

1   **Predicting depression onset in young people based on clinical, cognitive,**  
2   **environmental and neurobiological data**

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73     **Short/running title:** Predicting youth depression onset using multimodal data

74

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87

88 **ABSTRACT**

89

90 **Background:** Adolescent onset of depression is associated with long-lasting negative  
91 consequences. Identifying adolescents at risk for developing depression would enable the  
92 monitoring of risk-factors and the development of early intervention strategies. Using  
93 machine learning to combine several risk factors from multiple modalities might allow  
94 prediction of depression onset at the individual level.

95 **Methods:** A subsample of a multi-site longitudinal study in adolescents, the IMAGEN study,  
96 was used to predict future (subthreshold) major depressive disorder (MDD) onset in healthy  
97 adolescents. Based on 2-year and 5-year follow-up data, participants were grouped into: 1)  
98 developing an MDD diagnosis or subthreshold MDD and 2) healthy controls. Baseline  
99 measurements of 145 variables from different modalities (clinical, cognitive, environmental  
100 and structural magnetic resonance imaging [MRI]) at age 14 were used as input to penalized  
101 logistic regression (with different levels of penalization) to predict depression onset in a  
102 training dataset (N=407). The features contributing highest to the prediction were validated  
103 in an independent hold-out sample (3 independent IMAGEN sites; N=137).

104 **Results:** The area under the receiver operating characteristics curve (AUROC) for predicting  
105 depression onset ranged between 0.70-0.72 in the training dataset. Baseline severity of  
106 depressive symptoms, female sex, neuroticism, stressful life events and surface area of the  
107 supramarginal gyrus contributed most to the predictive model and predicted onset of  
108 depression with an AUROC between 0.68-0.72 in the independent validation sample.

109 **Conclusions:** This study showed that depression onset in adolescents can be predicted  
110 based on a combination multimodal data of clinical, life events, personality traits, brain  
111 structure variables.

112

113 **INTRODUCTION**

114 Major depressive disorder (MDD) usually has its onset in adolescence and young adulthood  
115 (1), which can have deleterious consequences for a young person's educational and  
116 occupational functioning, and personal and social life (2). Moreover, adolescent onset  
117 depression can have adverse economic consequences for society, since depression onset in  
118 adolescence is associated with poorer social and occupational functioning and recurrent or  
119 persistent mental illness in adulthood (4, 6). Predicting onset of depression at an early stage  
120 is of high clinical relevance, as it might guide the deployment of early interventions and  
121 preventions, thereby reducing the negative long-term consequences associated with  
122 adolescent onset depression.

123

124 Various studies have examined clinical, cognitive and environmental predictors of  
125 depression onset (7, 8). However, most of these studies examined cross-sectional  
126 associations and, hence, did not provide information on directionality (10, 11). Longitudinal  
127 studies are required to study the predictive value of these factors for the onset of depression,  
128 but only few studies exist that have investigated the longitudinal association between clinical,  
129 cognitive and environmental risk factors and subsequent onset of depression in young  
130 people. These studies have shown that risk factors such as anxiety symptoms, diagnosis of  
131 another psychiatric disorder, stressful life events and neuroticism precede the onset of  
132 depression (12–17). There are few studies that have examined the predictive characteristics  
133 of neuroimaging markers, and of those, most were conducted with small sample sizes (18–  
134 22). Our recent review showed that findings have been inconsistent, although there is some  
135 consistent preliminary evidence for blunted (ventral striatum) response to reward processing  
136 as a predictor for later depression (23).

