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# Depressive symptoms and single nucleotide polymorphisms predict clinical recurrence of inflammatory bowel disease — Source link 🖸

Sebastian Bruno Ulrich Jordi, Brian M. Lang, Bianca Auschra, Roland von Känel ...+8 more authors

Institutions: University of Zurich, University Hospital of Basel, University of Bern, Kantonsspital St. Gallen

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# **1 Depressive symptoms and single nucleotide polymorphisms**

# 2 predict clinical recurrence of inflammatory bowel disease

3 Short title: Depression and IBD course

# 4 Authors

- 5 Sebastian Bruno Ulrich Jordi <sup>1,2</sup>, Brian Matthew Lang <sup>3</sup>, Bianca Auschra <sup>4</sup>, Roland
- 6 von Känel<sup>4</sup>, Luc Biedermann<sup>2</sup>, Thomas Greuter<sup>2</sup>, Philipp Schreiner<sup>2</sup>, Gerhard
- 7 Rogler <sup>2</sup>, Niklas Krupka <sup>1</sup>, Michael Christian Sulz <sup>5</sup>, Benjamin Misselwitz <sup>1,2\*</sup>, Stefan
- 8 Begré <sup>6,7</sup>\* on behalf of the Swiss IBD cohort study group<sup>#</sup>

- <sup>1</sup> Clinic for Visceral Surgery and Medicine, Inselspital Bern and Bern University,
   Bern, Switzerland
- <sup>2</sup> Department of Gastroenterology and Hepatology, University Hospital Zurich and
   Zurich University, Zurich, Switzerland
- <sup>3</sup> Clinic for Transplantation Immunology and Nephrology (Swiss Transplant Cohort
- 15 Study), University Hospital of Basel, Basel, Switzerland
- <sup>4</sup> Department of Consultation-Liaison Psychiatry and Psychosomatic Medicine,
- 17 University Hospital Zurich and University of Zurich, Zurich, Switzerland
- <sup>5</sup> Department of Gastroenterology and Hepatology, Kantonsspital St. Gallen, St.
   Gallen, Switzerland
- <sup>6</sup> Neurology, Department of Biomedical Research, Bern University Hospital,
- 21 University of Bern, Bern, Switzerland
- <sup>22</sup> <sup>7</sup> ISFOM Institute of Stress Diseases and Stressmanagement, Zurich, Switzerland
- 23
- <sup>\*</sup> Both authors contributed equally and share last authorship
- 25 # Members of the SIBDCS study group: Claudia Anderegg; Peter Bauerfeind;
- 26 Christoph Beglinger; Stefan Begré; Dominique Belli; José M. Bengoa; Luc

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27 Biedermann; Beat Bigler; Janek Binek; Mirjam Blattmann; Stephan Boehm; Jan 28 Borovicka; Christian P. Braegger; Nora Brunner; Patrick Bühr; Bernard Burnand; 29 Emanuel Burri; Sophie Buyse; Matthias Cremer; Dominique H. Criblez; Philippe de 30 Saussure; Lukas Degen; Joakim Delarive; Christopher Doerig; Barbara Dora; Gian 31 Dorta; Mara Egger; Tobias Ehmann; Ali El-Wafa; Matthias Engelmann; Jessica Ezri; 32 Christian Felley; Markus Fliegner; Nicolas Fournier; Montserrat Fraga; Pascal Frei; 33 Remus Frei; Michael Fried; Florian Froehlich; Christian Funk; Raoul Ivano Furlano; 34 Suzanne Gallot-Lavallée; Martin Geyer; Marc Girardin; Delphine Golay; Tanja 35 Grandinetti; Beat Gysi; Horst Haack; Johannes Haarer; Beat Helbling; Peter 36 Hengstler; Denise Herzog; Cyrill Hess; Klaas Heyland; Thomas Hinterleitner; 37 Philippe Hiroz; Claudia Hirschi; Petr Hruz; Rika Iwata; Res Jost; Pascal Juillerat; 38 Vera Kessler Brondolo; Christina Knellwolf; Christoph Knoblauch; Henrik Köhler; 39 Rebekka Koller; Claudia Krieger-Grübel; Gerd Kullak-Ublick; Patrizia Künzler; 40 Markus Landolt; Rupprecht Lange; Frank Serge Lehmann; Andrew Macpherson; 41 Philippe Maerten; Michel H. Maillard; Christine Manser; Michael Manz; Urs Marbet; 42 George Marx; Christoph Matter; Valérie McLin; Rémy Meier; Martina Mendanova; 43 Christa Meyenberger; Pierre Michetti; Benjamin Misselwitz; Darius Moradpour; 44 Bernhard Morell; Patrick Mosler; Christian Mottet; Christoph Müller; Pascal Müller; 45 Beat Müllhaupt; Claudia Münger-Beyeler; Leilla Musso; Andreas Nagy; Michaela 46 Neagu; Cristina Nichita; Jan Niess; Natacha Noël; Andreas Nydegger; Nicole Obialo; 47 Carl Oneta; Cassandra Oropesa; Ueli Peter; Daniel Peternac; Laetitia Marie Petit; 48 Franziska Piccoli-Gfeller; Julia Beatrice Pilz; Valérie Pittet; Nadia Raschle; Ronald 49 Rentsch; Sophie Restellini; Jean-Pierre Richterich; Sylvia Rihs; Marc Alain Ritz; 50 Jocelyn Roduit; Daniela Rogler; Gerhard Rogler; Jean-Benoît Rossel; Markus 51 Sagmeister; Gaby Saner; Bernhard Sauter; Mikael Sawatzki; Michela Schäppi;

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52	Michael Scharl; Martin Schelling; Susanne Schibli; Hugo Schlauri; Sybille Schmid
53	Uebelhart; Jean-François Schnegg; Alain Schoepfer; Frank Seibold; Mariam Seirafi;
54	Gian-Marco Semadeni; David Semela; Arne Senning; Marc Sidler; Christiane
55	Sokollik; Johannes Spalinger; Holger Spangenberger; Philippe Stadler; Michael
56	Steuerwald; Alex Straumann; Bigna Straumann-Funk; Michael Sulz; Joël Thorens;
57	Sarah Tiedemann; Radu Tutuian; Stephan Vavricka; Francesco Viani; Jürg Vögtlin;
58	Roland vVon Känel; Alain Vonlaufen; Dominique Vouillamoz; Rachel Vulliamy; Jürg
59	Wermuth; Helene Werner; Paul Wiesel; Reiner Wiest; Tina Wylie; Jonas Zeitz;
60	Dorothee Zimmermann.
61	
62	Correspondence
63	Stefan Begré, MD
64	Address: ISFOM - Institute of Stress Diseases and Stressmanagement,
65	Weinbergstrasse 139, CH-8006 Zurich, Switzerland
66	E-mail address: stefan.begre@dbmr.unibe.ch
67	
68	
69	Authors' contributions
70	The study was conceived and designed by S.B.U.J., B.M. and S.B.
71	Statistical analyses were performed by S.B.U.J. and B.M.L. The manuscript was
72	written by S.B.U.J., B.M. and S.B. The following authors have contributed to data
73	interpretation and revised the manuscript for important intellectual content: B.M.L.,
74	R.v.K., B.A., L.B., T.G., P.S., G.R., N.K. and M.C.S. All authors approved the final
75	version of the manuscript.

