

Hair cortisol and its association with psychological risk factors for psychiatric disorders: a pilot study in adolescent twins

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Abstract

Measuring cortisol in hair is a promising method to assess long-term alterations of the biological stress response system, and hair cortisol concentrations (HCC) may be altered in psychiatric disorders and in subjects suffering from chronic stress. However the pattern of associations between HCC, chronic stress and mental health require clarification. Our exploratory study (1) assessed the association between HCC and perceived stress, symptoms of depression and neuroticism, and the trait extraversion (as a control variable) and (2) made use of the twin design to estimate the genetic and environmental covariance between the variables of interest. Hair samples from 109 (74 female) subjects (age range 12–21 years, mean 15.1) including 8 monozygotic (MZ) and 21 dizygotic (DZ) twin-pairs were analysed. Perceived stress was measured with the Perceived Stress Scale and/or the Daily Life and Stressors Scale, neuroticism and extraversion with the NEO-Five Factor Inventory or the Junior Eysenck Personality Questionnaire, and depressive symptoms with the Somatic and Psychological Health Report. We found a modest positive association between HCC and the three risk-factors perceived stress, symptoms of depression, and neuroticism ($r=0.22-0.33$) but no correlation with extraversion (-0.06). A median-split revealed that the associations between HCC and risk factors were stronger ($0.47-0.60$) in those subjects with HCC >11.36 pg/mg. Furthermore, our results suggest that the genetic effects underlying HCC are largely shared with those that influence perceived stress, depressive symptoms and neuroticism. These results of our proof of principle study warrant replication in a bigger sample but raise the interesting question of the direction of causation between these variables.

Keywords: Hair cortisol, Stress, Twin-Study, Depression, Neuroticism; Genetics

Introduction

The adequate treatment of psychiatric disorders is burdened by their complex aetiology. The joint action of multiple genetic and environmental factors leads to onset of illness, which usually starts with unspecific symptoms in adolescence (Van Os et al., 2008; Lesch 2004). Thus, the identification of biological and psychological markers that underlie the development and progression of illness and link to relevant pathophysiology is of crucial importance to understanding the mechanisms leading to psychiatric disorders and hence to their prevention and treatment.

Chronic stress and an impaired ability to cope with it, both at the psychological and biological level, are thought to play a key role in the development and maintenance of psychiatric disorders (de Kloet et al., 2005). The most common biological marker to measure stress-response is cortisol, which is released by the hypothalamic-pituitary-adrenal (HPA) axis. The HPA axis is the main endocrine mediator of the stress-response, and it is modulated by and exerts feedback on brain regions involved in psychiatric disorders e.g. the amygdala, the hippocampus, and the prefrontal cortex (de Kloet et al., 2005; Dedovic et al., 2009; Herman et al., 2003).

Studies in healthy population samples show an increase in cortisol in blood, urine and saliva during and after exposure to short-term stressors and a decrease once the stressor disappears (Miller et al., 2007). The regulation of the HPA axis has been reported to be altered in patients suffering from depression, schizophrenia and anxiety disorders supporting the hypothesis of impaired ability to cope with stress in psychiatric disorders (Bradley & Dinan, 2010; Faravelli et al., 2012; Herbert, 2013; Jones & Moller, 2011). The measurement of cortisol in blood, saliva or urine is a sensitive method to assess acute stress-responses and reflects short-term cortisol

release. However, single cross-sectional samples may not provide a valid measure of chronic stress and long-term alterations of HPA-functioning as they are sensitive to daily circumstances such as circadian rhythm, nutrition, etc.