137

138 Most of the longitudinal studies investigating clinical, environmental and neurobiological risk  
139 factors for the onset of depression in adolescence have examined these risk factors in  
140 isolation. It remains to be investigated whether a combination of risk factors may yield better

141 predictive performance, and which risk factors are most predictive. In addition, most of the  
142 studies have used a traditional group comparison approach. However, a statistically  
143 significant variable at group level will not necessarily be useful for individual prediction, due  
144 to low effect size or because of its redundancy with respect to other variables. Conversely,  
145 even seemingly insignificant variables may become important when combined with other  
146 variables. Some studies have however used a multimodal approach to predict depression,  
147 and have identified important predictors such as sex, neuroticism, rumination, negative  
148 affect, low self-esteem, childhood abuse and familial history of mood disorders among others  
149 (3, 5, 7, 9). Machine learning-based predictive models are also well suited for combining  
150 large amount of data and different data modalities into a single model. In addition, contrary  
151 to traditional multivariate prediction methods, they are optimized for evaluating the model's  
152 predictive value for previously unseen individuals ("new" individuals). Thus, they allow  
153 evaluation of the predictive model at the level of the individual.

154

155 A recent machine learning study in 15-year old adolescents using psychosocial variables as  
156 predictors showed that school failure, social isolation, involvement in physical fights, drug  
157 use, running away from home and maltreatment were predictive of MDD onset within 3 to 4  
158 years after baseline, with a receiver operating characteristics curve (AUROC) between 0.76  
159 and 0.79 (24). Importantly, the predictive model was externally validated in two separate  
160 datasets. With regard to neurobiological risk factors, Foland-Ross and colleagues showed  
161 that cortical thickness can predict onset of depression within 5 years after a baseline scan  
162 with 70% accuracy when 55% of the girls developed depression (25). Thickness of the right  
163 precentral and medial orbitofrontal cortex, left anterior cingulate cortex and insula were the  
164 most predictive features in their predictive model.

165

166 These machine learning studies are an important first step towards the development of a  
167 predictive model that enables identification of adolescents at risk for depression. A critical  
168 next step is to elucidate whether we can predict depression onset in adolescents using a

169 combination of risk factors found in these studies described above (neurobiological, clinical,  
170 cognitive and environmental). Therefore, in the current study we examined the predictive  
171 value of a multimodal data, using of clinical, cognitive, environmental and neurobiological  
172 variables, for the onset of MDD, including subthreshold MDD. We included subthreshold  
173 MDD as the DSM diagnostic criteria for adolescent MDD have low diagnostic validity and  
174 specificity, with unclear diagnostic boundaries (26, 27). In addition, earlier studies have  
175 shown that subthreshold MDD is associated with a higher risk for developing future MDD  
176 and other adverse effects that are associated with MDD (28), highlighting the clinical  
177 importance of considering subthreshold MDD when predicting onset of depression in  
178 adolescence. We employed a machine learning method (penalized logistic regression) as  
179 this machine learning algorithm is appropriate to identify, in combination with a feature  
180 selection approach, the optimal set of measures that prospectively predict onset of  
181 depression over 5 years in a subsample of 407 subjects from the IMAGEN study who were  
182 aged 14 at baseline(29). The predictive model was validated in an independent hold-out  
183 sample from the IMAGEN study (N=137), and specificity for depression onset was tested in  
184 a sample with risky alcohol use (N=268). To our knowledge, this is the first machine learning  
185 study in adolescents that combines a number of different modalities to predict depression  
186 onset.

187

## 188 **METHODS**

### 189 ***Participants***

190 The IMAGEN cohort study is a multisite study, in which the baseline (BL) sample consisted  
191 of 2223 adolescents (around 14 years old) who were followed-up at age 16 (follow-up 1;  
192 FU1), 19 (FU2) and 22 (FU3, these data are still being collected) (29). The participants were  
193 recruited from schools, and their diversity in terms of academic performance, socioeconomic  
194 status and behavioral and emotional functioning was maximized. Exclusion criteria included:  
195 receiving treatment for schizophrenia or bipolar disorder, IQ<70, autism diagnosis, nutritional

196 or metabolic diseases, neurological conditions (e.g. brain tumor, epilepsy) and other medical  
197 diagnoses. The data were collected from 8 sites in Europe (France, Germany, Ireland and  
198 United Kingdom). Ethics was approved by local ethics committees. Participants' parents  
199 signed informed consent and participants gave written assent. Participants older than 18  
200 gave informed consent at FU2. Detailed information about the study protocol can be found in  
201 prior literature (29).

