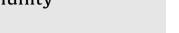
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Discovery of serum biomarkers predicting development of a subsequent depressive episode in social anxiety disorder

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ABSTRACT

Although social anxiety disorder (SAD) is strongly associated with the subsequent development of a depressive disorder (major depressive disorder or dysthymia), no underlying biological risk factors are known. We aimed to identify biomarkers which predict depressive episodes in SAD patients over a 2-year follow-up period. One hundred sixty-five multiplexed immunoassay analytes were investigated in blood serum of 143 SAD patients without co-morbid depressive disorders, recruited within the Netherlands Study of Depression and Anxiety (NESDA). Predictive performance of identified biomarkers, clinical variables and self-report inventories was assessed using receiver operating characteristics curves (ROC) and represented by the area under the ROC curve (AUC). Stepwise logistic regression resulted in the selection of four serum analytes (AXL receptor tyrosine kinase, vascular cell adhesion molecule 1, vitronectin, collagen IV) and four additional variables (Inventory of Depressive Symptomatology, Beck Anxiety Inventory somatic subscale, depressive disorder lifetime diagnosis, BMI) as optimal set of patient parameters. When combined, an AUC of 0.86 was achieved for the identification of SAD individuals who later developed a depressive disorder. Throughout our analyses, biomarkers yielded superior discriminative performance compared to clinical variables and self-report inventories alone. We report the discovery of a serum marker panel with good predictive performance to identify SAD individuals prone to develop subsequent depressive episodes in a naturalistic cohort design. Furthermore, we emphasise the importance to combine biological markers, clinical variables and self-report inventories for disease course predictions in psychiatry. Following replication in independent cohorts, validated biomarkers could help to identify SAD patients at risk of developing a depressive disorder, thus facilitating early intervention.

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1. Introduction

Social anxiety disorder (SAD; also referred to as "social phobia") is among the most common anxiety spectrum disorders with a 12month prevalence ranging between 2% and 7% (Kessler et al., 2012; Wittchen et al., 2011). Defined by a marked fear of social situations, the affected individual avoids situations associated with exposure to possible scrutiny by others (American-Psychiatric-Association, 2013). The age of onset is usually in childhood or adolescence (Ballenger et al., 1998). Despite the associated distress

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http://dx.doi.org/10.1016/j.bbi.2015.04.011 0889-1591/© 2015 Published by Elsevier Inc. and impairment, only half of the patients fulfilling diagnostic criteria for SAD ever seek help. This results in a median delay of over two decades until correct diagnosis and initial treatment, the longest delay amongst all psychiatric disorders investigated in the US National Comorbidity Survey Replication (Wang et al., 2005). In addition to the characteristic chronic course, SAD patients frequently (20-30%) present with co-morbid major depressive disorder (MDD) Stein et al., 1990; Merikangas and Angst, 1995; Lewinsohn et al., 1997, with SAD being the most prevalent co-morbid anxiety disorder in patients suffering from depressive disorders (Pini et al., 1997; Rush et al., 2005). Co-morbidity is associated with a more severe and chronic disease course and worse clinical outcome (Stein et al., 2001; Beesdo et al., 2007). The vast majority of SAD patients present initially with social anxiety symptoms (Kessler et al., 1999) and develop a co-morbid depression on average within 5 years (Beesdo et al., 2007).

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82 Consistent with these findings, SAD has been shown to be an 83 important predictor/risk factor of a subsequent depressive disorder 84 independent of the age of onset (Stein et al., 2001; Beesdo et al., 85 2007). Furthermore, apart from a SAD lifetime history, distinct psychological constructs within the SAD symptom spectrum (e.g. 86 87 behavioral inhibition) have also been shown to be predictive of 88 the future onset of depression (Beesdo et al., 2007). Other charac-89 teristics of anxiety disorders that have been linked to an increased 90 risk of developing a depressive disorder include the level of anxiety-associated impairment (Bittner et al., 2004) and the presence 91 92 of multiple anxiety disorders (Woodward and Fergusson, 2001) 93 or panic attacks (Goodwin, 2002). However, little is known about 94 the molecular mechanisms involved in the onset of either anxiety or depressive disorders. Changes in cortisol awaking response 95 96 (Adam et al., 2010) and serum interleukin 6 (Khandaker et al., 97 2014) have been reported to be predictors for the future onset of 98 depression in adolescence. So far no biomarkers have been associ-99 ated with the development of depressive episodes in anxiety disorder patients. Awareness of factors that predict increased 100 susceptibility for the onset of a depressive disorder within the 101 102 SAD patient population could lead to an improved clinical outcome 103 due to early intervention (Kessler et al., 1999).

104 In the present study, we investigated molecular changes in 105 serum collected from patients diagnosed with SAD without current 106 co-morbid depressive disorders with the objective to identify a 107 molecular biomarker panel aiding in the prediction of the onset 108 of a depressive disorder within a 2-year follow-up period. We ana-109 lyzed the discriminative power of serum protein changes at the time of baseline clinical assessment of 165 analytes using multi-110 111 plexed immunoassays. Biomarkers were initially identified in 72 112 patients diagnosed solely with SAD and the analysis was then expanded to include SAD patients with other co-morbid anxiety 113 disorders (totalling 143 SAD patients) in order to account for mul-114 115 tiple anxiety spectrum co-morbidities in SAD individuals. A selec-116 tion of these candidate biomarkers was combined into an 117 optimized panel of four serum analytes. Finally, we evaluated the 118 predictive performance of the identified biomarker panels alone 119 and in combination with psychiatric and somatic patient variables 120 selected from structured patient assessments in order to determine 121 their potential for clinical application.