A relatively new method which enables the efficient assessment of cortisol-release over a longer period is its measurement in hair of the scalp (Stalder et al., 2012). Scalp hair grows at an average rate of one cm per month so cortisol assayed from a 3cm hair segment closest to the scalp likely reflects the average release over the past 3 months and is therefore a potential biomarker of long-term altered HPA functioning. Initial studies showed altered HCC in unaffected adults with high levels of chronic stress and in cases suffering from psychiatric disorders (including bipolar disorders, depression and anxiety disorder) compared with controls (reviewed in Herane Vives et al., 2015). More recent studies found similar associations in children and adolescents for objective stress and HCC (Rippe et al., 2016; Vliegthart et al., 2016; Simmons et al. , 2016)

These alterations in HCC may reflect causal mechanisms such as genetic determinants which jointly influence the level of HCC, and moderate affective symptoms and perceived stress (pleiotropy), or may be a direct consequence of the pathogenesis such that affective symptoms and perceived stress increase HCC; or both mechanisms may operate . However, few studies have assessed the association between HCC, affective symptoms and perceived stress. Furthermore, to our knowledge no study has assessed the association between HCC and the personality traits of neuroticism and extraversion, where the first is a risk factor for the development of psychiatric disorders and the latter is not (Kendler et al., 2006; Van Os & Jones, 2001)

Many previous studies assessing the relationship between perceived stress and HCC found no significant or low correlations (Staufenbiel et al., 2013; Heinze et al., 2016). This so-called lack of psychoendocrine variance, with endocrine response not coinciding with psychological stress might be due to most studies assuming a linear association between hair cortisol and psychological markers. It is possible though that a pathological threshold exists for cortisol as it does for other metabolite biomarkers such as creatinine, where only higher values are associated with renal damage and lower levels have no clinical implications (Singhal & Saha, 2014).

The primary aim of our pilot study was to assess the association of HCC with psychological risk factors for psychiatric disorders - perceived stress, symptoms of depression and neuroticism and also the personality trait extraversion for which there is no expected relation with cortisol. Our secondary aim was to explore the genetic and environmental causation of these associations making use of the MZ and DZ twins in our sample.

Materials and methods

Sample

Our sample comprised 135 adolescent or young adult twins (95 females) who had taken part in either the Brisbane Longitudinal Twin Study (BLTS; Wright and Martin, 2004), or the Queensland Twin Imaging Study (QTIM study; Blokland et al., 2011) between 2009 and 2012. In the BLTS, psychiatric symptoms are assessed at ages 12, 14 and 16 years (Hansell et al., 2012). The QTIMS is a follow-up study of the BLTS cohort, which involves investigation of twins aged 21-28 years. Zygosity was initially determined by a combination of standard questions

(Martin & Martin, 1975) and a photograph of the twin pairs and subsequently confirmed by extensive genotyping. The sample consisted of 8 monozygotic (MZ) and 24 dizygotic (DZ) complete twin pairs and 66 individual twins and 5 siblings. In both the MZ and DZ groups, female twin pairs were predominant.

Data of those subjects with a weight of < 7.5mg of hair cortisol were excluded (n = 26 which left 8 MZ pairs, 21 DZ pairs, and 51 unpaired twins or sibs for further analysis (for details see *Hair Cortisol Analysis*). The mean age of the sample was 15.1 years (sd=1.99; range 12–21 years).

We are well aware that this is too small a sample for reliable heritability estimation but our principal interest is in the relation between HCC and our psychological variables and the genetic and environmental causes of any such correlation for which the power of analysis is greater given the multivariate nature of the data.

Self-report measures

For participants aged 12 and 14 years, perceived stress in the past month was measured using the 10-item Perceived Stress Scale (PSS; Cohen et al., 1983). The trait variables Neuroticism and extraversion were measured using the Junior Eysenck Personality Questionnaire (JEPQ; Eysenck 1972) 20 items for neuroticism, 24 items for extraversion. For participants aged 16 years or older, perceived stress was measured using the 30-item Daily Life and Stressors Scale (DLSS;(Kearney et al., 1993), and neuroticism and extraversion were measured using the NEO-Five Factor Inventory revised version (Neo-FFI-R; McCrae & Costa Jr., 2004) each with 12 items for neuroticism and extraversion. Depressive symptoms at all ages were assessed using the

34-item Somatic and Psychological Health Report (SPHERE; Hickie et al., 2001; Hansell et al., 2012).

