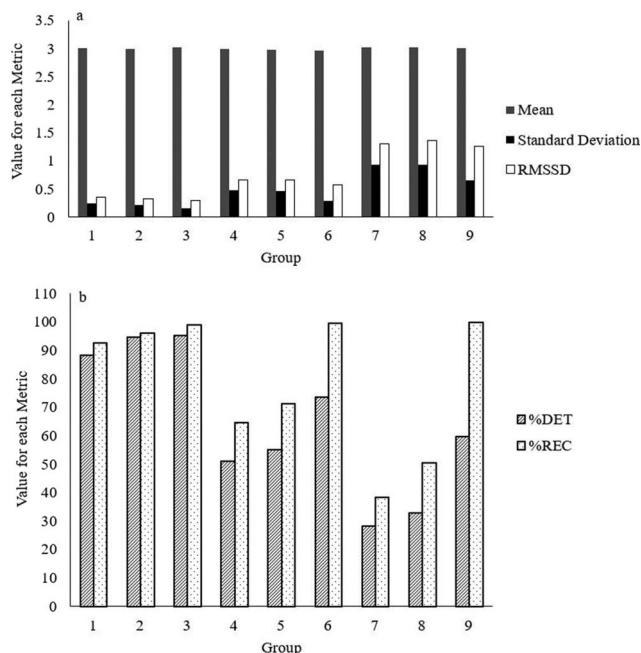


Figure 3. Mean, standard deviation, root mean squared successive difference (RMSSD), percent determinism (%DET), and percent recurrence (%REC) values by group. Note that the mean is about the same across all groups while standard deviation and RMSSD are each similar for Groups 1 through 3, then 4 through 6, and then 7 through 9. %DET and %REC help distinguish within these sets of groups.

metrics of predictability that consider the role of time in assessing patterns of affective experiences. Over the nine groups, mean levels were stable and did not distinguish between the groups. And, as noted by the other measures (*SD*, *RMSSD*, %DET, and %REC), the same mean level of affect can be associated with different levels of variability and predictability. This demonstrates that studies relying only on mean level of affect across time may be overlooking important information.

In addition, these results demonstrate that relying solely on the *SD* or *RMSSD* approach for assessing affective change is insufficient for capturing the finer details about the patterns of variation across time. For Groups 4 through 6 (see Figure 3a), *RMSSD*

(along with the means) was kept constant. Based on these values, previous researchers may have assumed that each of these groups were equal. However, once the RQA measures are taken into consideration, a substantial difference in affective patterning emerges and it becomes apparent that these groups are in fact not equal. When the same patterns of affect are repeated more often, there is an increase in %DET and %REC (see Figure 3b). For example, Group 4 and Group 6 have nearly identical means and *RMSSD* as one another. However, Group 4 is different from Group 6 because the same pattern of affect was repeated seven times (i.e., more predictability), as noted by %DET and %REC. Repeated patterns, and the predictability that follows, vary from person to person and are indicative of individual differences in affective experiences. This is critical because those differences in patterns of variation may have distinct implications for various outcomes. The addition of RQA metrics adds more information about the dynamic nature of affect and demonstrates how the relation between affective experiences over time may influence how affective profiles are categorized and understood.

There are several limitations to Study 1. First, although the normal distributions were intended to mirror similar means and *SDs* appearing in the affect literature (e.g., Jenkins et al., 2018), the smaller and larger *SDs* may be less likely. Similarly, the repeated nature of the data may or may not be ecologically valid. It is likely that repeated affect could occur from week to week with, for example, affect on Mondays looking similar to affect on other Mondays and affect on Fridays looking similar to affect on other Fridays. Therefore, the medium repeated data (in which data from 7 days are repeated across the next 7 days) may closely reflect this week by week repeated structure. Additionally, a no repeated condition (low repeat) was included that would mirror the natural environment if affect did not follow a specific repeated structure. It is important to note that even when data were not purposefully repeated, random repeated days could have occurred.

Regardless of these limitations, this study demonstrates that RQA measures may add more detailed information above and beyond measures of variability. Thus, the next step is to explore if this new methodology offers additional explanatory power in terms of its association with real world outcomes. If RQA helps determine the nonlinear dynamics that underlie affective change, researchers can potentially (a) build dynamical models of affective change and make corresponding predictions from that mode and