202

203 At each timepoint, participants filled out a psychiatric symptom self-assessment using the  
204 Development and Well-Being Assessment (DAWBA) (30). We used the self-report version of  
205 the DAWBA instead of the clinical version in order to be consistent with previous reports.  
206 Three groups were created based on the DAWBA self-assessment: (1) healthy controls who  
207 did not meet criteria for any mental disorder or subthreshold MDD at any of the assessments  
208 (N=430), (2) those who developed subthreshold MDD at follow-up (FU1 and/or FU2; N=177)  
209 or full-threshold MDD at follow-up (N=71). We excluded participants who met criteria for a  
210 psychiatric diagnosis, or subthreshold MDD, at baseline. Full-threshold MDD and  
211 subthreshold MDD were defined based on earlier research in the IMAGEN sample  
212 (Supplementary Methods and Supplementary Figure 1) (31, 32). We will use the term  
213 'depression' when referring to the combined group of subthreshold MDD and full-threshold  
214 MDD. We kept a subset of healthy controls (N=134) for a sensitivity analysis with regard to  
215 predicting onset of risky alcohol use (described below). Therefore, N=296 healthy controls  
216 were included for the main analysis (Supplementary Figure 2).

217

218 To investigate whether our model's performance was specific to the prediction of onset of  
219 depression or was broadly predictive of psychopathology, additional groups (non-  
220 overlapping) were defined based on onset of risky alcohol use at FU. A risky alcohol group  
221 (N=134) was defined by having a total score of 8 or above on the AUDIT at FU1 and/or FU2,  
222 while not meeting criteria for any other psychiatric disorder (including MDD and subthreshold  
223 MDD) at BL and FU. The healthy controls for this analysis were a randomly selected

224 subsample (to match the number of participants in the risky alcohol use group, N=134) of  
225 those participants that did not meet criteria for psychiatric disorders and had a score lower  
226 than 8 on the AUDIT at BL and FU.

227

228 **Predictor variables**

229 All measures were collected at multiple timepoints, however only baseline variables were  
230 included as predictors in this study. Demographic (n=2), clinical (n=7), cognitive (n=24),  
231 personality (n=9), environmental (n=22), substance use (n=4), developmental (n=1) and  
232 structural MRI (i.e. surface-based morphometry) (n=76) variables were used as predictors.

233 In total, 145 predictors were included from these different modalities, described in the  
234 Supplementary Methods and Supplemental Table S2.

235

236 **Statistical analysis**

237 *Splitting the sample into training and validation sets*

238 The dataset for the main analysis was divided into a training dataset (N=407) and  
239 independent validation dataset (N=137) based on recruitment site. Data from three randomly  
240 selected recruitment sites (Dublin, Mannheim and Paris) were kept separately as the  
241 independent validation set (between-site split). The other five sites formed the training  
242 dataset. A between-site split instead of within-site split was chosen to examine if the model  
243 would generalize to completely new sites, which is especially relevant for neuroimaging, as  
244 machine learning models can be influenced by scanner effects. The age, sex and diagnosis  
245 distribution did not differ between training and validation set. The group labels we aimed to  
246 predict were 1) healthy controls versus 2) those who developed depression at follow-up.