Hair Cortisol Analysis

Sample collection and preparation were performed as described elsewhere (Kirschbaum et al., 2009). In brief, a swatch of hair about 3mm in cross-section from the back of the head (posterior vertex position) was cut with fine scissors as close as possible to the scalp. As hair grows at an average speed of one cm per month, the 3 cm segment proximal to the scalp can be assumed to reflect cortisol secretion during the preceding 3 months (LeBeau et al., 2011) and our sample was trimmed from the distal end to this length. Hair-washing and steroid extraction were performed as described in (Davenport et al., 2006). The hair samples were then micro centrifuged at 10,000 rpm for 2 min. A 1 ml volume of the clear supernatant was then transferred into a new cryo vial in order to allow the alcohol to evaporate at 60 °C under a constant stream of nitrogen until the samples were completely dry. Finally, 0.4 ml of phosphate buffer was added and the tube was vortexed for 15s before cortisol was assayed using a commercial immunoassay kit as described by Kirschbaum et al. (2009).

Ideally, the HCC assays should be performed on a hair sample weighing at least 10 mg. However, for some individuals (n=46, mainly males) this amount of hair had not been collected. Preliminary analysis of the entire sample of observations revealed a strong negative correlation ($r=-0.49$) between the weight of the hair sample being analysed and the HCC. We have no explanation for this effect, which was observed in men and women and both did not differ in average cortisol levels ($t_{133}=0.24$, $p=0.81$). On inspection this relation appeared to apply only to the smallest samples and to asymptote above a sample weight of 7.5mg so we used this as a lower limit for inclusion in the study.

Data Analysis

Due to skewness, hair cortisol (mean = 14.07 pg/mg, SD= 6.48, range 1.61 - 41.33) was log transformed for all analyses.

To compare neuroticism scores between participants, the neuroticism and extraversion sum-scores of the NEO-FFI and the JEPQ-scores were separately z-transformed and then combined. For the assessment of perceived stress measured by the PSS (Cohen et al., 1983) and the DLSS (Kearney et al., 1993) and depressive symptoms measured by the SPHERE (Hickie et al., 2001), item response theory (IRT) analyses were performed (Wray et al., 2008) to harmonize data. The details and advantages of IRT analysis are described elsewhere (Rietschel et al., 2014).

Preliminary analysis of the association between hair cortisol and the different psychological measures (for example, ratings of perceived stress and neuroticism) was complicated by the fact that our twin study contained two children from the same family, leading to non-independent observations. As such, we randomly split the data so that no family member would be in the same group, Spearman correlations were then performed to examine the relationship between neuroticism, extraversion, symptoms of depression, perceived stress and hair-cortisol, using pairwise deletion of missing data. Afterwards correlations were meta-analytically combined using Hedges method which adjusts estimated standard errors and sample size. To distinguish subjects with high versus low hair cortisol a median split was carried out.

Model Fitting

To allow use of data from all individual twins, including those without co-twins, and from participants with missing outcome measures, structural equation models (SEM) were fitted using the full information maximum likelihood (FIML) method implemented in Mx (Neale, 2015). Details of the twin design and analytical methods, including assumption testing and multivariate modelling (below), are described elsewhere (Neale & Cardon, 1992). For multivariate analysis a Cholesky decomposition for the ACE model was compared with Cholesky decompositions for simpler AE, CE, and E models. Despite evidence for a nonlinear relationship between HCC and the affect variables, we carried out the variance components analysis assuming only linear relations given our small sample size. The fit of each model was assessed by the differences in log likelihood between the sub and the full models. For all models, sex and age were fitted as fixed effects.