Table 2
Mean, SD, RMSSD, and Recurrence Quantification Measures by Group

Group	Group name	Mean	Variability		Predictability	
			<i>SD</i>	<i>RMSSD</i>	%DET	%REC
1	Low variability–Low predictability	3.01 ^a [3.00, 3.02]	.25 ^{ab} [.24, .26]	.36 ^a [.34, .38]	92.60 ^a [90.97, 94.24]	88.41 ^a [86.16, 90.65]
2	Low variability–Medium predictability	3.00 ^a [2.99, 3.02]	.22 ^{ab} [.21, .23]	.32 ^a [.30, .34]	96.13 ^{ab} [94.76, 97.51]	94.55 ^a [92.26, 96.84]
3	Low variability–High predictability	3.03 ^a [2.99, 3.06]	.16 ^b [.13, .18]	.30 ^a [.26, .35]	99.00 ^b [98.94, 99.06]	95.15 ^a [92.11, 98.19]
4	Medium variability–Low predictability	2.99 ^a [2.96, 3.01]	.48 ^c [.46, .50]	.67 ^b [.65, .70]	64.57 ^c [61.29, 67.86]	50.99 ^{bc} [48.33, 53.65]
5	Medium variability–Medium predictability	2.98 ^a [2.95, 3.02]	.46 ^c [.43, .49]	.67 ^b [.62, .72]	71.30 ^d [67.56, 75.04]	55.08 ^b [51.30, 58.85]
6	Medium variability–High predictability	2.97 ^a [2.90, 3.04]	.29 ^a [.25, .34]	.57 ^b [.48, .65]	99.44 ^b [99.33, 99.55]	73.62 ^d [68.31, 78.92]
7	High variability–Low predictability	3.03 ^a [2.98, 3.08]	.94 ^d [.90, .98]	1.30 ^c [1.23, 1.37]	38.30 ^e [35.33, 41.27]	28.13 ^e [26.68, 29.58]
8	High variability–Medium predictability	3.03 ^a [2.95, 3.10]	.94 ^d [.88, .99]	1.37 ^c [1.29, 1.46]	50.41 ^f [47.08, 53.73]	32.75 ^e [30.76, 34.73]
9	High variability–High predictability	3.01 ^a [2.87, 3.15]	.66 ^c [.56, .76]	1.27 ^c [1.08, 1.47]	99.73 ^b [99.63, 99.82]	59.62 ^c [55.02, 64.21]

Note. *SD* = standard deviation; *RMSSD* = root mean squared successive difference; %DET = percent determinism; %REC = percent recurrence. Column values with similar superscript letters indicate no significant difference ($p > .05$). The 95% confidence intervals are presented in parentheses.

(b) develop a machine-learning model or nonlinear function approximator (i.e., neural network) model to predict affective change. However, the key for either modeling approach is to first identify the degree to which there is a deterministic process underlying affective change (even if this process is chaotic or aperiodic). This latter goal is the aim of Study 2.

Study 2

Building on the findings of Study 1, in Study 2 we link affect mean, variability, and predictability to depressive and somatic symptom reports as most previous studies assessing affect variability have concentrated on similar outcomes (Gruber et al., 2013; Houben, Van Den Noortgate, & Kuppens, 2015; Human et al., 2015; Peeters et al., 2006). Capturing averages, variability, and predictability of affective experiences may provide researchers with a better understanding of how the intricacies of the affective experience influence mental and physical health. Furthermore, because these factors are not perfectly correlated, it may be advantageous to understand how they interact to predict certain health outcomes. The initial simulation study (Study 1) provided a foundational rationale for why these factors are important, and the following study (Study 2) applied the RQA methodology to real health outcomes.

Method

Participants. Study 2 used data from the “Daily Life Study” conducted from 2011 to 2014 at the University of Otago, New Zealand. Participants included 1,482 college students ($M_{\text{age}} = 19.76$, $SD_{\text{age}} = 2.43$). Sixty-seven percent of the participants were female. Participants were 78% Caucasian, 10% Asian, 5% Māori/Pacific Islander, 3% Indian, and 4% were another ethnicity or mixed ethnicity.

Procedure. Participants completed an initial survey asking about demographics and depressive symptoms. Participants then completed daily diaries for 13 consecutive days. The diaries consisted of several questionnaires including affect and stress measures. After the 13 consecutive days, participants then completed a follow up survey asking about physical health symptoms. All study procedures were approved by the University of Otago, New Zealand Ethics Committee.

Affect mean, variability, and predictability. State emotion adjectives were assessed each day for 13 days on a scale from 1 (*not at all*) to 5 (*extremely*) describing how much each of the words reflected how the participant felt that day. Nine NA words (nervous, dejected, irritable, hostile, sad, angry, unhappy, anxious, and tense) were averaged to create a daily NA value (Cronbach’s α range for each of the 13 days = .87 to .91) and nine PA words (happy, excited, cheerful, pleasant, calm, energetic, enthusiastic, content, and relaxed) were averaged to create a daily PA value (Cronbach’s α range for each of the 13 days = .88 to .92). Affect words were selected to capture a range of high to low intensities on the affective circumplex (Barrett & Russell, 1999). Day averages were then averaged over the 13 days to create an overall PA and NA mean value. Then, SD s and RMSSD values over the 13 time points were calculated for PA and NA with the formulas used in Study 1 to create measures of variability.² Finally, the RQA metrics, %DET and %REC, were calculated using the methods and

software described in Study 1 to create measures of predictability. All PA and NA daily mean values were rounded to the nearest integer value to allow for Categorical-RQA. These calculations resulted in mean, variability, and predictability values: NA mean (NA_{MEAN}), PA mean (PA_{MEAN}), NA SD (NA_{SD}), PA SD (PA_{SD}), NA RMSSD (NA_{RMSSD}), PA RMSSD (PA_{RMSSD}), NA %DET ($NA_{\%DET}$), PA %DET ($PA_{\%DET}$), NA %REC ($NA_{\%REC}$), and PA %REC ($PA_{\%REC}$). %REC and %DET were divided by 100 to be similar in range to the NA and PA means, SD s, and RMSSD values.