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# 77 Abbreviations

- 78 **aHR**: adjusted hazard ratio
- 79 **AIC**: Akaike information criterion
- 80 **BMI**: body mass index
- 81 **CD**: Crohn's disease
- 82 CDAI: Crohn's Disease Activity Index
- 83 CHRNA5: cholinergic receptor nicotinic alpha 5 subunit
- 84 **CI:** confidence interval
- 85 **EIM:** extraintestinal manifestations
- 86 ENS: enteric nervous system
- 87 HADS: Hospital Anxiety and Depression Scale
- 88 HR: hazard ratio
- 89 **IBD**: inflammatory bowel disease
- 90 IC: indeterminate colitis
- 91 LD: linkage disequilibrium
- 92 **MTWAI**: Modified Truelove and Witts Severity Index
- 93 NAChR: nicotinic acetylcholine receptors
- 94 **OR**: odds ratio
- 95 **PCLO**: piccolo presynaptic cytomatrix protein
- 96 **PSC:** primary sclerosing cholangitis
- 97 **R**<sup>2</sup>: square of the correlation coefficient

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- 98 SIBDC: Swiss IBD cohort
- 99 **SNP**: single nucleotide polymorphism
- 100 **t**<sub>n</sub>: follow-up time *n* of recording
- 101 **TNF**: tumor necrosis factor
- 102 **UC**: ulcerative colitis

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# 104 Abstract

# 105 Background and Aims

Inflammatory bowel disease (IBD) patients are at high risk for depression. We
 examined interrelations between genetic risk factors for depression, depressive
 symptoms and IBD flares.

109

# 110 Methods

111 In 1973 patients (1137 Crohn's disease, 836 ulcerative colitis) of the Swiss IBD 112 cohort study (SIBDC), 62 single nucleotide polymorphisms (SNPs) preselected for 113 associations with depression, stress, pain and smoking were screened for cross-114 sectional associations with depression (hospital anxiety and depression subscale for 115 depression, HADS-D≥11). Logistic regression and Cox proportional hazards models 116 were built to test for effects of depressive symptoms on disease course and genetic 117 risk factors on depression and disease course. As endpoints we used active disease 118 (CDAI≥150 or MTWAI≥10) and two published composite flare definitions: FNCE: 119 physician reported flare, non-response to therapy, new complication or 120 extraintestinal manifestation and AFFSST: active disease, physician reported flare, 121 fistula, stenosis and new systemic therapy.

122

### 123 **Results**

Depressive symptoms were a strong risk factor for disease related endpoints including active disease (adjusted hazard ratio, aHR: 3.25, p<0.001), AFFSST (aHR: 1.62, p<0.001) and FNCE (aHR: 1.35, p=0.019). Rs588765's TC alleles and rs2522833's C allele were associated with depressive symptoms at baseline (odds

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128	ratio, OR: 0.43, q=0.050 and OR: 1.73, q=0.059, respectively). Rs588765-TC
129	remained protective regarding presence of depression (aHR: 0.67, p=0.035) and
130	was associated with fewer active disease states (aHR: 0.72, p=0.045) during follow-
131	up.
120	

132

# 133 Conclusions

- 134 In IBD, genetics, depressive symptoms and inflammatory activity are intimately
- 135 related: Depressive symptoms were a predictor of clinical deterioration and
- 136 rs588765-TC was protective for depression and high IBD activity.

137

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# 141 Introduction

Inflammatory bowel diseases (IBD) are frequent conditions affecting approximately 2.2 million people in Europe and 1.5 million Americans with an increasing incidence worldwide.<sup>1</sup> IBD, comprising the subtypes Crohn's disease (CD), ulcerative colitis (UC) and indeterminate colitis (IC) are immunologically mediated diseases which can severely impair affected individuals' lives. The disease course of IBD is variable with periods of quiescent disease which are interrupted by IBD flares, i.e. episodes of more active inflammation.

The exact cause of IBD remains elusive, but likely involves an interplay of genetic, microbial, immunological and environmental factors.<sup>2</sup> A significant part of IBD risk is heritable and more than 230 single nucleotide polymorphisms (SNP) associated with IBD have been identified.<sup>3</sup> These SNPs partially explain an individual's vulnerability to develop IBD but have not been useful to predict the clinical disease course in IBD patients.<sup>4</sup> Many environmental factors have been implicated as trigger factors for IBD flares including diet, medication and smoking.<sup>5</sup>

Psychiatric conditions such as stress, depression and anxiety have been associated with IBD and the risk to experience depression is higher for IBD patients compared to healthy individuals (incidence rate ratio: 1.6).<sup>6</sup> Furthermore, stress and depression overlap with pain<sup>7</sup>, pain sensitivity<sup>8</sup> and smoking.<sup>9</sup> Stress, depression and anxiety have been suggested as triggers of flares and clinical deterioration in IBD.<sup>10,11</sup>

Depressive disorders are heritable conditions with heritability rates of approximately 40%<sup>12</sup> and many individual SNPs have been associated with the risk for depression.<sup>13</sup> An example for a possible genetic origin of the observed overlap of depression, smoking and pain sensitivity are neuronal nicotinic acetylcholine

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receptors (nAChRs). Recent findings suggest that nAChRs are involved in psychiatric disorders such as depression although mechanistic details remain unclear.<sup>14</sup> Moreover, nAChRs had been associated with nicotine and alcohol dependence,<sup>15,16</sup> also among SIBDC patients.<sup>17</sup> Currently, nAChRs are investigated as pharmacological targets for depression, anxiety, pain and nicotine addiction.<sup>18</sup> For these reasons, SNPs within nAChR genes might influence the occurrence of flares in IBD via their association with depression and/ or their link to smoking.

However, the role of genetic risk factors for depressive symptoms has not yet been clarified for individuals at high risk for depression such as IBD patients and also the interplay between genetic and environmental risk factors of depression remains unclear.

We are using data from the Swiss IBD cohort study (SIBDC)<sup>19</sup> to explore the relationship between depression, individual SNPs and the risk for IBD deterioration. SIBDC was started in 2006 as a prospective cohort study that recruits nationwide in Switzerland.<sup>19,20</sup> The impact of depressive symptoms, anxiety and social support on IBD flares has been studied in SIBDC before;<sup>10,21</sup> we were aiming to validate, generalize and extend these findings, and add a genetic context.

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# 183 Methods

## 184 **Design of the SIBDC**

The SIBDC was started in 2006 as a prospective cohort study that recruits nationwide in Switzerland.<sup>19,20</sup> New patients are continuously recruited and patients are followed yearly. IBD-related clinical data is collected by physicians at enrolment and yearly follow-ups. In addition, clinical, sociodemographic and psychosocial data are acquired using patient questionnaires at enrolment and yearly follow-ups. All SIBDC patients consenting to genetic analysis with appropriate biomaterial available were genotyped and only genotyped patients were included in our study.

SIBDCS is funded by the Swiss National Science Foundation and has been approved by the local ethics committee of each participating center (institutional review board approval No. EK-1316, approved on February 5, 2007 and KEK Zurich, March 9, 2020; BASEC 2018-02068). SIBDC goals and methodology are described elsewhere.<sup>19,20</sup> All patients provided written informed consent prior to inclusion into the SIBDCS. Analysis of patient data for the current study was approved by the scientific board of SIBDCS.