247

248 *Prediction of depression onset at follow-up in training dataset*

249 Penalized (to prevent overfitting) logistic regression was performed on the training dataset  
250 including all predictors to predict depression onset at follow-up (Figure 1) (33). We tested

251 model performance across four different values for  $\alpha$  (1 to 0.25, with 0.25 decreases) in the  
252 penalized logistic regression. When  $\alpha$  is 1, it means that the Lasso penalty was applied, and  
253 when  $\alpha$  decreased a combination of Lasso and Ridge penalties were applied. Lasso  
254 facilitates feature selection as it shrinks coefficients of features to zero, thereby removing  
255 these features from the model. Multiple values of  $\alpha$  were used to examine which features  
256 were selected consistently. The hyperparameter  $\lambda$  value, the weight of the penalty, was  
257 determined by selecting the optimal  $\lambda$  associated with the minimum Brier score in an inner  
258 cross-validation loop. Using the R package ‘glmnet’, a sparse model that uses feature  
259 selection was created (34). We applied a 10-fold cross-validation (CV), which was repeated  
260 10 times. For the CV, the training data was divided into 10 sets, and within each cross-  
261 validation fold, 9 sets formed the training set while the 10<sup>th</sup> was held out for testing. We  
262 ensured that the distribution of scanning sites within each group was the same across all 10  
263 CV folds in order to correct for possible site effects. In each CV fold, a random subsample of  
264 healthy controls was selected to match the number of participants in the depression group.  
265 All variables were scaled and centered in the fold and missing values were imputed in the  
266 training sets based on data of the 5 nearest neighbors (35). The parameters of the training  
267 set were used to impute the test set separately to prevent data leakage.

268  
269 To identify features that contributed most to the prediction, the models (at different levels of  
270  $\alpha$ ) were fitted 10 times in random subsamples within the 10 folds (75% of the training dataset  
271 in the fold). Features that were selected in 90% or more of the 100 repeats were identified  
272 (36).

273  
274 *Replication in independent validation dataset*  
275 The features that were identified as most predictive (i.e., selected at least 90% of the times  
276 in the random subsamples) were subsequently used to build a Ridge logistic regression ( $\alpha =$   
277 0) model using the whole training dataset. The Ridge regression approach ensured that all

278 features were used in the model. This model was then applied to the independent validation  
279 dataset (3 recruitment sites as a separate hold-out sample) to evaluate the predictive value  
280 of this subset of features for onset of depression in participants from independent sites.

281

282 *Performance measures*

283 Performance of the models was examined using the AUROC, sensitivity, specificity and  
284 balanced accuracy (average of sensitivity and specificity). The AUROC represents the  
285 probability that a subject from the depression group is ranked lower than a randomly  
286 selected HC subject across all classification thresholds. An AUROC higher than 0.5 is  
287 performing better than chance level. Permutation testing was used to test if the models  
288 performed statistically better than chance level prediction (1000 permutations with randomly  
289 permuted group labels). A non-parametric significance level p-value was estimated as the  
290 proportion the randomly permuted groups that had a higher AUROC than the AUROC for the  
291 original groups.

292

293 **Prediction of future risky alcohol use**

294 To evaluate if the features that were selected in the training set were specific to predicting  
295 onset of depression or whether they predict onset of psychopathology more generally, we  
296 used the Ridge model with the selected feature to predict risky alcohol use at follow-up.

297

298 **Prediction of MDD**

299 To assess if we could predict onset of MDD, we did an exploratory penalized logistic  
300 regression in a CV predicting MDD in the whole IMAGEN dataset (8 sites) (see  
301 Supplemental Material), once excluding subthreshold depression (N = 349) and once with  
302 those with subthreshold depression included in the HC group (N=513).

303

304 [Figure 1]

305

306 **RESULTS**

307 Demographic and clinical characteristics of the healthy controls and participants who  
308 developed depression can be found in Table 1 and Supplemental Table S3.

309

310 [Table 1]

311

312 **Prediction of depression onset**

313 In the training dataset, depression onset (subthreshold and full-threshold MDD combined)  
314 could be predicted with an AUROC ranging between 0.70 and 0.72 across different levels of  
315  $\alpha$  (Table 2). This was significantly different from chance level (all p-values = 0.001).