Distress. Distress was assessed each day with the question “Overall, how much stress (e.g., because of hassles, demands, or other stressors) have you been under today?” rated on a scale of 0 (*no stress*) to 4 (*a great deal of stress*). The response for each day was averaged over the 13 days and used as a control variable in all analyses as has been done in previous research on affect variability (e.g., Gruber et al., 2013).

Depressive symptoms. The Center for Epidemiologic Studies Depression Scale (CES-D; Radloff, 1977), a 20 item measure, was used to assess depressive symptoms in the initial survey. Items included statements such as “I did not feel like eating; my appetite was poor,” “I thought my life had been a failure,” and “My sleep was restless.” Items were scored using the following scale to assess how often the items reflected how they felt during the past week: rarely or none of the time (0), some or a little of the time (1), occasionally or a moderate amount of the time (2), or most or all of the time (3; positive items reverse scored). Scores were summed with higher values indicating higher levels of depressive symptoms (Cronbach’s $\alpha = .89$).

Self-reported somatic symptoms. During the follow up survey, participants were asked whether they felt like they had a cold or flu in the past 2 weeks (rated on a scale from 0 = *not at all* to 4 = *very*). Additionally, they were asked whether they “felt physically ‘run down’,” “felt tired,” and “felt refreshed when [they] woke up in the mornings” (each rated on a scale from 0 = *not at all* to 4 = *very*; the refreshed item was reverse coded). These four items were summed, with higher values reflecting a greater number of self-reported somatic symptoms.

Statistical approach. Depressive and somatic symptoms were used as the dependent variables in all analyses. Because of the dependent variables being count variables and skewed, poisson regression in Stata 15 (StataCorp, 2017) was used. Mean, variability, and predictability and the interactions among them were used as independent variables in a series of 10 models. Models 1 through 5 reflect the NA results while Models 6 through 10 reflect the PA results. Model 1 used mean and variability as the independent variables of interest. Model 2 added the Mean \times Variability interaction term to Model 1. Model 3 added the predictability metric to Model 2. Model 4 added the Mean \times Predictability interaction term to Model 4. Model 5 added the Variability \times Predictability and the Mean \times Variability \times Predictability interaction terms to Model 4. Models 6 through 10 mirrored Models 1 through 5 but used the PA variables. All predictor variables were

² We reanalyzed the data using autocorrelation as a measure of emotional inertia (see supplemental material B). The pattern of results remained similar with the exception of a few deviations. We describe these in supplemental material B.

mean centered to allow for ease of interpretation in interaction terms. Distress was adjusted for in all models. Effect sizes as well as 95% CIs are presented throughout.

Results

Descriptive statistics. Table 3 presents the means and *SD*s of the affect metrics as well as their associations. PA mean was higher than NA mean, $t(1,298) = 59.90, p < .001$, 95% CI of the difference [1.29, 1.37], Cohen's $d = 2.74$, 95% CI of Cohen's d [2.63, 2.85]. PA was more variable compared with NA as evidenced by the variability measures (PA_{SD} vs. NA_{SD}, $t(1,298) = 15.44, p < .001$, 95% CI of the difference [0.08, 0.10], Cohen's $d = 0.45$, 95% CI of Cohen's d [0.38, 0.53]; PA_{RMSSD} vs. NA_{RMSSD}, $t(1,298) = 13.87, p < .001$, 95% CI of the difference [0.09, 0.12], Cohen's $d = 0.41$, 95% CI of Cohen's d [0.33, 0.49]). NA was more predictable compared with PA as evidenced by the RQA measures (NA_{%DET} vs. PA_{%DET}, $t(1,298) = 11.52, p < .001$, 95% CI of the difference [6.18, 8.72], Cohen's $d = 0.40$, 95% CI of Cohen's d [0.32, 0.48]; NA_{%REC} vs. PA_{%REC}, $t(1,298) = 12.41, p < .001$, 95% CI of the difference [6.73, 9.25], Cohen's $d = 0.42$, 95% CI of Cohen's d [0.34, 0.49]). An interesting finding was that NA mean and affect variability measures were all positively associated while NA mean was negatively associated with affect predictability measures. In other words, individuals higher in NA were more likely to have variable affect but less likely to have predictable affect. On the other hand, PA mean was positively associated with affect predictability measures and negatively associated with affect variability measures. These PA findings indicate that individuals higher in PA have more predictable, but less variable affect.