199

### 200 Patient characteristics

We exported clinical and sociodemographic variables acquired at enrolment and follow-ups: Disease activity was measured using the Crohn's disease activity index CDAI or the modified Truelove and Witts activity index (MTWAI) for CD and UC/IC patients, respectively. For intercomparable measures, CDAI or MTWAI score values were Z-score normalized to a disease activity score (disease activity score =  $(s_i - \bar{s}) / \sigma_s$ ) or cutoff values for active disease were used (see below). The following

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207 binary variables were created by distinguishing between patients that had any of the 208 respective included criteria and patients that did not: Complications included 209 colorectal cancer, colon dysplasia, intestinal lymphoma, osteopenia/osteoporosis, 210 anemia (not as an adverse event of medical therapies), deep vein thrombosis, pulmonary embolism, nephrolithiasis, gallstones,<sup>22</sup> malabsorption syndrome, 211 212 massive hemorrhage, perforation or peritonitis. Extraintestinal manifestations (EIM) 213 comprised peripheral arthritis/arthralgia, uveitis/iritis, pyoderma gangrenosum, 214 erythema nodosum, aphthous oral ulcers/stomatitis, ankylosing spondylitis/sacroiliitis 215 and primary sclerosing cholangitis (PSC). Any stenosis localized in the esophagus, 216 duodenum/jejunum, ileum, large bowel, rectum or anus was summarized in the 217 stenosis variable. The fistula variable comprised IBD-related fistula and abscesses 218 and anal fissure, whereas the variable *surgery* included any abdominal or fistula and 219 abscess-related surgery. TNF inhibitors referred to treatments with infliximab, 220 certolizumab pegol or adalimumab (no patient with golimumab treatment was 221 reported at the time of data export). The variable *number of current therapies* was 222 created by counting presently administered therapies which could be any of the 223 following: any steroids, any antibiotics, 5-aminosalicylic acid, sulfasalazine, 224 azathioprine, 6-mercaptopurine, methotrexate, cyclosporine, tacrolimus, infliximab, 225 adalimumab, certolizumab pegol, cholestyramine, *E. coli* Nissle (Mutaflor<sup>®</sup>), ursodeoxycholic acid, bisphosphonates, the probiotic VSL#3<sup>®</sup>, or any other 226 227 medication that was noted by physicians. A patient was considered a smoker when 228 he or she declared to have a current habit of smoking independent of the quantity.

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# 231 Outcome measures

Depressive symptoms were measured with the Hospital Anxiety and Depression Scale's (HADS) subscale for depression (HADS-D).<sup>23</sup> Following well-established criteria, the presence of depression was defined as HADS-D  $\geq$ 11, indicating probable moderate or severe depression.<sup>24-26</sup> This was done for both a constant (at baseline) and a time-varying depression variable. In this study, the term *depression* is used with reference to this definition unless stated otherwise.

Active disease was defined by CDAI ≥150<sup>27</sup> or MTWAI ≥10<sup>28</sup>, respectively. An IBD 238 239 flare was measured as a composite endpoint of clinical variables. IBD is a complex 240 disease and we considered two different flare definitions, following the examples of two prior SIBDC studies.<sup>10,21</sup> The first composite clinical endpoint *active disease*, 241 242 physician reported flare, fistula, stenosis, surgery or new systemic therapy 243 (AFFSST) was reached upon occurrence/persistence of active disease, defined by 244 CDAI  $\geq 150^{27}$  or MTWAI  $\geq 10^{28}$ , a physician reported flare, new fistula, new stenosis, 245 new surgery, intake of systemic steroids and/or start of therapy with a new TNF 246 inhibitor (also changes between TNF inhibitors) at follow-up.<sup>10</sup>

The second composite clinical endpoint *physician reported flare, non-response to* any administered therapy with consequent transition to a more aggressive medication, new complication or new *EIM* (FNCE) was defined as the occurrence of any one of these events at a follow-up examination.<sup>21</sup>

For both composite endpoints we counted the first occurrence of each defining feature after enrolment as event. Both composite endpoints were reached upon the first occurrence of any of these events.

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For data analyses with survival techniques, all endpoints (the individual flare features, the FNCE and AFFSST composite endpoints and depression) were coded with right censoring. Thereby, a time interval  $t_{n-1}-t_n$  was defined where  $t_n$  corresponds to the n<sup>th</sup> follow-up visit when the event was documented first. This way, we describe a time interval during which an event had occurred [ $t_{n-1}$ ,  $t_n$ ] and then assumed an event's occurrence at  $t_{n-1}+(t_n-t_{n-1})/2$  on average.

260

### 261 Selection of candidate SNPs

262 Allelic state of a number of SNPs were determined in SIBDC patients. We selected 263 all 95 SNPs available without a link to inflammation or IBD diagnosis. These non-264 inflammatory SNPs had been selected with three objectives: 48 SNPs were selected 265 for an association with stress and/or depressive symptoms, based on a thorough 266 literature search in PubMed and SNPedia. Moreover, 30 SNPs with a suspected 267 association with pain were included. Furthermore, 16 SNPs associated with smoking in all 3 genome wide association studies available in 2015 were included.<sup>17,29-31</sup> 268 Both, pain<sup>7</sup> and smoking<sup>11,37</sup> have a strong reciprocal associations with depression 269 270 and anxiety.

For calculation of linkage disequilibrium (LD) we used the square of the correlation coefficient ( $r^2$ ) between two SNPs. Groups were formed whenever two or more SNPs shared an  $r^2$  value of 0.5 or higher, representing moderate LD.<sup>32</sup> The SNP with the strongest correlation to group members was chosen as the representing tag-SNP for the respective group. If there were only two SNPs in a group, the SNP with the lowest number of missing data points in the respective group was selected as tag SNP.

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# 279 Screening for depression-associated SNPs

280 For the screening of SNPs associated with depression, we used methods of the SNPassoc R package<sup>33</sup> version 1.9-2, developed and established for whole genome 281 282 association studies. We calculated the association between each of our SNP and 283 depression while simultaneously selecting the genetic model (codominant, dominant, 284 recessive, overdominant, log-additive) for each SNP which best describes this 285 relationship via likelihood ratio test comparison. We filtered the list of SNPs to those 286 with a Bonferroni corrected p-value (q-value) <0.1. Associations were confirmed by 287 multivariable analyses with depressive symptoms at enrolment as endpoint. For 288 multivariable analyses with depressive symptoms as dependent variable in this 289 study, a set of control variables was defined (set 1: sex, diagnosis, age, time since 290 diagnosis, BMI, smoking status, alcohol consumption, standardized CDAI/MTWAI 291 score, abdominal surgery prior to enrolment, systemic steroids, TNF inhibitors, 292 number of current therapies). These variables were tested for association with 293 depressive symptoms at enrolment using univariable logistic regression analysis.

Models included all control variables and were calculated for every SNP individually. To minimize the risk of obtaining significance due to overfitting we also created a simpler secondary model by automated variable elimination. In this process the optimal model regarding the AIC was calculated. For this we used the *glmulti* R package<sup>34</sup> version 1.0.7.1.

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### 300 Time-to-event analyses

To confirm associations with SNPs at enrolment (cross-sectional data), we constructed time-to-event models using the *Survival* R package<sup>35</sup> version 3.1-12. We implemented univariable and multivariable Cox proportional hazards models with the endpoint of interest being presence of depressive symptoms (HADS  $\geq$ 11) at a followup examination, independent of a patients HADS score value at enrolment.