316

317 [Table 2, Table 3]

318

319 **Feature selection**

320 With an  $\alpha$  of 1, 4 features were selected in the feature selection procedure (Supplemental  
321 Table 3), as well as one recruitment site (Dresden). The features selected were depression  
322 score at baseline, sex and lifetime frequency of events in the family (sum score of the  
323 presence or absence of events such as parents divorced, abused alcohol, fought or argued,  
324 remarried or had money problems) and distress (seeing therapist, thought about suicide,  
325 face broke out in pimples, ran away, gained a lot of weight, got poor grades in school)  
326 categories. At  $\alpha$  of 0.75 and 0.50, the same features were selected but with the addition of  
327 surface area of the supramarginal gyrus. Being bullied at school, neuroticism and verbal  
328 comprehension were additionally selected when  $\alpha$  was 0.25.

329

330 **Generalization to independent validation dataset**

331 The features that were selected in the penalized logistic regression were used to predict  
332 depression onset in the independent validation dataset (3 independent IMAGEN sites), and  
333 an AUROC ranging between 0.68 and 0.72 was achieved (Table 4).

334

335 [Table 4]

336

337 **Generalization to onset of risky alcohol use at follow-up**

338 Demographic and clinical characteristics of the participants that had risky alcohol use at  
339 follow-up can be found in Supplemental Table S4. The model was able to discriminate  
340 between participants with risky alcohol use at follow-up and healthy controls with AUROC of  
341 0.62 when using the features selected at different levels of  $\alpha$  in the model predicting onset of  
342 depression (Table 3).

343

344 **DISCUSSION**

345 In a large longitudinal sample of young people, we were able to prospectively predict  
346 depression onset with an AUROC ranging between 0.70 and 0.72 using penalized logistic  
347 regression applied to a large set of clinical, cognitive, developmental, personality and  
348 neurobiological characteristics. Importantly, our prediction model was validated in an  
349 independent validation sample consisting of participants of the IMAGEN study assessed at  
350 independent sites (AUROC range 0.68-0.72), confirming the validity of the predictive model  
351 and its generalizability to independent recruitment sites.

352

353 Monitoring risk factors identified in this study could lead to early identification of those at risk  
354 for developing depression, which could help the development of risk-factor specific  
355 strategies for prevention of onset of depression. However, the question arises if an AUROC  
356 of 0.72 is high enough for a predictive model to be clinically relevant. Of note, the AUROC  
357 range is concordant with validated prognostic studies in psychosis (0.73-0.79), bipolar  
358 disorder (0.76) and cardiovascular disease (0.76-0.79) (37-39). The clinical utility of a

359 machine learning model should be assessed by considering the cost-effectiveness of  
360 monitoring risk factors for depression identified by the prediction model. Due to the high  
361 levels of disability that depression can cause, with consequences for not only the individual  
362 but for the broader community, monitoring low-cost risk factors such as clinical  
363 characteristics or life events that can predict depression onset in adolescents with an  
364 AUROC of 0.70 might be sufficient.

365

366 The relative contribution of the predictors should be interpreted with caution as the model  
367 performance is based on multivariate data, and features with small weights still contribute to  
368 the overall performance of the model. However, using only the subset of features that made  
369 substantial contributions to the prediction in the training set to predict depression onset in an  
370 independent validation dataset yielded similar AUROCs as in the training set (0.68-0.72).  
371 Higher depressive symptoms at baseline, being bullied at school, neuroticism, female sex  
372 and more negative life events were found to be among the largest contributors to depression  
373 onset, which is in line with previous research that examined these pre-existing risk factors in  
374 isolation, using multivariate non-machine learning methods or a machine-learning method (7,  
375 14, 17, 24, 28). We found that a higher level of depressive symptoms was an important  
376 predictor for subsequent onset of depression, even though participants with subthreshold  
377 depression at baseline were excluded and thus the mean level of depressive symptoms at  
378 baseline was low (mean: 0.75, on a scale from 0-14). This may be due to shared method  
379 variance. The selection of negative life events seems to suggest that early life stress is an  
380 important predictor of depression onset, and that experiencing stressful life events could be  
381 a valid prospective risk factor to monitor. Additionally, the use of machine learning methods  
382 including internal and external validation in the current study strengthens the hypothesis that  
383 the predictive characteristics could be extrapolated to new individuals (40). However, the  
384 performance of the predictive model will likely have to be improved for it to be clinically  
385 useful. Future studies could focus on sex-specific predictors of depression, which might help  
386 improve the performance.