The variability metrics of *SD* and RMSSD were very highly correlated with each other (NA: $r = .90, p < .001$, 95% CI [0.89, 0.91]; PA: $r = .82, p < .001$, 95% CI [0.80, 0.84]; see Table 3). This was also true for the predictability metrics (NA: $r = .84, p < .001$, 95% CI [0.82, 0.86]; PA: $r = .80, p < .001$, 95% CI [0.78, 0.82]; see Table 3). No other correlations between any of the variables were higher than these associations. Therefore, for ease of presenting the results, the following analyses use *SD* as the metric for variability and %DET as the metric for predictability. *SD* and %DET were selected over RMSSD because they resulted in greater differentiation among the nine groups in Study 1 (see Table 2).

Main effects of mean affect levels. All associations between mean levels of affect and the outcome variables were consistent with previous literature. Specifically, greater NA mean was associated with more depressive and somatic symptoms (see first rows in Tables 4 and 5). Higher PA mean was associated with fewer depressive symptoms and somatic symptoms (see eighth row in Tables 4 and 5).

Main effects of affect variability. Greater amounts of both PA and NA variability were associated with more depressive symptoms (NA_{SD}: $b = 0.19, z = 4.59, p < .001$, 95% CI [0.11, 0.27]; PA_{SD}: $b = 0.31, z = 7.64, p < .001$, 95% CI [0.23, 0.39]) and more somatic symptoms (NA_{SD}: $b = 0.22, z = 3.53, p < .001$, 95% CI [0.10, 0.34]; PA_{SD}: $b = 0.48, z = 8.17, p < .001$, 95% CI [0.36, 0.59]; see Tables 4 and 5 Models 1 and 6). In other words, individuals who had more affect variability had worse health outcomes.

Main effects of affect predictability. Higher NA predictability was associated with fewer depressive symptoms ($b = -0.16, z = -3.25, p = .001$, 95% CI [-0.26, -0.06]) and fewer somatic symptoms ($b = -0.25, z = -3.31, p = .001$, 95% CI [-0.39, -0.10]; see Tables 4 and 5 Model 3). PA predictability, on the other hand, was not associated with either depressive symptoms ($b = 0.00, z = 0.08, p = .938$, 95% CI [-0.10, 0.11]) or somatic symptoms ($b = 0.10, z = 1.40, p = .162$, 95% CI [-0.04, 0.25]; see Tables 4 and 5 Model 8). In summary, individuals with more predictable NA, had better health outcomes while their PA predictability did not matter.

Interactions between variability and mean levels. NA variability interacted with NA mean to predict depressive symptoms ($b = -0.69, z = -10.63, p < .001$, 95% CI [-0.81, -0.56]) but not somatic symptoms ($b = -0.15, z = -1.58, p = .114$, 95% CI [-0.34, 0.04]; see Tables 4 and 5 Model 2). Specifically, at higher levels of NA mean, less NA variability was associated with greater depressive symptoms (see Figure 4). PA variability interacted with PA mean to predict depressive symptoms ($b = 0.33, z = 4.17, p < .001$, 95% CI [0.18, 0.49]) but not somatic symptoms ($b = 0.16, z = 1.36, p = .175$, 95% CI [-0.07, 0.38]; see Tables 4 and 5 Model 7). At lower levels of PA mean, less PA variability was associated with more depressive symptoms (see Figure 5). This demonstrates that, as expected from previous characterizations of depressive symptoms (Watson, Clark, & Carey, 1988), individuals

Table 3
Mean, *SD*, and Pearson's Correlation of Affect Metrics

Affect Metric	Mean	<i>SD</i>	NA _{SD}	NA _{RMSSD}	NA _{%DET}	NA _{%REC}	PA _{MEAN}	PA _{SD}	PA _{RMSSD}	PA _{%DET}	PA _{%REC}
NA _{MEAN}	1.67	.47	.64***	.57***	-.52***	-.59***	-.36***	.17***	.16***	-.12***	-.12***
NA _{SD}	.41	.22		.90***	-.66***	-.76***	-.24***	.44***	.39***	-.30***	-.32***
NA _{RMSSD}	.52	.28			-.65***	-.69***	-.21***	.40***	.44***	-.27***	-.29***
NA _{%DET}	71.67	19.03				.84***	.26***	-.29***	-.29***	-.22***	-.23***
NA _{%REC}	58.36	21.11					.29***	-.34***	-.31***	.25***	.28***
PA _{MEAN}	3.00	.50						-.12***	-.10***	.06***	.09***
PA _{SD}	.50	.18							.82***	-.63***	-.75***
PA _{RMSSD}	.63	.25								-.59***	-.61***
PA _{%DET}	64.22	18.32									.80***
PA _{%REC}	50.38	17.25									

Note. *SD* = standard deviation; RMSSD = root mean squared successive difference; %DET = percent determinism; %REC = percent recurrence.
*** $p < .001$.