306 Clinical disease course was described by the absence or presence of either active 307 disease or flares according to composite endpoint definitions (see above). To assess 308 the impact of depressive symptoms and depression-associated SNPs on the clinical course of IBD.<sup>10</sup> we coded a time-varying control variable for depression (HADS-D 309 score ≥11) that reflected a patient's depression status over time.<sup>36</sup> To ensure an 310 311 unambiguous time sequence of time-varying depression status and clinical endpoint 312 measurements (e.g. flares), a conservative coding approach was chosen: only if a 313 recording of depressive symptoms preceded a flare, an association was made. 314 Depression status was defined forwardly, while flares were defined backwardly. For 315 example, measures for depression at time point  $t_n$  defined the time-varying 316 depression variable for the interval  $t_n$  until  $t_{n+1}$ , while recordings of clinical variables at 317 time point  $t_n$  defining a clinical endpoint variable (e.g. flare) where assumed to have 318 happened at the time point in the middle between the last and the current follow-up 319 (calculated as  $t_{n-1}+(t_n-t_{n-1})/2$ ). As a result, concurrent depression and flares will not be 320 detected as an association.

For these time-to-event analyses with clinical endpoints we adapted the set of control variables by removing variables that could be proxy-variables for clinical endpoints (set 2: *sex, diagnosis, time since diagnosis, age, BMI, disease related surgery prior to enrolment, smoking status, daily alcohol consumption*). As before,

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- 325 univariable and multivariable Cox proportional hazards models were implemented.
- 326 All calculations were performed using  $R^{37}$  version 3.6.1.

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# 327 **Results**

# 328 Study participants

329 For our analysis, data from 1973 genotyped SIBDC patients (1137 with CD, 836 with 330 UC/IC) were available with regular follow-ups up to 4250 days. Baseline 331 characteristics of our cohort sample are presented in **Table 1**, confirming a mixed 332 study population with mild, moderate and severe disease. Depression (HADS-D  $\geq$ 11) 333 at enrolment tended to be more prevalent in CD compared to UC (9.5% in CD vs. 6.5% in UC/IC, p=0.052), consistent with earlier findings.<sup>38</sup> Smoking,<sup>39</sup> fistulae, 334 335 stenosis, EIMs, complications, previous abdominal surgery and therapy with TNF 336 inhibitors was more common in CD patients and CD patients had significantly longer 337 disease courses and lower BMIs. On the other hand, UC/IC patients had a higher 338 number of different therapies at enrolment and were more often treated with 339 systemic steroids.

Depressive symptoms at enrolment were significantly associated with several control variables including BMI (p=0.01), daily alcohol consumption (p=0.017), disease activity (p<0.001) and number of current therapies (p=0.004, **Table 2**).

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# 344 Screening of SNPs for an association with depressive symptoms

SNPs were tested for linkage disequilibrium (Supplementary figure 1) and SNPs with a pair-wise correlation coefficient  $r^2 \ge 0.5$  aggregated in 16 groups. From each group only one representative SNP was used henceforward, reducing the number of SNPs from 95 to 62 (Supplementary table 1).

All SNPs were screened for an association with depression (HADS-D  $\geq$ 11) at enrolment in three different models: univariable, complete multivariable and reduced

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351 multivariable model after automated parameter elimination (Supplementary table 352 2). Of 62 SNPs tested, two SNPs, rs588765 and rs2522833, were significantly 353 associated with depressive symptoms at enrolment with q<0.1 (Bonferroni 354 corrected). In the univariable model, for rs2522833 the association was strongest in 355 a log-additive inheritance model with the C allele as an associated risk allele (odds 356 ratio, OR: 1.71, CI: 1.25-2.34, q=0.043). For rs588765, an overdominant genetic 357 pattern yielded the strongest significance identifying TC as an associated protective 358 allele combination (OR: 0.45, CI: 0.28-0.74, q=0.066, **Table 3** and **Supplementary** 359 figure 2). For both SNPs, the associations detected in the univariable models stayed 360 significant (q<0.1) in multivariable models correcting for possible confounders 361 (rs2522833: q=0.045; rs588765: q=0.061, **Supplementary figure 3**), also after 362 parameter elimination (rs2522833: q=0.059; rs588765: q=0.050, Supplementary 363 figure 4). This indicates a robust association between both SNPs and depressive 364 symptoms

365

### 366 SNPs effects on depression-free survival

367 To assess the effects of both SNPs on the hazard for continuance or new onset of 368 depressive symptoms (HADS-D ≥11) over time, we implemented Cox proportional 369 hazards models. In this time-to-event analyses, the TC allele combination of 370 rs588765 had significant protective effects regarding depressive symptoms in the 371 univariable model (hazard ratio (HR)): 0.68, confidence interval (CI): 0.46-0.97, 372 p=0.042). The association with rs588765-TC remained robust in the multivariable 373 analysis (adjusted hazard ratio (aHR): 0.67, CI: 0.46-0.97, p=0.035; Table 4 and 374 Figure 1; panel A). Protective effects of rs588765-TC were independent of disease

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375	type and similar for CD and UC patients, although non-significant in each subgroup
376	(CD: aHR: 0.70, p=0.155; UC: aHR: 0.58, p=0.100).
377	For the C-allele of rs2522833, no significant association was detected (aHR: 1.10,

- 378 CI: 0.86-1.41, p=0.455, Table 4 and Supplementary figure 5; panel A).
- 379

# 380 Depressive symptoms as a hazard for IBD disease course

- 381 Patients with depressive symptoms (HADS-D ≥11) had a significantly higher hazard
- of experiencing active disease (CDAI ≥150/MTWAI ≥10) during follow-up (HR: 3.18,
- p<0.001, Table 5 and Supplementary figure 6; panel A); this association remained
- significant after correction for confounders (aHR: 3.25, p<0.001, **Table 5** and **Figure**
- 385 **2; panel A)**. We also observed a significant negative impact of depressive symptoms
- on the endpoints new surgery (aHR: 1.92, p>0.001), systemic steroid therapy (aHR:
- 387 1.55, p=0.033) and primary non-response to therapy (aHR: 1.79, p=0.044, Figure 2;
- panel E-G, univariable results: Supplementary figure 7). In other words, patients
  were more likely to experience the respective endpoints when depressive symptoms
  were present before, even when corrected for possible confounders.
- Overall, for all clinical endpoints, patients with depressive symptoms had a higher hazard for clinical deterioration, even though this effect did not reach significance in all cases (**Figure 2**). Furthermore, depressive symptoms also tended to increase the hazard for individual EIMs and significant effects for new PSC (aHR: 3.88, p=0.032) and new uveitis/iritis (aHR: 2.01, p=0.046) in multivariable models were observed (**Figure 3; panel E** and **G**, univariable results: **Supplementary figure 8**).
- 397 To further investigate the impact of depressive symptoms on IBD disease course,<sup>10</sup> 398 we assessed two previously used composite endpoints indicating clinical

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deterioration: AFFSST<sup>10</sup> (**a**ctive disease, physician reported flare, new fistula, new stenosis, surgery or new systemic therapy, see methods) and FNCE<sup>21</sup> (physician reported flare, **n**on-response to therapy, new complication or **E**IM).

Patients with depressive symptoms were also more likely to reach these composite endpoints: in the univariable analysis, the AFFSST endpoint (HR: 1.57, CI: 1.23-1.99, p<0.001, **Table 5**) as well as the FNCE endpoint (HR: 1.37, CI: 1.07-1.74, p=0.012, **Table 5**) were significantly associated with preceding depressive symptoms. Correction for control variables barely changed results and significance was maintained (**Figure 4**).