Results

Relation between HCC and our affect measures

Preliminary inspection of scatterplots of HCC with behavioral measures suggested nonlinear relationships. We therefore performed a median split at $HCC=11.36\text{pg/mg}$ and estimated correlations for the high and low halves of the sample as well as the whole. Table 1 reveals the following: a) correlations between HCC and the three risk-factors perceived stress, symptoms of depression, and neuroticism are modest and significant in the whole sample but high and significant for the group of high HCC ($HCC>11.36\text{pg/mg}$); b) there are no significant associations between HCC and the three risk factors in the low-cortisol group ($HCC\leq 11.36\text{pg/mg}$); c) there is no significant association between extraversion and HCC.

“Please insert Table 1 about here”

Twin correlations

The twin-correlations between the psychological variables perceived stress, neuroticism, extraversion, and HCC are displayed in Table 2. As expected, correlations between MZ twins are higher than for DZ twins for neuroticism, perceived stress and extraversion. However, unexpectedly the correlation for HCC is higher in the DZ ($r=.63$) than in the MZ ($r=.20$) group, though not significantly so given these small sample sizes.

“Please insert Table 2 about here”

Genetic covariance between HCC and affect variables

Rather than focus on the zero estimate of heritability for HCC (which is likely a stochastic result due to low numbers), we are more interested in evidence for genetic covariance between HCC and the psychological variables. To investigate this we fit a series of Cholesky decomposition models including variously A, C and E as sources of covariation. The aim of the analyses was to determine whether genetic influences specific to HCC exist after perceived stress, neuroticism and depressive symptoms have been accounted for. Therefore, perceived stress was used as the first, depressive symptoms as the second, neuroticism as the third and HCC as the fourth latent factor. We began by fitting an ACE Cholesky model but no significant worsening of the fit was observed when either the C matrix was fixed to zero ($\chi^2_{10} = 4.51$) or the A matrix was fixed to zero ($\chi^2_{10} = 4.07$). In fact both A and C matrices could be fixed to zero without fatally affecting fit ($\chi^2_{20} = 18.22$) underlining the fundamental lack of power of our experiment. Nevertheless, the point estimates from the ACE Cholesky decomposition are of

great interest. In Table 3 shows that the estimates of factor loadings for the A matrix are substantial, those for the C matrix are negligible except for C(4,4), the specific loading on HCC, reflecting the low r_{mz} for HCC noted above. Indeed, even the cross loadings for the E matrix are considerably smaller than their corresponding elements in the A matrix. Notwithstanding the deficiencies in power of our study, our results point strongly to the predominant importance of genetic factors in shaping the correlations observed between HCC and the psychological variables.

“Please insert Table 3 about here”

Nevertheless, because A and C estimates tend to be intimately confounded in the twin model, and because the point estimates from Figure 1 suggest that the dominant source by far is A, for simplicity, in Figure 1 we show the standardized path coefficients for the AE Cholesky decomposition (these must be squared to obtain variance components), remembering that, to a small extent, the A estimates are confounded with some shared environmental effects. Under this simplified model, genetic factors would account for more than half the total variance for all four variables. Further, the first genetic factor (A1), which loads primarily on perceived stress and accounts for 63.3 % of its variance, also accounts for 32.2% of the variance in depressive symptoms, 20 % of the variance for neuroticism and 11.4% of the variance for HC. The second genetic factor (A2), which loads primarily on depressive symptoms (35.7%), also accounts for 24.2 % of the variance in neuroticism and 15.4% in HCC. The third genetic factor (A3), which loads primarily on neuroticism (23.2%) also accounts for 11.6% of the variance in HC. This leaves a specific genetic contribution (A4) to HCC that accounts for 24.9 % of its variance. Decomposition of the nonshared environmental covariance shows that the first factor (E1) accounts for 36.7 % of the variance for neuroticism

and also accounts for 7.6 % of the variance in depressive symptoms, 9.6 % of the variance in perceived stress and 4% of the variance for HC. The second factor (E2), which loads on depressive symptoms (24.6 %), also accounts for 6% of the variance in perceived stress and 0.3% of the variance for HCC. The third factor (E3) loads on neuroticism and accounts for 17% of its variance and 3% on HCC. The specific nonshared environmental contribution to HCC (E4) accounts for 29.9 % of its variance.