Table 4
Mean Level, Variability, and Predictability Metrics of Affect Predicting Depressive Symptoms

Predictor	Models									
	1	2	3	4	5	6	7	8	9	10
NA _{MEAN}	.56***	.61***	.60***	.60***	.58***					
NA _{SD}	.19***	.32***	.24***	.21***	.23***					
NA _{MEAN} × NA _{SD}		-.69***	-.66***	-.55***	-.51***					
NA _{%DET}			-.16**	-.21***	-.22***					
NA _{MEAN} × NA _{%DET}				.20	.08					
NA _{SD} × NA _{%DET}					.67**					
NA _{MEAN} × NA _{SD} × NA _{%DET}					-.16					
PA _{MEAN}						-.48***	-.48***	-.48***	-.48***	-.47***
PA _{SD}						.31***	.35***	.35***	.34***	.37***
PA _{MEAN} × PA _{SD}							.33***	.33***	.29**	.30**
PA _{%DET}								.00	-.00	-.01
PA _{MEAN} × PA _{%DET}									-.07	-.06
PA _{SD} × PA _{%DET}										.48*
PA _{MEAN} × PA _{SD} × PA _{%DET}										.39
Distress	.00	-.01	-.01	-.02	-.02	.18***	.18***	.18***	.18***	.18***
Constant	2.60***	2.66***	2.66***	2.67***	2.68***	2.37***	2.37***	2.37***	2.37***	2.38***
Observations	1,301	1,301	1,301	1,301	1,301	1,300	1,300	1,300	1,300	1,300

Note. SD = standard deviation; %DET = percent determinism.
* $p < .05$. ** $p < .01$. *** $p < .001$.

with worse psychological health outcomes had consistently high NA and/or consistently low PA.

Interactions between predictability and mean levels. NA predictability interacted with NA mean to predict somatic symptoms ($b = 0.40, z = 2.55, p = .011, 95\% \text{ CI } [0.09, 0.70]$) and, marginally, depressive symptoms ($b = 0.20, z = 1.96, p = .051, 95\% \text{ CI } [-0.00, 0.39]$; see Tables 4 and 5 Model 4). Specifically, at higher levels of NA mean, higher NA predictability was associated with marginally more depressive symptoms (see Figure 6a) and more somatic symptoms (see Figure 6b). PA variability did not interact with PA mean to predict depressive symptoms ($b = -0.07, z = -0.70, p = .483, 95\% \text{ CI } [-0.27, 0.13]$) or somatic symptoms ($b = -0.03, z = -0.21, p =$

.835, 95% CI [-0.32, 0.26]; see Tables 4 and 5 Model 9). In summary, NA predictability tended to be worse for health outcomes at higher levels of mean NA. This might represent someone who experiences a lot of day to day NA that occurs in regular patterns. In contrast, the effect of PA predictability did not change based on levels of PA mean.

Interactions between variability and predictability. NA predictability interacted with NA variability to predict depressive symptoms ($b = 0.67, z = 2.96, p = .003, 95\% \text{ CI } [0.23, 1.11]$) but not somatic symptoms ($b = 0.03, z = 0.10, p = .918, 95\% \text{ CI } [-0.63, 0.70]$; see Tables 4 and 5 Model 5). At higher levels of NA variability, more NA predictability was associated more depressive

Table 5
Mean Level, Variability, and Predictability Metrics of Affect Predicting Somatic Symptoms

Predictor	Models									
	1	2	3	4	5	6	7	8	9	10
NA _{MEAN}	.13***	.13***	.12***	.11***	.10**					
NA _{SD}	.22***	.24***	.12	.07	.06					
NA _{MEAN} × NA _{SD}		-.15	-.10	.11	.06					
NA _{%DET}			-.25***	-.31***	-.27***					
NA _{MEAN} × NA _{%DET}				.40*	.42*					
NA _{SD} × NA _{%DET}					.03					
NA _{MEAN} × NA _{SD} × NA _{%DET}					-.68					
PA _{MEAN}						-.19***	-.19***	-.19***	-.19***	-.17***
PA _{SD}						.48***	.48***	.55***	.55***	.57***
PA _{MEAN} × PA _{SD}							.16	.15	.13	.14
PA _{%DET}								.10	.10	.10
PA _{MEAN} × PA _{%DET}									-.03	-.02
PA _{SD} × PA _{%DET}										.65*
PA _{MEAN} × PA _{SD} × PA _{%DET}										.83
Distress	.16***	.16***	.15***	.15***	.15***	.19***	.19***	.19***	.19***	.19***
Constant	1.69***	1.70***	1.70***	1.71***	1.72***	1.64***	1.65***	1.64***	1.64***	1.66***
Observations	1,290	1,290	1,290	1,290	1,290	1,289	1,289	1,289	1,289	1,289

Note. SD = standard deviation; %DET = percent determinism.
* $p < .05$. ** $p < .01$. *** $p < .001$.