387

388 With regard to brain measures, we found that lower surface area of the supramarginal gyrus  
389 contributed to the model's predictive performance. Prior research has shown that cortical  
390 surface area alterations may play a particular role when depressive symptoms are  
391 experienced early in adolescence (41, 42). Given that cortical surface area, compared with  
392 cortical thickness, has a higher genetic heritability (43), is determined earlier in development,  
393 and is less strongly affected by later environmental influences (44), cortical surface area  
394 reductions may represent a pre-existing risk factor for depression, shaped by genetic factors  
395 and/or early life adversity (45). Of note, surface area of the supramarginal gyrus, involved in  
396 complex higher order cognitive processes, was not identified to be associated with MDD in  
397 adolescents in a large consortium study (N=505 adolescents) (41, 46, 47). Since the  
398 supramarginal gyrus was not selected at the highest  $\alpha$  thus not affecting the AUROC, and  
399 has the supramarginal gyrus been identified as an important brain region in adolescent  
400 depression in previous literature, the predictive role of the surface area of the supramarginal  
401 gyrus is most likely marginal. This is in contrast to a previous study by Foland-Ross and  
402 colleagues who found a similar AUROC including only cortical thickness measures to predict  
403 depression onset in a relatively small sample (N=34) of young adolescent girls (25). An  
404 important difference between the Foland-Ross study and the current study is that we also  
405 included participants with subthreshold depression, and included multimodal predictors with  
406 other modalities that might be more informative than cortical thickness. Given that no other  
407 surface area regions, cortical thickness or subcortical volumes measures were identified in  
408 our feature selection approach and as it is costly to acquire structural neuroimaging  
409 measures, structural imaging might not be a useful predictor for depression onset in young  
410 people. However, this does not implicate that structural brain changes in young people with  
411 depression cannot provide information about the underlying mechanisms of depression.

412

413 The model was not specific to predicting depression onset at follow-up, but could also  
414 successfully predict risky alcohol use in an independent sample, with a slightly lower

415 AUROC (0.62). This may not be surprising, given the high comorbidity between alcohol use  
416 disorder and MDD, with an increase in comorbidity in young adulthood (46). In addition, a  
417 risky lifestyle, including risky alcohol use, in adolescence is predictive of depressive  
418 symptoms (47). Lastly, risky alcohol use occurred in the depression group, which might have  
419 contributed to the lack of specificity of the predictors. Beyond this, comorbidity of mental  
420 disorders is common; most people who experience mental illness will be diagnosed with  
421 more than one psychiatric disorder during their lifetime and an early age of onset of the first  
422 psychiatric disorder has been associated with having more comorbid psychiatric disorders  
423 during the lifetime (48). We anticipate that our model could be similarly predictive for the  
424 onset of mental disorders other than depression or alcohol abuse, in line with previous  
425 longitudinal studies showing that other psychiatric disorders are associated with similar risk  
426 factors as the risk factors identified in the current study such as bullying, neuroticism,  
427 depressive symptoms and stressful life events (28, 49–51). However, since the prevalence  
428 of other disorders such as bipolar disorder and psychosis were limited in the IMAGEN  
429 sample, the hypothesis about the non-specificity for depression of the model requires further  
430 investigation in other samples.