“Please insert figure 1 about here”

Genetic correlation between HCC and affect variables

Also for simplicity, the genetic and environmental correlations between the four variables, derived from the AE Cholesky analysis, are shown in Table 4.

We can see that the correlations between the three psychological variables largely derive from genetic sources. The correlations between these three and HCC are also much more strongly genetically than environmentally influenced.

“Please insert table 4 about here”

Discussion

The aim of this exploratory study was to assess the association between hair cortisol concentration (HCC) and three affect variables we might reasonably expect to be associated with perceived stress, symptoms of depression, neuroticism. We also included extraversion as a control variable with no a-priory expectation of association. Our sample included a small

number of complete monozygotic and dizygotic twin pairs and these enabled us to explore the genetic and environmental contributions to covariation between our variables. In particular, we were interested in the causes of covariation between HCC and the affect variables. Finally we were interested in whether relationships between HCC and the affect variables are linear and we found some evidence that the relationship is much stronger in those with high HCC, suggesting there might be some threshold effect.

Our results show moderate but significant associations between HCC and neuroticism, perceived stress, and depressive symptoms, respectively ($r=0.22-0.33$). Interestingly, splitting the sample into low versus high-HCC revealed that lower levels of HCC seem to be unrelated to these psychological variables whereas the associations in the high-HCC group are substantial and significant ($r=0.47-0.60$). Furthermore, our study suggests strongly that these associations are genetic in origin. As expected however, extraversion is not associated with HCC in either the high or the low-HCC group and this accords with previous studies of other cortisol markers (Munafo 2005; Wilson et al., 2015).

Positive correlations between perceived stress and cortisol markers have been reported previously, even though the reports are not consistent (Staufenbiel et al., 2013; Vanaelst et al., 2012).. These inconsistencies together with our finding that the association is stronger in the high-HCC group might indicate a potential pathological threshold for HCC although, mindful of our small sample size, we have to regard this finding most tentatively and clearly it needs to be replicated in a much larger dataset. To our knowledge, no previous study has suggested the existence of such a threshold, although others have suggested a more complex non-linear relationship (Wells et al., 2014).

In monkeys, the heritability for HCC has been estimated at 30% (Fairbanks et al., 2011), while other twin studies suggest that genetic factors account for 62% of the variance of saliva and serum cortisol (Bartels et al., 2003). Due to small sample size we are not able reliably to assess the heritability of HCC; the DZ correlation for HCC is high ($r=0.63$) and most likely the very low MZ correlation ($r=0.20$) reflects our having only 8 MZ twin pairs; we are currently collecting a much larger sample which will give us a more reliable estimate.

Most pertinent to the main aim of our study, our results suggest that the association between HCC and the affect variables seems to be largely driven by genetic factors. This supports the hypothesis of a common genetic association between all four measures, whereas environmental factors do not seem to contribute much to covariation. This is consistent with our results from a much larger twin sample (of which the present study uses a subsample) in which we estimated high genetic covariances between neuroticism, depression and perceived stress (Rietschel et al., 2014).

The most obvious limitation of our study is the small sample size; this will be rectified in our forthcoming study with a sample six times larger than here. The much greater limitation however, is that even if the relationships we observe here are replicated with increased significance it will not elucidate the direction of causation in the correlation between these variables. While it is possible that perceived stress is environmentally induced and in turn induces high HCC, the reverse could also be true. Another suggestion from our present results though is compatible with a pleiotropic explanation, in which the same genes simultaneously influence both HCC and the psychological variables. Elucidation of this conundrum might be advanced by sensitive longitudinal studies, or as a by-product of large scale GWAS studies which enable construction of powerful instrumental variables for use in Mendelian

randomization analyses (Richmond et al., 2016). Larger samples of MZ twins with varying degrees of discordance for HCC and the other measures here may also shed light on the nature and direction of causation between these variables.