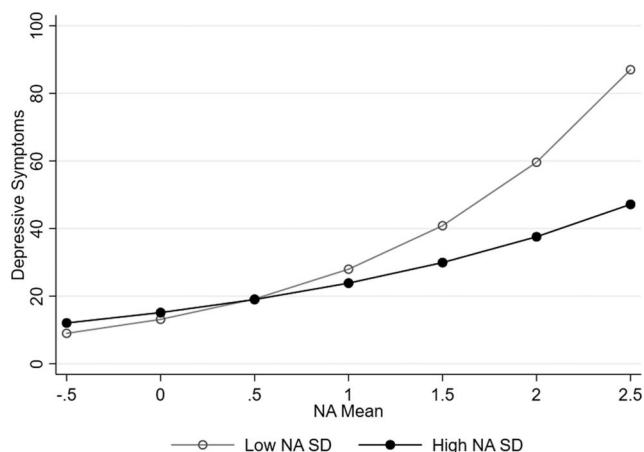


Figure 4. Interaction between NA_{MEAN} and NA_{SD} predicting depressive symptoms. Lines represent adjusted predictions and so may therefore exceed the range of the dependent variable in some cases.

symptoms (see Figure 7). PA predictability interacted with PA variability to predict depressive symptoms ($b = 0.48$, $z = 2.37$, $p = .018$, 95% CI [0.08, 0.87]) and somatic symptoms ($b = 0.65$, $z = 2.24$, $p = .025$, 95% CI [0.08, 1.22]; see Tables 4 and 5 Model 10). At higher levels of PA variability, more PA predictability was associated with more depressive symptoms (see Figure 8a) and more somatic symptoms (see Figure 8b). As previously shown (see Tables 4 and 5 Models 1 and 6), greater affect variability, whether it be NA or PA, was associated with worse outcomes. In these analyses of the interaction between variability and predictability, we see again that high levels of variability are detrimental and this is particularly true when affect is predictable.

Three-way interactions between variability, predictability, and mean. There were no three-way interactions between variability, predictability, and mean levels (all $ps > .05$; see Tables 4 and 5 Models 5 and 10).

Discussion

This study shows for the first time that measures of affect predictability add important new information in regard to health outcomes. Critically, measures of predictability differed from measures of variability in their association with affect valence. For example, this study revealed that individuals with higher average NA had more variable but less predictable affect. Conversely, those with greater mean PA had less variable but more predictable affect. This implies that individuals high in NA generally have greater and more erratic fluctuations in affect compared with those who have low NA. Additionally, individuals high in PA generally have less intense and more stable fluctuations in affect compared with those who have low PA.

These factors of variability and predictability were not only associated with PA and NA differentially, but they also had different associations with health outcomes. In general, more variable NA and PA was associated with worse outcomes (i.e., more depressive and somatic symptoms), consistent with past studies on similar topics (Gruber et al., 2013; Hardy & Segerstrom, 2017). On the other hand, our newly studied predictability metrics revealed

that NA patterns that are more consistent were associated with better outcomes (i.e., fewer depressive and somatic symptoms) while PA predictability mattered less.

Drawing on the discrepancies found between predictability and variability, it is also informative to examine how these factors interact with mean levels of affect to predict those same health outcomes. Although variability generally resulted in less favorable outcomes, at higher levels of mean NA, lower NA variability actually became associated with worse outcomes. For example, individuals with higher mean levels of NA had more depressive symptoms when they had less variable NA. It is possible that less variation for those high in mean NA indicates they are constantly in a negative state, which amplifies the detrimental effects that NA has on health. In contrast, someone with high mean NA and with greater NA variability may at least get to have some “breaks” from their typical high levels of negativity, which ultimately leads to more favorable health outcomes. For PA, variability had a similar effect whereby at lower levels of mean PA, less variability was associated with worse outcomes. For these individuals, less variability means that they are consistently low in positivity. In these ways, less variability has similar effects but depends on the valence and mean level of affect.

More NA predictability was generally associated with better outcomes. However, higher levels of NA predictability were associated with worse outcomes for those with high levels of NA. High predictability of NA for those already high in NA could signify that a person is “stuck” in a negative situation that they are continually experiencing. Someone who feels poorly now and expects to continue feeling poorly in the future will likely exhibit the least desirable outcomes, consistent with dynamic systems approaches to resilience (Pincus & Metten, 2010). PA predictability, on the other hand, did not have a main effect on the health outcomes nor did it interact with mean PA. However, the interaction between variability and predictability for both NA and PA was significant such that at high levels of variability, high predictability was worse for health. In summary, consistent NA predictability was generally better for health, except when NA mean was high. In addition, consistent PA and NA predictability were each worse for health when variability levels were high.