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### 409 Effects of SNPs on IBD activity over time

We tested whether rs588765-TC or rs2522833's C allele could be directly implicated as protective or risk factors for clinical deterioration (**Table 6**). In univariable models, the TC allele combination of rs588765 showed significant protective effects against active disease (CDAI  $\geq$ 150/ MTWAI  $\geq$ 10, HR: 0.78, CI: 0.61-0.99, p=0.042). These significant protective effects of rs588765's TC alleles remained significant after controlling for confounders (**Figure 1; panel B**).

In contrast, the C allele of rs2522833 was not a significant risk factor for active
disease (univariable: HR: 1.08, CI: 0.92-1.28, p=0.344, multivariable:
Supplementary figure 5; panel B).

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### 420 Genetic inheritance patterns of rs588765 and rs2522833

421 In a sensitivity analysis we examined the impact of different allele variants of 422 rs588765 on outcomes of interest. Similarly as for initial screening, the

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423	overdominance model (TC vs. TT or CC) seemed to provide the best fit for Cox
424	proportional models with depressive symptoms over time and active disease as
425	endpoints (Supplementary tables 3 and 4, Supplementary figure 8; panel A and
426	C). Similarly, for rs2522833, the log-additive inheritance model which uses the
427	number of C alleles (0, 1 or 2) seemed to be the best fit regarding active disease
428	(Supplementary table 4, Supplementary figure 8; panel D), but not regarding
429	depressive symptoms over time (Supplementary table 3, Supplementary figure 8;
430	panel B). No formal statistical testing comparing the fits of genetic models was done.

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# 431 **Discussion**

432 In our cross-sectional analysis of depressive symptoms in IBD patients, we identified 433 variants of two SNPs (rs588765-TC, rs2522833's C allele) that were significantly 434 (q<0.10) associated with depression in IBD. The presence of depressive symptoms 435 was a strong risk factor for subsequent clinical deterioration in time-to-event 436 analyses, confirming and extending earlier findings by showing robust negative 437 effects on numerous clinical endpoints as well as two previously published composite endpoints for clinical deterioration.<sup>10,21</sup> Finally, we could demonstrate that 438 439 the TC allele of rs588765 was protective regarding depression and IBD activity in a 440 time-to-event analysis.

IBD patients are a high-risk group for developing depression.<sup>6</sup> In line with these findings, we observed an elevated point prevalence of depressive symptoms in our cohort (77/814, 9.5% among CD patients and 40/618, 6.5% among UC/IC patients) compared to the general population. For instance, point prevalence in a non-clinical Swiss sample with comparable assessment was estimated to be only 3.1%,<sup>40</sup> similar to findings in the USA<sup>41</sup> and the UK.<sup>26</sup>

447 We found a significant negative association between depressive symptoms and 448 patients with a specific variant in one of the preselected SNPs (rs588765-TC, ORs of 449 0.43, q=0.05) in a cross-sectional analysis with enrolment data. This is an 450 observation that, to the best of our knowledge, had not been described before. 451 Moreover, the time-to-event analysis indicated a protective effect of the same allele 452 combination regarding new or persistent depressive symptoms over time (aHR: 0.67, 453 p=0.035), thus independently confirming protective properties of rs588765-TC. 454 Finally, protective effects of rs588765-TC regarding depression translated to

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protection regarding IBD flares (CDAI ≥150/MTWAI ≥10, aHR: 0.72, p=0.045). All analyses were corrected for smoking status and alcohol consumption indicating that protection regarding depression and flares is independent of these features.

458 Rs588765 is part of the chr15q25.1 region and represents a well-established tag-SNP on the CHRNA5-CHRNA3-CHRNB4 gene cluster<sup>42,43</sup> that encodes the nAChR 459 460 subunits  $\alpha 5$ ,  $\alpha 3$ , and  $\beta 4$ . Rs588765 is located in the neuronal acetylcholine receptor subunit  $\alpha$ 5 gene (CHRNA5).<sup>44</sup> CHRNA5's gene product serves as a subunit of 461 pentameric nAChRs.<sup>45</sup> Rs588765 impacts the expression of CHRNA5 in human 462 463 brain tissue<sup>46</sup> with subjects homozygous with two minor alleles (TT) showing the highest CHRNA5 mRNA expression.<sup>46</sup> NAChRs are ligand-gated ion channels that 464 465 mediate signal transmission at synapses and modulate the release of different neurotransmitters.<sup>47</sup> Addition of an a5 subunit to nAChRs a4g2, the most frequent 466 467 receptor subtype in the brain, increases the rate of receptor desensitization and calcium permeability.<sup>47</sup> Similarly to the brain, CHRNA5 seems to mediate excitatory 468 nicotinic cholinergic transmission in the enteric nervous system (ENS).<sup>48</sup> Rs588765 469 had been associated with nicotine and alcohol dependence,<sup>15,16</sup> also among SIBDC 470 patients.<sup>17</sup> Moreover, the associated CHRNA5-CHRNA3-CHRNB4 region has also 471 472 been suggested to contain risk genes for affective disorders such as major depressive disorder <sup>49-51</sup> and recent findings suggest that nAChRs in general are 473 involved in psychiatric disorders such as depression.<sup>14</sup> 474

The genetic model of overdominance we stated best fitting for rs588765 refers to advantageous effects of the heterozygous variant compared to homozygous variants.<sup>52</sup> Examples include protective effects of hemoglobin mutation against malaria of in human sickle cell anaemia.<sup>53</sup> For rs588765, functional interactions of genetic variants associated with the T and C allele, respectively, seem possible as a

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480 variant of epistasis. Another plausible explanation for overdominance is the concept
481 of an inverted U-shaped response curve.<sup>54</sup>

482 The second SNP (rs2522833) associated with depressive symptoms is part of the 483 piccolo presynaptic cytomatrix protein (PCLO) gene. The presynaptic cytomatrix 484 describes a complex electron-dense meshwork of proteins involved in transmitter release.<sup>55,56</sup> Moreover, PCLO has been implicated to potentially contributing to short-485 term neuronal plasticity.<sup>48</sup> The PCLO gene has been associated with depressive 486 conditions such as major depressive disorder<sup>57</sup> and the C allele of rs2522833 487 specifically might increase an individual's vulnerability regarding depression.<sup>58</sup> Our 488 489 current results align well with these earlier findings and the C allele was a significant 490 (q<0.1) log-additive risk factor for depressive symptoms at baseline (ORs: 1.73, 491 q=0.059). However, our study did not detect significance for rs2522833 in the time-492 to-event analyses.

493 In our analysis, depressive symptoms remained a strong and highly significant risk 494 factor for future clinical deterioration according to several clinical measures including 495 the well-established disease scores CDAI and MTWAI and two pre-defined, partially overlapping composite flare definitions (AFFSST<sup>10</sup> and FNCE<sup>21</sup>, respectively). 496 497 Depressive symptoms also increased the likelihood for future occurrence of nearly all 498 other IBD-related endpoints tested over time (Figures 2 and 3), even though without 499 uniform statistical significance. This suggests that the specific definition for a flare is 500 largely irrelevant for the association with depression.