122 2. Subjects and methods

123 2.1. Study sample

Data collected within NESDA, a longitudinal naturalistic 124 125 cohort study were used (Penninx et al., 2008). The full baseline 126 dataset comprises 2981 persons aged 18-65 years, including 127 those with lifetime or current anxiety or depressive disorder 128 (n = 2329; 78%) and healthy controls (n = 652; 22%). Individuals were recruited from the community (n = 564; 19%), primary care 129 (n = 1610; 54%) and specialized mental health care (n = 807;130 27%) between September 2004 and February 2007 at clinical 131 132 sites in Amsterdam, Groningen and Leiden. Recruitment from community samples included sampling from two previously 133 134 studied cohorts with diagnoses based on the Composite International Diagnostic Interview (CIDI) World-Health-135 Organization, 1997 outcomes (for more information on CIDI 136 137 see Section 2.2). Of 662 individuals formerly recruited within 138 the Netherlands Mental Health Survey and Incidence Study (NEMESIS) Bijl et al., 1998, 359 (54%) refused to participate 139 and 303 (46%) were included. The second recruitment involved 140 141 individuals previously enrolled in the Adolescent at Risk for 142 Anxiety and Depression (ARIADNE) Landman-Peeters et al., 143 2005 study. 394 former participants were contacted, 11 (3%)

could not be traced, 122 (31%) refused participation and 261 144 (66%) were included into NESDA. For the initial screening in 145 the primary care setting 23750 questionnaires (Kessler-10 146 Kessler et al., 2003 with additional questions for anxiety spec-147 trum disorders and psychotropic medication) were sent out to 148 general practitioners (GPs) in the vicinity of the clinical sites 149 and 10706 were returned (45%). During the subsequent screen-150 ing stages 4316 (40%) people refused participation, 4532 (43%) 151 were not contacted (due to random selections in the screened 152 individuals) and 248 (2%) met NESDA exclusion criteria (see 153 Section 2.2), resulting in 1610 (15%) included participants. Of 154 1413 people contacted for inclusion based on structured psychi-155 atric interviews in outpatient clinics around the clinical sites, 156 606 (45%) refused and 807 (57%) agreed to participate in 157 NESDA. See Penninx et al. (2008) for more details on the 158 NESDA sample and recruitment process. Ethical approval was 159 granted by local ethic boards of all participating centres and 160 all participants gave written informed consent. Exclusion criteria 161 included a clinical diagnosis of any psychotic disorder, bipolar 162 disorder, severe addictive disorder, obsessive compulsive disor-163 der or non-fluency in Dutch. Baseline data were collected in a 164 4 h interview, gathering information on clinical psychopathol-165 ogy, psychiatric characteristics, use of medication and sociode-166 mographics as well as physical and psychosocial testing. 167 Additionally the baseline evaluation included an overnight fast-168 ing blood draw. Processed serum samples were stored immedi-169 ately at -80 °C until further use (Penninx et al., 2008). A 2-year 170 follow-up assessment was conducted with a response rate of 171 87 1% 172

In the presented study, we used blood analyte abundance infor-173 mation available for a subset of NESDA individuals (n = 1840, 62%174 of the total baseline assessment), who participated in the 2-year 175 follow-up and had sufficient serum available at baseline (approxi-176 mately 1 mL) for multiplexed immunoassay analysis. For our 177 study, we included only individuals with a baseline diagnosis of 178 SAD during the 6 months prior to sampling. We excluded SAD 179 patients with a current depressive disorder at baseline (6 months 180 recency) to be able to investigate subsequent depressive disorder 181 onset or recurrence. This resulted in 143 SAD patients, of whom 182 72 patients had pure SAD and 71 patients had a diagnosis of SAD 183 and another co-morbid anxiety disorder. 184

2.2. Diagnoses, clinical characteristics and self-report inventories

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Diagnoses of anxiety disorders [SAD, panic disorder with agora-186 phobia (PDA), panic disorder without agoraphobia (PD), agorapho-187 bia (AP) and generalized anxiety disorder (GAD)] and depressive 188 disorders (major depressive disorder (MDD) and dysthymia) were 189 established using the Composite International Diagnostic 190 Interview (CIDI) lifetime version 2.1 (World-Health-Organization, 191 1997) by specially trained clinical research staff. This instrument 192 has been used worldwide for WHO field research and has been 193 shown to possess high validity for the detection of anxiety and 194 depressive disorders (Wittchen et al., 1989; Wittchen, 1994) as 195 well as high inter-rater (Wittchen et al., 1991) and test-retest reli-196 ability (Wacker et al., 2006). The CIDI assessment was used again 197 to evaluate the course of disorders after 2 years, determining the 198 presence of a DSM-IV classified anxiety or depressive disorder dur-199 ing the period between baseline and follow-up. Severity of anxiety-200 related symptoms at baseline was assessed with the Fear 201 Questionnaire (FQ) Marks and Mathews, 1979 and the Beck 202 Anxiety Inventory (BAI) Beck et al., 1988. Severity of depression-re-203 lated symptoms was assessed with the Inventory of Depressive 204 Symptomatology (IDS) Rush et al., 1996. Psychotherapy informa-205 tion was based on self-reported participation in any form of psy-206 chiatrist- or psychotherapist-guided psychotherapy during the 207

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