431

432 When the analysis was restricted to patients with MDD and those with subthreshold  
433 depression were excluded, the AUROC was higher than in the main analysis. Unfortunately,  
434 the sample size of the MDD group was too small to allow validation in an independent  
435 dataset. This increase in AUROC when excluding subthreshold depression could be due to  
436 the fact that adolescents who will develop MDD are more differentiated from healthy  
437 adolescents than adolescents developing subthreshold depressive symptoms. When those  
438 who developed subthreshold depression were treated as healthy controls, the AUROC  
439 decreased. These findings further support the postulation that depression based on a cut-off  
440 for a diagnosis of MDD is arbitrary, as young people with a full-threshold MDD diagnosis  
441 cannot reliably be distinguished from those with subthreshold depression as indicated by our  
442 findings.

443

444 The current study has major strengths, including its large sample size, longitudinal design,  
445 and integration of predictors across multiple modalities. However, an important limitation is  
446 that the diagnostic information was based on the self-report DAWBA, a measure that only  
447 captures a period of four weeks prior to each follow-up assessment. Since there was no  
448 information available on possible depressive episodes in the periods between the follow-ups,  
449 we may have missed depression in the healthy control group, which might have impacted  
450 the classification performance of our model (though likely in the direction of weakening it). It  
451 could also have led to underdiagnosing depression at baseline, potentially leading to a less  
452 healthy group at baseline. Additionally, the DAWBA is clinically reliable (30), although due to  
453 the use of a self-report measure, symptoms might have been underreported (52).

454

455 There are still challenges with translating these types of models into clinical practice,  
456 including that the rate of depression is high in the selected sample. Participants with a  
457 psychiatric diagnosis were removed from the healthy control group, which limits the clinical  
458 utility of the model as people in the general population might show non-depression  
459 psychiatric diagnoses. Therefore, future studies should test if a predictive model works in the  
460 general population that includes people who have already experienced episodes of mental ill  
461 health. Additionally, the depression group might include young people with comorbid  
462 diagnoses such as anxiety disorders with similar risk factors, which could increase the  
463 predictive power. However, in a sensitivity analysis, excluding those who developed  
464 comorbid anxiety disorder at follow-up, showed that the predictive performance measures  
465 were similar.

466

467 In conclusion, the current study showed that depression onset in adolescents can be  
468 predicted based on multimodal data, including clinical, cognitive, life events, personality traits  
469 and neurobiological variables. The variables contributing most to the predictive model were  
470 found to be depressive symptoms at baseline, neuroticism, cognition, supramarginal gyrus

471 surface area and stressful life events. Since the model was also predictive of onset of risky  
472 alcohol use, these risk factors may likely be predictive more generally of onset  
473 psychopathology during adolescence.

474

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509

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522

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- 675
- 676

677 **Table 1. Demographics and clinical characteristics of the groups in the training and**  
 678 **validation datasets.**

	Training depression (n=180)	Training control (n=227)	Validation depression (n=68)	Validation control (n=69)
<b>Age</b>				
Mean (SD)	14.5 (0.54)	14.4 (0.44)	14.4 (0.59)	14.4 (0.61)
<b>Sex</b>				
Female	121 (67%)	104 (46%)	46 (68%)	29 (42%)
Male	59 (33%)	123 (54%)	22 (32%)	40 (58%)
<b>Site</b>				
Berlin	34 (19%)	17 (8%)	NA	NA
Dresden	17 (9%)	62 (27%)	NA	NA
Hamburg	35 (19%)	45 (20%)	NA	NA
London	47 (26%)	53 (23%)	NA	NA
Nottingham	47 (26%)	50 (22%)	NA	NA
Dublin	NA	NA	21 (31%)	11 (16%)
Mannheim	NA	NA	20 (29%)	27 (39%)
Paris	NA	NA	27 (40%)	31 (45%)
<b>Depression score at BL</b>				
<b>(DAWBA)*</b>				
Mean (SD)	1.07 (1.23)	0.59 (0.80)	0.82 (0.88)	0.36 (0.57)

679 DAWBA: development and well-being assessment, MDD: major depressive disorder, N:  
680 sample size, SD: standard deviation. \* score based on number of depressive symptoms  
681 present according to youth self-report DAWBA, ranges between 0-14.