A growing body of evidence indicates that stress in general<sup>11,21</sup> and depressive symptoms specifically<sup>10</sup> are associated with clinical flares in IBD patients. Whether the psychological state of an IBD patient can indeed directly increase gut inflammation remains an important question in IBD research.<sup>59</sup> Our results support a

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505 direct impact of depressive symptoms on gut inflammation: i) we show strong 506 associations between rs588765, depressive symptoms and IBD flares (Tables 3-6); 507 ii) depressive symptoms remain a strong risk factor for flares according to various 508 definitions (CDAI/MTWAI: aHR: 3.25, p<0.001, AFFSST: aHR: 1.62, p<0.001, FNCE: 509 aHR: 1.35, p=0.019) and iii) the TC allele of rs588765 with protective effects 510 regarding depressive symptoms (aHR: 0.67, p=0.035) was also protective against 511 IBD flares (CDAI/MTWAI: aHR: 0.72, p=0.045). The combination of clinical and 512 genetic data strengthens this point; however, indirect effects cannot be entirely 513 excluded. For instance, depression might be a marker for subclinical inflammation or 514 might affect reporting of symptoms by patients. Another interesting possibility could 515 be an effect of rs588765-TC on CHRNA5 expression in the gut and the brain 516 simultaneously, thus independently explaining depressive symptoms and flares. 517 Further mechanistic studies would be necessary to better understand our 518 observation.

519 Our study has several strengths and limitations. Strengths include the excellent 520 clinical characterization regarding IBD-related outcomes and the sequential 521 measurements of the key variable *depressive symptoms* and the clinical endpoint 522 variables over a long time enabling the combination of cross-sectional and time-to-523 event analyses. Limitations are the relatively small sample size for genetic analyses, 524 yielding in lower statistical power for some analyses. Moreover, missing data points, 525 e.g. for therapy or surgery data, further reduced statistical power for some analyses. 526 Furthermore, even though serial measurements for HADS-D were available and a 527 HADS-D score of ≥11 represents a well-established cut off strongly indicating depression,<sup>23-26</sup> a diagnosis of depression normally requires direct clinical 528 529 assessment by a psychiatrist. Furthermore, for validation of our data, independent

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confirmation in another IBD cohort would have been desirable. Finally, in 2019, a
large meta-analysis for the risk of depression including 807,553 individuals was
published. This study identified 102 independent variants (not rs588765, rs2522833)
of which 87 were replicated in an independent sample.<sup>13</sup> Therefore, future studies
with a larger number of SNPs will now be possible.

535 In conclusion, our study confirms an intimate relationship between depressive symptoms and IBD disease course. The prevalence of depression among IBD 536 patients is high and active disease further increases the hazard for depression.<sup>6</sup> Vice 537 538 versa, depressive symptoms also independently increased the hazard for IBD flares 539 over time. Moreover, two SNPs (rs588765-TC, rs2522833's C allele) were 540 associated with depressive symptoms among IBD patients and one of those SNPs 541 (rs588765-TC) was also associated with high IBD activity. These findings are in 542 accordance with a multidirectional relationship between genetics, depression and 543 IBD disease course. Therefore, a comprehensive therapeutic concept for IBD patients, targeting inflammation and depression might further improve patient care.<sup>60</sup> 544

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# 691 Tables

# 692 **Table 1**

Variables at enrolment	CD	UC/IC	P-value	
Median (1 <sup>st</sup> quartile-3 <sup>rd</sup> quartile				
Diagnosis		1137 (57.6%)	836 (42.4%)	
N=1973				
Sex	Male	567/1137 (49.9%)	452/836 (54.1%)	0.072
N=1973	Female	570/1137 (50.1%)	384/836 (45.9%)	
Age (years)		37.2 (24.6-50.0);	38.5 (26.8-49.0);	0.458
N=1973		6.5-85.5	4.6-81.8	
Time since diagnosis (years)		6.7 (1.7-15.4); 0-	4.9 (1.4-11.7); 0-	< 0.001
N=1970		52.5	49.5	
BMI		22.5 (19.9-25.7);	23 (20.7-25.9);	0.003
N=1942		12.3-48.1	13.1-48.8	
Smoking status	Smoker	301/929 (32.4%)	73/714 (10.2%)	< 0.001
N=1643	Non-smoker	628/929 (67.6%)	641/714 (89.8%)	
Daily alcohol consumption	Yes	703/820 (85.7%)	512/614 (83.4%)	0.251
N=1434	No	117/820 (14.3%)	102/614 (16.6%)	
Disease activity	CDAI	34.0 (11.0-80.0);		-
N=1973	N= 1137	0.0-450.0		
	MTWAI		2.0 (-0.0-3.4);	-
	N=836		0.0-19.0	
	Standardized	-0.3 (-0.7-0.6); -0.9-	-0.3 (-0.9-0.6); -	0.009
	CDAI/MTWAI #	7.3	0.9-4.6	
Active disease (CDAI ≥150 or	Yes	83/1137 (7.3%)	71/836 (8.5%)	0.373
MTWAI ≥10)				
N=1973				
	No	1054/1137 (92.7%)	765/836 (91.5%)	
Systemic steroids: yes	Yes	199/1124 (17.7%)	219/832 (26.3%)	< 0.001
N=1956	No	925/1124 (82.3%)	613/832 (73.7%)	
TNF inhibitors: yes	Yes	344/1124 (30.6%)	95/832 (11.4%)	< 0.001
N=1956	No	780/1124 (69.4%)	737/832 (88.6%)	
Number current therapies		1 (1-2): 0-6	2 (1-3): 0-7	< 0.001
N=1956			_ ( ,	
Disease related surgery prior	Yes	550/1137 (48.4%)	93/836 (11.1%)	< 0.001

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to enrolment	No	587/1137 (51.6%)	743/836 (88.9%)	
N=1973				
Fistula at or prior to	Yes	468/1137 (41.2%)	45/836 (5.4%)	< 0.001
enrolment	No	669/1137 (58.8%)	791/836 (94.6%)	
N=1973				
Stenosis at or prior to	Yes	366/1137 (32.2%)	18/836 (2.1%)	< 0.001
enrolment	No	771/1137 (67.8%)	818/836 (97.6%)	
N=1973				
EIM at or prior to enrolment	Yes	498/1137 (43.8%)	265/836 (31.7%)	< 0.001
N=1973	No	639/1137 (56.2%)	571/836 (68.3%)	
Complications at or prior to	Yes	562/1137 (49.4%)	356/836 (42.6%)	0.003
enrolment	No	575/1137 (50.6%)	480/836 (57.4%)	
N=1973				
rs588765	CC	448/1137 (39.4%)	278/836 (33.2%)	0.018
N=1973	TC	509/1137 (44.8%)	406/836 (48.6%)	
	TT	180/1137 (15.8%)	152/836 (18.2%)	
rs2522833	AA	419/1137 (36.9%)	300/836 (35.9%)	0.883
N=1973	AC	524/1137 (46.1%)	388/836 (46.4%)	
	СС	194/1137 (17.1%)	148/836 (17.7%)	
HADS score (A and D)		10 (5-16); 0-36	9 (5-15); 0-39	0.117
N=1432				
Depression (HADS-D ≥11)	Yes	77/814 (9.5%)	40/618 (6.5%)	0.052
N=1432	No	737/814 (90.5%)	578/618 (93.5%)	
Anxiety (HADS-A ≥11)	Yes	156/814 (19.2%)	93/619 (15.0%)	0.048
N=1432	No	658/814 (80.8%)	526/619 (85.0%)	

# Standardization of score values was obtained by norming values in reference to the respective score mean and standard deviation (disease activity score =  $(s_i - \bar{s}) / \sigma_s$ ). This results in comparable values between the different scores (CDAI/ MTWAI) that indicate a patients score value in reference to the mean score value of the respective score and taking each score's internal variance into account.