In conclusion our pilot-study provides new evidence that HCC is related to psychological variables, which elevate the risk for psychiatric disorders but not extraversion. Further, these correlations are seen most strongly in the higher range of HCC values suggesting some sort of threshold effect. Most interestingly, analysis of these associations in MZ and DZ twins suggest that they are largely driven by genetic rather than either familial or idiosyncratic environmental influences. However, the direction of causation between these variables remains unclear.

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Conflicts of interest

None.

Ethical standards

The authors assert that all procedures contributing to this work comply with ethical standards of the relevant national and institutional committees on human experimentation and with the Helsinki Declaration of 1975, as revised in 2008. The project was approved by the QIMR Human Research Ethics Committee.

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Tables

Table I Correlations (95% C.I.) between psychological variables and HCC in the whole sample (n=109), and after median split (HCC >\≤11.36pg/mg)

	Whole sample (n=109)	High-cortisol (n=53)	Low-cortisol (n=56)
Stress	0.22 (0.03, 0.41)	0.47 (0.23, 0.78)	0.09 (-0.19, 0.36)
Depression	0.33 (0.15, 0.53)	0.60 (0.41, 0.97)	0.03 (-0.24, 0.30)
Neuroticism	0.28 (0.10, 0.48)	0.49 (0.26, 0.81)	0.15 (-0.12, 0.42)
Extraversion	-0.06 (-0.13, 0.25)	-0.07 (-0.21, 0.35)	0.12 (-0.15, 0.39)

Table 2: Pearson correlations between all variables in DZ (21 pairs) and MZ (8 pairs) twins (MZ in lower, DZ in upper triangle) corrected for age and sex

	Twin 1					Twin 2					
	Stress	Depression	Neuroticism	Extraversion	Haircortisol	Stress	Depression	Neuroticism	Extraversion	Haircortisol	
	1	0.53	0.66	-0.26	0.19	0.24	-0.04	0.64	0.05	-0.63	
	0.68	1	0.46	-0.09	0.36	0.35	0.05	0.11	-0.03	0.21	
	0.84	0.81	1	-0.10	-0.10	0.25	-0.05	0.04	0.09	-0.01	
Twin1	Extraversion	0.23	0.01	0.48	1	-0.09	-0.23	-0.34	-0.31	-0.15	0.19
	Hair cortisol	0.54	0.87	0.72	0.31	1	-0.10	0.17	-0.09	0.09	0.63
	Stress	0.61	0.74	0.71	-0.04	0.52	1	0.73	0.69	-0.08	0.28
Twin2	Depression	0.72	0.73	0.86	0.17	0.56	0.90	1	0.76	-0.15	0.29
	Neuroticism	0.53	0.68	0.75	0.05	0.46	0.80	0.91	1	-0.18	0.31
	Extraversion	0.40	0.35	0.32	0.08	0.45	0.48	0.66	0.30	1	-0.08
	Hair cortisol	0.15	0.09	0.25	0.65	0.20	0.15	0.10	0.13	-0.42	1

Table 3 Cholesky decomposition for ACE model. Latent factor loadings are standardized to unit variance and must be squared to obtain standardised variance components

A	1	2	3	4
Stress	0.77			
Depression	0.57	0.60		
Neuroticism	0.43	0.49	0.48	
Hair cortisol	0.37	0.37	0.34	0.26
C				
Stress	0.20			
Depression	0.04	0.02		
Neuroticism	0.11	0.02	0.09	
Hair cortisol	0.21	0.05	0.18	0.40
E				
Stress	0.61			
Depression	0.28	0.49		
Neuroticism	0.31	0.25	0.41	
Hair cortisol	0.19	0.08	0.17	0.49