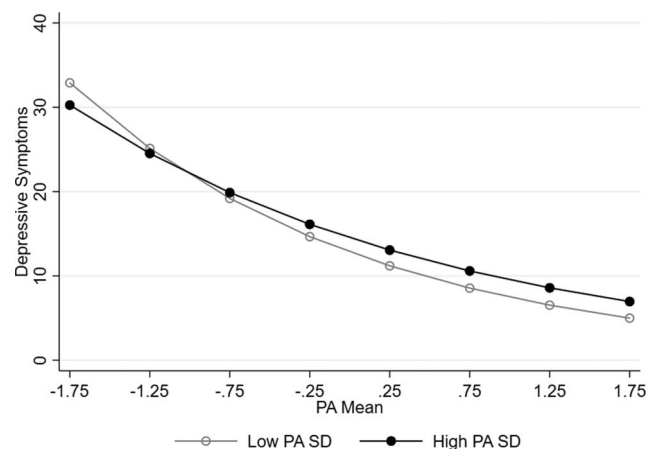


Figure 5. Interaction between PA_{MEAN} and PA_{SD} predicting depressive symptoms.

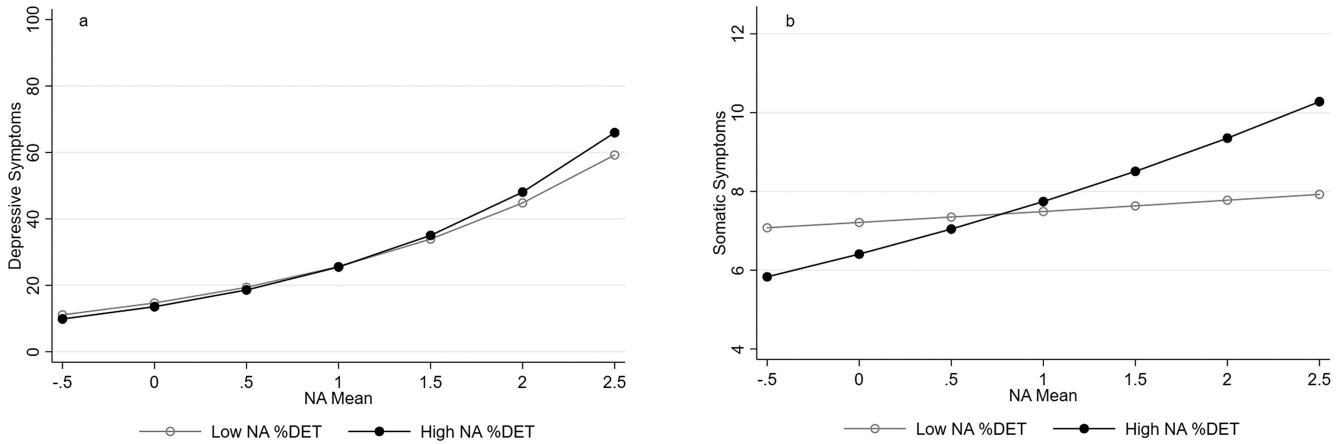


Figure 6. Interaction between NA_{MEAN} and $NA_{\%DET}$ predicting depressive and somatic symptoms. Lines represent adjusted predictions and so may therefore exceed the range of the dependent variable in some cases.

Based on these findings, researchers should consider using predictability metrics in addition to variability metrics, and this is especially true for NA. Furthermore, using interaction terms will provide opportunities to see how variability and predictability play divergent roles for different levels of mean affect. When we consider the nature of repeated patterns of affect (i.e., RQA metrics) in addition to measures of variability, the conclusions drawn about how affect changes wellness outcomes are altered. Assessing variability on its own is important, but additionally assessing predictability provides a clearer and more interesting picture about how fluctuations in affect influence mental and physical health.

Further, these findings demonstrate the importance of using multiple metrics of variability to test the robustness of findings. For example, *SD* and *RMSSD* were very highly correlated ($r_s > .80$, see Table 3) demonstrating that they are likely tapping the same construct, namely, affect variability. While these metrics are mathematically similar, they each independently reveal a more nuanced understanding of affect variability by providing conceptually different information about how affect changes over time. *RMSSD* determines changes in affect from one time point to the next, which illuminates how the

temporal ordering of affective events influences variability. On the other hand, *SD* represents the magnitude of change in affect but ignores the sequential order of events. On their own, they each provide only one piece of the interconnected puzzle, but when taken together we can understand how the magnitude and sequential changes of affect influence health.

There are a number of limitations in this work. First, we are unable to make causal conclusions about how affect is related to health, which leaves open the possibility of reverse causation. More variability could conceivably lead to higher levels of depressive symptoms, but it is also possible that having high levels of depressive symptoms could lead to more variable affect. Similarly, high somatic symptoms could have led to changes in affect variability and predictability. As in other observational studies on affect, our study design does not allow us to answer these types of directional questions. Nevertheless, this study adds substantially to the affect literature by demonstrating that predictability plays at least some role in the affect-health association. A second limitation is the relatively short time frame of data collection. It is possible that adding an additional week of data could have helped us more accurately assess predictability. However, data collection during a third week would add additional strain on participant adherence. Future research may consider examining different time spans of data collection. Additionally, it must be acknowledged that variability and predictability are highly correlated. This high correlation partially accounts for why there were sometimes no significant effects of PA predictability. However, these metrics are not perfectly correlated, which allows the RQA metrics to further differentiate certain cases and provide additional information. Another limitation is that our assessment methodology only focused on self-reported affect and did not take into account the variety of factors (e.g., stress, daily experiences) that may have contributed to fluctuations in affect. In future studies, it may be informative to monitor daily activities and contexts to paint a more refined picture of what drives these affective changes. Finally, the results of our study are not generalizable to the population at large because of our limited study sample. The participants were primarily Caucasian undergraduates, so our conclusions only apply to these types