693 **Table 1: Study participants.** Characteristics of study participants at enrolment stratified for

diagnosis. Abbreviations: CD: Crohn's disease, UC: ulcerative colitis, IC: indeterminate

695 colitis, BMI: body mass index, CDAI: Crohn's Disease Activity Index, MTWAI: Modified

696 Truelove and Witts Severity Index, TNF: tumor necrosis factor, T: thymine, C: cytosine, A:

adenine, HADS: Hospital Anxiety and Depression Scale, HADS-D: subscale for depressive

698 symptoms, HADS-A: subscale for anxiety symptoms.

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#### Table 2 700

#### Univariable logistic regression analyses - odds ratios for depression in IBD patients

Dependent variable: depression (HADS depression sub score ≥11)			
Sex: male	0.91 (CI: 0.58-1.43)		
Diagnosis: UC/IC (vs. CD)	0.79 (CI: 0.50-1.26)		
Age (years)	1.01 (CI: 1.00-1.03)		
Time since diagnosis (years)	1.00 (CI: 0.97-1.02)		
BMI	1.07 (CI: 1.02-1.11) ***		
Smoking status: smoker	1.42 (Cl: 0.90-2.26)		
Daily alcohol consumption: yes	3.12 (CI: 1.24-7.83) **		
Disease activity index (standardized CDAI/MTWAI)	1.80 (CI: 1.53-2.13) ***		
Disease related surgery prior to enrolment	1.00 (CI: 0.64-1.62)		
Systemic steroids: yes	1.55 (CI: 0.95-2.54) *		
TNF inhibitors: yes	1.30 (CI: 0.78-1.59)		
Number current therapies	1.01 (CI: 1.00-1.03) ***		

#### 701 Table 2: Variables associated with depression (HADS-D ≥11) at baseline.

702 Univariable logistic regression models showing the association of possible

703 confounders for multivariable models with depression. Results are presented as

704 odds ratios; p-values are indicated as follows: \*: p<0.1, \*\*: p<0.05, \*\*\*: p<0.01. 1043

705 patients were analyzed. Abbreviations: CI: 95% confidence interval, IBD:

706 inflammatory bowel disease, HADS: Hospital Anxiety and Depression Scale, UC:

707 ulcerative colitis, IC: indeterminate colitis, CD: Crohn's disease, BMI: body mass

708 index, CDAI: Crohn's Disease Activity Index, MTWAI: Modified Truelove and Witts

709 Severity Index, TNF: tumor necrosis factor.

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# 711 **Table 3**

### Association analysis - odds ratios for depression in IBD patients

	Dependent variable: depression at enrolment (HADS depression sub score ≥11)			
	Univariable	Multivariable without variable	Multivariable with variable	
		elimination	elimination	
rs588765: TC (vs. CC/TT)	0.45 (CI: 0.28-	0.43 (CI: 0.26-0.72) *	0.43 (Cl: 0.25-0.72) **	
	0.74) *			
rs2522833: number of C	1.71 (CI: 1.25-	1.75 (Cl: 1.26-2.44) **	1.73 (Cl: 1.25-2.39) *	
alleles (0-2)	2.34) **			

Table 3: SNPs associated with depression (HADS-D ≥11). Table showing the analysis results of the two SNPs significantly (q<0.1) associated with depression at enrolment in IBD patients. Results are presented as odds ratios; q-values are indicated as follows: \*: q<0.1, \*\*: q<0.05, \*\*\*: q<0.01. Analyses: logistic regression analysis. Abbreviations: CI: 95% confidence interval, IBD: inflammatory bowel disease, HADS: Hospital Anxiety and Depression Scale, T: thymine, C: cytosine.

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# 719 **Table 4**

Cox proportional hazards model - hazard ratios for new or persistent depression among IBD patients

#### over time

	Endpoint variable: depression (HADS depression sub score ≥11)			
	Univariable	Multivariable	Multivariable	
		(rs588765)	(rs2522833)	
rs588765: TC (vs. CC/TT)	0.68 (CI: 0.46-0.99)	0.67 (CI: 0.46-0.97)		
	**	**		
rs2522833: number of C alleles (0-2)	1.12 (CI: 0.87-1.44)		1.10 (CI: 0.86-1.41)	
Sex: male		0.80 (Cl: 0.54-1.19)	0.79 (CI: 0.54-1.18)	
Diagnosis: UC/IC (vs. CD)		0.82 (Cl: 0.51-1.30)	0.80 (CI: 0.50-1.27)	
Time since diagnosis (years)		0.99 (Cl: 0.97-1.01)	0.99 (CI: 0.97-1.01)	
Age (years)		1.03 (CI: 1.01-1.04)	1.03 (CI: 1.01-1.04)	
		***	***	
BMI		1.00 (Cl: 0.95-1.04)	1.00 (CI: 0.96-1.04)	
Smoking status: smoker		1.63 (CI: 1.09-2.42)	1.65 (CI: 1.11-2.44) **	
		**		
Daily alcohol consumption: yes		1.21 (Cl: 0.71 -2.07)	1.24 (CI: 0.73 -2.12)	
Disease activity (standardized		1.53 (CI: 1.31-1.79)	1.52 (CI: 1.30-1.78)	
CDAI/MTWAI)		***	***	
Disease related surgery prior to		1.15 (CI: 0.74-1.78)	1.14 (CI: 0.73-1.78)	
enrolment				
Systemic steroids: yes		0.99 (Cl: 0.60-1.63)	0.99 (CI: 0.60-1.64)	
TNF inhibitors: yes		1.26 (Cl: 0.80-1.99)	1.25 (CI: 0.79-1.96)	
Number current therapies		1.07 (CI: 0.87-1.31)	1.06 (CI: 0.86-1.30)	

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# Table 4: Predictors for new or persistent depression (HADS-D $\geq$ 11) over time.

721 Results are presented as hazard ratios for new or persistent depression among IBD 722 patients over time; p-values are indicated as follows: \*: p<0.1, \*\*: p<0.05, \*\*\*: p<0.01. 723 919 patients were analyzed and 114 times the endpoint was reached. Analysis: Cox 724 proportional hazards model. Abbreviations: CI: 95% confidence interval, IBD: 725 inflammatory bowel disease, HADS: Hospital Anxiety and Depression Scale, T: 726 thymine, C: cytosine, UC: ulcerative colitis, IC: indeterminate colitis, CD: Crohn's 727 disease, BMI: body mass index, CDAI: Crohn's Disease Activity Index, MTWAI: 728 Modified Truelove and Witts Severity Index, TNF: tumor necrosis factor.