682

683 **Table 2.** Performance measures in penalized logistic regression for four different  $\alpha$  (Ridge  
684 towards Lasso penalty) to predict depression onset in the training set.

685

$\alpha$	AUROC	SD AUROC	Sensitivity	Specificity	Accuracy
0.25	0.70	0.10	0.66	0.66	0.66
0.5	0.70	0.08	0.66	0.65	0.65
0.75	0.72	0.08	0.67	0.66	0.66
1	0.71	0.07	0.65	0.66	0.66

686 SD: standard deviation across folds

687

688 **Table 3. Selected features in penalized logistic regression for prediction of depression  
689 onset at different  $\alpha$  levels in training dataset.**

Predictor category	Parameter threshold			
	$\alpha = 0.25$	$\alpha = 0.5$	$\alpha = 0.75$	$\alpha = 1$
Clinical	DAWBA depression	DAWBA depression	DAWBA depression	DAWBA depression
Life events	LEQ family lifetime	LEQ family lifetime	LEQ family lifetime	LEQ family lifetime
	LEQ distress lifetime	LEQ distress lifetime	LEQ distress lifetime	LEQ distress lifetime
	Bullied at school			
Personality	Neuroticism			

<b>Cognitive</b>	WISC-IV Similarities			
<b>Biological</b>	Sex	Sex	Sex	Sex
	Supramarginal gyrus surface area	Supramarginal gyrus surface area	Supramarginal gyrus surface area	

690 DAWBA: development and well-being assessment, LEQ: life events questionnaire, WISC-IV:

691 Wechsler intelligence scale for children – fourth edition.

692

693 **Table 4.** Performance measures of Ridge logistic regression with the features that were  
 694 selected in the training dataset across different levels of  $\alpha$  to predict depression at follow-up  
 695 in the independent validation dataset.

696

Number of features (selected at which $\alpha$ in training set)	AUROC	Sensitivity	Specificity	Accuracy
---	-------	-------------	-------------	----------

#### Predicting depression in independent validation dataset

8 ( $\alpha=0.25$ )	0.72	0.51	0.83	0.67
5 ( $\alpha=0.50$ and $0.75$ )	0.68	0.49	0.77	0.63
4 ( $\alpha=1$ )	0.71	0.50	0.81	0.66

#### Predicting risky alcohol use in independent dataset

8 ( $\alpha=0.25$ )	0.62	0.41	0.74	0.57
5 ( $\alpha=0.50$ and $0.75$ )	0.62	0.43	0.79	0.61
4 ( $\alpha=1$ )	0.62	0.42	0.78	0.60

697 SD: standard deviation

698

699 **Figure 1 Statistical procedure for penalized logistic regression.** 1. Baseline predictors  
700 from different domains were used to predict subthreshold MDD or MDD onset at follow-up. 2.  
701 Penalized logistic regression with 10-fold cross-validation was applied to the training dataset  
702 (5 sites) and repeated 10 times with 4 different levels of  $\alpha$ . Permutation testing was used to  
703 test the statistical significance of the model. 3. Features that were selected in 90% or more  
704 in 100 random subsamples of the training data were selected to be tested in the  
705 independent validation set. 4. The selected features from step 3 were used as input to Ridge  
706 logistic regression in the whole training set that was then used to predict depression onset in  
707 the validation set (3 independent sites), to test the generalizability of the model in the three  
708 sites that were left out from the training set. 5. The same Ridge model was used to evaluate  
709 its predictive value for onset of risky alcohol use in unseen individuals.

710