Table 4: Additive Genetic and Unshared Environmental Correlations between psychological variables and HCC corrected for age and sex

		Environmental			
		Stress	Depression	Neuroticism	Hair Cortisol
Genetic	Stress		0.32	0.56	0.15
	Depression	0.86	-	0.34	0.01
	Neuroticism	0.77	0.98	-	0.10
	Hair Cortisol	0.37	0.53	0.43	-

Figure 1

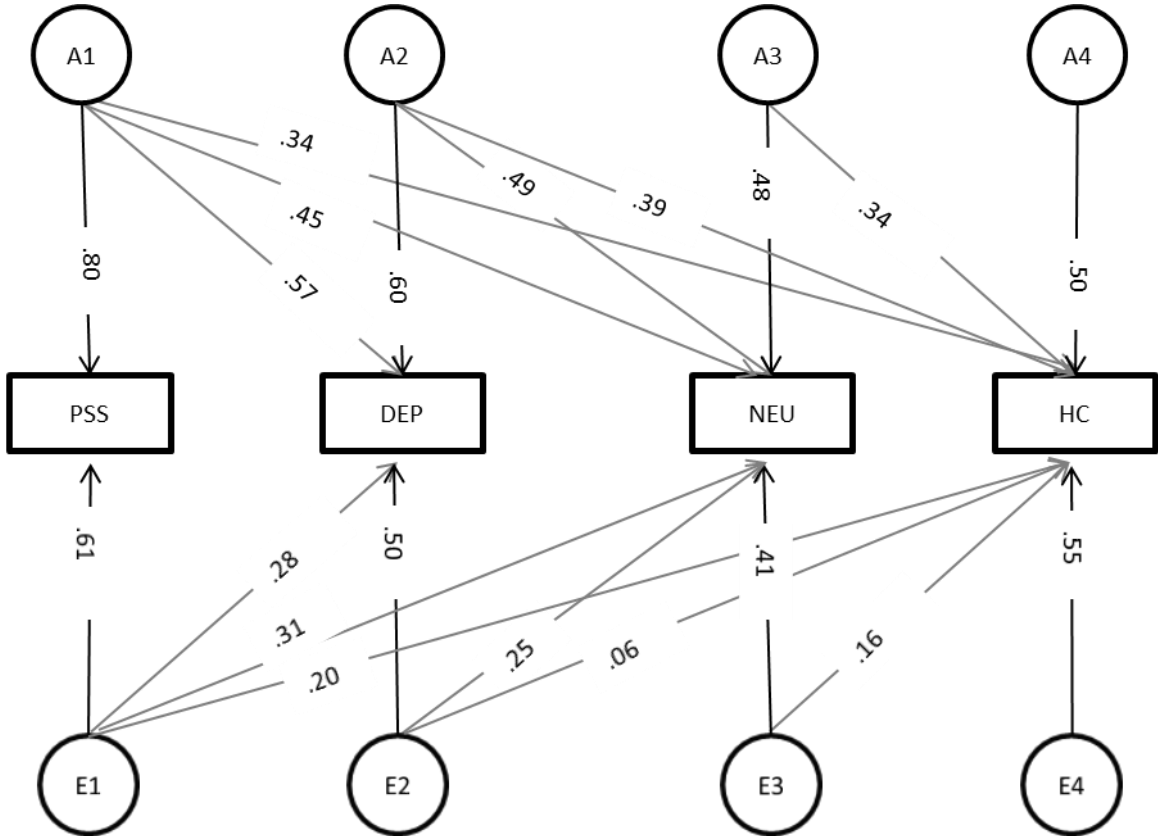


Figure 1. Cholesky decomposition for AE model. Latent factor loadings are standardized to unit variance and must be squared to obtain standardised variance components. A1–A4 additive genetic factors, E1–E4 unique environmental factors.