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# 730 Table 5

# Cox proportional hazards model - hazard ratios for active disease and clinical deterioration

#### (flares) in IBD patients over time

	Endpoint variable: active disease (CDAI/MTWAI)		Endpoint variable: combined clinical endpoint (AFFSST)		Endpoint variable: combined clinical endpoint (FNCE)	
	Univariable	Multivariable	Univariable	Multivariable	Univariable	Multivariable
Preceding	3.18 (CI:	3.25 (CI:	1.57 (CI:	1.62 (CI:	1.37 (CI:	1.35 (CI:
depression	2.19-4.60)	2.10-5.03)	1.23-1.99)	1.27 -2.08)	1.07-1.74) **	1.05 -1.72) **
over time (yes)	***	***	***	***		
Sex: male		0.86 (CI:		0.95 (CI:		0.89 (CI:
		0.62-1.20)		0.83-1.10)		0.77-1.03)
Diagnosis:		1.73 (CI:		0.89 (CI:		1.09 (CI:
UC/IC (vs. CD)		1.21-2.47)		0.76-1.05)		0.93-1.28)
		***				
Time since		1.01 (CI:		1.00 (CI:		1.00 (CI:
diagnosis		0.99-1.03)		0.99-1.01)		0.99-1.01)
(years)						
Age (years) #		1.00 (CI:		0.99 (CI:		1.00 (CI:
		0.99-1.01)		0.98-1.00) ***		0.99-1.00)
		***				
ВМІ		0.98 (CI:		1.00 (CI:		1.00 (CI:
		0.95-1.02)		0.98-1.01)		0.98-1.02)
Disease related		1.60 (CI:		1.02 (CI: 0.87		0.95 (CI:
surgery prior to		1.10 -2.32) **		-1.20)		0.80 -1.12)
enrolment						
Smoking		1.43 (CI:		1.04 (CI:		1.14 (CI:

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status: smoker	1.00-2.03) **	0.88-1.22)	0.97-1.34)
Daily alcohol	0.79 (CI:	1.17 (CI: 0.96	1.11 (Cl:
consumption:	0.51 -1.22)	-1.43)	0.90 -1.36)
yes			

# Values of 1 with simultaneous significance were obtained by rounding and are not equal to 1

731

732 Table 5: Predictors for active disease and clinical deterioration (flare) in IBD 733 patients over time. Results are presented as hazard ratio for active disease or 734 clinical deterioration (flare) respectively, in IBD patients over time. Active disease 735 was defined as CDAI ≥150 or MTWAI ≥10 in CD and UC patients, respectively. A 736 flare was defined by the AFFSST or FNCE composite endpoint; p-values are 737 indicated as follows: \*: p<0.1, \*\*: p<0.05, \*\*\*: p<0.01. Analysis: Cox proportional 738 hazards model. Abbreviations: CI: 95% confidence interval, IBD: inflammatory bowel 739 disease, HADS: Hospital Anxiety and Depression Scale, UC: ulcerative colitis, IC: 740 indeterminate colitis, CD: Crohn's disease, BMI: body mass index, CDAI: Crohn's 741 Disease Activity Index, MTWAI: Modified Truelove and Witts Severity Index, 742 AFFSST: active disease, physician reported flare, new fistula, new stenosis, surgery 743 or new systemic therapy, FNCE: physician reported flare, non-response to therapy, 744 new complication or EIM.

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# 746 **Table 6**

# Cox proportional hazards model - hazard ratios for active disease in IBD patients over time

	Endpoint variable: active disease (CDAI/MTWAI)				
	rs588765		rs2522833		
	Univariable	Multivariable	Univariable	Multivariable	
	(rs588765)	(rs588765)	(rs2522833)	(rs2522833)	
rs588765: TC (vs. CC/TT)	0.78 (Cl: 0.61-	0.72 (Cl: 0.52 -			
	0.99) **	0.99) **			
rs2522833: number of C			1.08 (CI: 0.92-	1.12 (CI: 0.90 -	
alleles (0-2)			1.28)	1.39)	
Sex: male		0.85 (CI: 0.61-		0.84 (Cl: 0.61-	
		1.18)		1.17)	
Diagnosis: UC/IC (vs. CD)		1.73 (Cl: 1.19-		1.70 (Cl: 1.17-	
		2.50) ***		2.46) ***	
Time since diagnosis		1.01 (CI: 0.99-		1.01 (CI: 0.99-	
(years)		1.03)		1.03)	
Age (years)		1.00 (CI: 0.98-		1.00 (CI: 0.98-	
		1.01)		1.01)	
BMI		0.99 (Cl: 0.95-		0.99 (Cl: 0.95-	
		1.03)		1.03)	
Disease related surgery		1.66 (CI: 1.14 -		1.63 (CI: 1.12 -	
prior to enrolment		2.42) ***		2.38) **	
Smoking status: smoker		1.63 (CI: 1.15-		1.64 (CI: 1.16-	
		2.31) ***		2.33) ***	

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Daily alcohol consumption:	0.90 (Cl: 0.59 -	0.88 (CI: 0.58 -
yes	1.38)	1.35)

Table 6: Predictors for active disease in IBD patients over time. Results are 747 748 presented as hazard ratio for active disease in IBD patients over time. Active disease 749 was defined as CDAI ≥150 or MTWAI ≥10 in CD and UC patients, respectively; pvalues are indicated as follows: \*: p<0.1, \*\*: p<0.05, \*\*\*: p<0.01. Analysis: Cox 750 751 proportional hazards models. Abbreviations: CI: 95% confidence interval, IBD: 752 inflammatory bowel disease, CDAI: Crohn's Disease Activity Index, MTWAI: Modified 753 Truelove and Witts Severity Index, C: cytosine, T: thymine, UC: ulcerative colitis, IC: 754 indeterminate colitis, CD: Crohn's disease, BMI: body mass index.

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# 756 **Figures**

### 757 **Figure 1**



759 Figure 1: Kaplan Meier curves showing how rs588765's genotype impacts the 760 hazards for depression and IBD activity over time. A: Percentage of IBD patients 761 without depression stratified for rs588765 genotype (overdominant). B: Percentage 762 of IBD patients without active disease defined as CDAI ≥150 or MTWAI ≥10 in CD 763 and UC patients, respectively and stratified for rs588765 genotype (overdominant). 764 Analysis: multivariable Cox proportional hazards models. Abbreviations: HR: hazard 765 ratio, CI: confidence interval, T: thymine, C: cytosine, IBD: inflammatory bowel 766 disease, CDAI: Crohn's Disease Activity Index, MTWAI: Modified Truelove and Witts 767 Severity Index.

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# 769 Figure 2



Figure 2: Kaplan Meier curves showing how depression increases the hazards for future adverse outcomes over time. Percentage of IBD patients without experiencing the respective outcome (A-H) stratified for depression. Analysis: multivariable Cox proportional hazards models. Abbreviations: HR: hazard ratio, CI: confidence interval, TNF: tumor necrosis factor, CDAI: Crohn's Disease Activity Index, MTWAI: Modified Truelove and Witts Severity Index.

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# 778 Figure 3



Figure 3: Kaplan Meier curves showing how depression increases the hazards for future EIMs over time. Percentage of IBD patients without experiencing the respective outcome (A-H) stratified for depression. Analysis: multivariable Cox proportional hazards models. Abbreviations: HR: hazard ratio, CI: confidence interval, PSC: primary sclerosing cholangitis.

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# 786 Figure 4



788 Figure 4: Kaplan Meier curves showing how depression increases the hazard 789 for future clinical deterioration (flare, composite endpoints) over time. A: 790 Percentage of IBD patients without a flare according to the AFFSST-definition, 791 stratified according to the presence of depression. B: Percentage of IBD patients 792 without a flare according to the FNCE-definition, stratified according to the presence 793 multivariable Cox proportional hazards depression. Analysis: models. of 794 Abbreviations: HR: hazard ratio, CI: confidence interval, IBD: inflammatory bowel 795 disease, AFFSST: active disease, physician reported flare, new fistula, new stenosis, 796 surgery or new systemic therapy, FNCE: physician reported flare, non-response to 797 therapy, new complication or EIM.