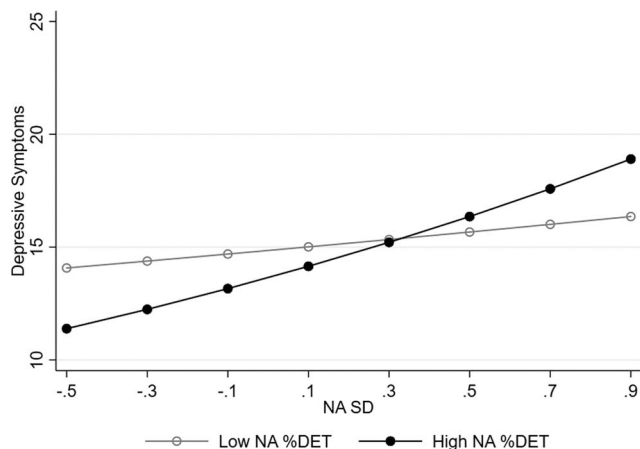


Figure 7. Interaction between $NA_{\%DET}$ and NA_{SD} predicting depressive symptoms.

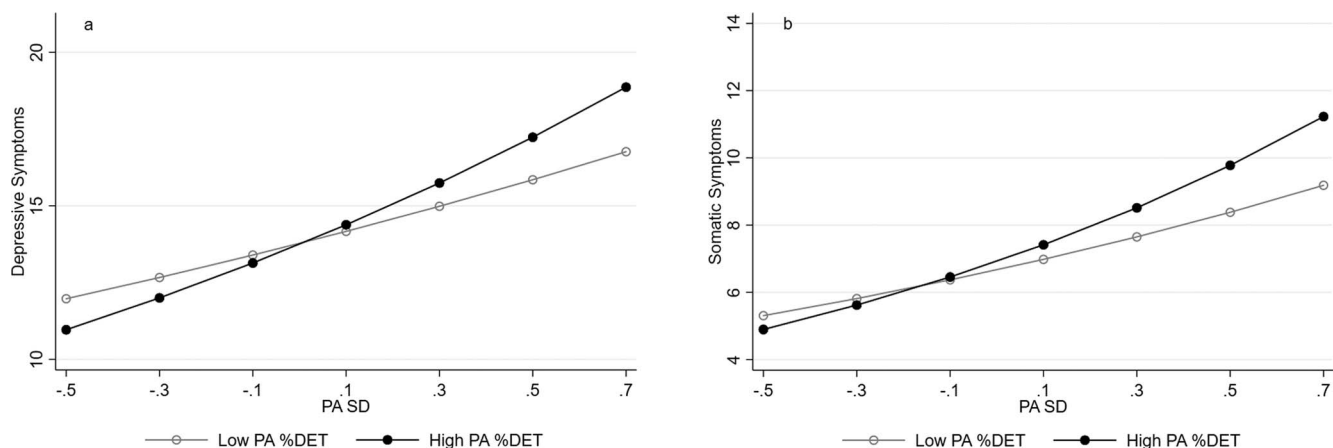


Figure 8. Interaction between $PA_{\%DET}$ and PA_{SD} predicting depressive symptoms and somatic symptoms.

of individuals. However, these methods may be extended to other populations and so future research may address this gap.

There are important conclusions about the divergent impacts of variability and predictability that deserve consideration. In general, it was found that the most desirable outcomes stemmed from those who had high PA, low NA, low variability, and high predictability. PA mostly did not interact with variability or predictability (high PA was good in almost all scenarios), but there were some interesting findings in regard to NA. The results indicate that for individuals high in NA mean, it is good to have high variability (possibly because one gets breaks from the negativity) and it is bad to have high predictability (possibly because the negativity is unchangeable). As demonstrated by these findings, the additional predictability measures add explanatory depth to how the dynamics of affect are associated with health.

These studies are the first to demonstrate how RQA metrics (%DET and %REC) can add interesting new information about the association between affective experiences and health outcomes by elucidating temporal patterns that are often overlooked when relying only on variability or mean levels. When assessing psychosomatic connections, the vast majority of studies rely on indicators of mean affect (Pressman & Cohen, 2005). Our findings should implore future researchers to also consider the role of predictability as well as encourage the growing interest in affect variability and health. Affect unfolds over time, so the consideration of temporal patterns is critical to capture the dynamic nature of affective experiences (Jenkins, 2017).

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