

# 1 Comorbidities of keloid and hypertrophic scars in UK Biobank

2 **Title:** Comorbidities of keloid and hypertrophic scars in UK Biobank

3 **Subtitle:** Comorbidities of excessive scarring

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24 **Key points**

25 Question: What diseases are people with keloid and hypertrophic scars (excessive scarring) at risk of?

26 Findings: This cross-sectional UK Biobank study identified associations of excessive scarring with  
27 hypertension, vitamin D deficiency and atopic eczema. Atopic eczema was also significantly associated  
28 when accounting for confounders, and the associations varied with ethnicity. A phenome-wide scan  
29 of this predominantly white cohort identified a range of previously unreported correlations;  
30 musculoskeletal disease and pain symptoms were prominent non-dermatological associations.

31 Meaning: Excessive scarring may share underlying predispositions with systemic diseases; future  
32 research should aim to understand whether there are causal relationships linking the observed  
33 associations.

34 **Tweet**

35 What diseases associate with keloid/hypertrophic scars? This UK Biobank study confirmed excessive  
36 scarring associates with hypertension, vitamin D deficiency and atopic eczema, and discovered  
37 musculoskeletal disease & pain symptoms are also comorbidities.

## 38 **Abstract**

39 Importance: Keloids and hypertrophic scars (excessive scarring) are relatively understudied disfiguring  
40 chronic skin conditions with high treatment resistance.

41 Objective: To evaluate established comorbidities of excessive scarring in Europeans, with comparisons  
42 across ethnic groups, and to identify novel comorbidities via a phenome-wide association study  
43 (PheWAS).

44 Design: This cross-sectional study used UK Biobank (UKB) data and fitted logistic regression models for  
45 testing associations between excessive scarring and a variety of outcomes, including previously  
46 studied comorbidities and 1518 systematically defined disease categories. Additional modelling was  
47 performed within subgroups of participants defined by self-reported ethnicity.

48 Setting: Multi-centre population-based cohort study.

49 Participants: Of 502701 UKB participants, analyses were restricted to 230078 individuals with linked  
50 primary care records.

51 Exposures: Keloid or hypertrophic scar diagnoses

52 Main outcomes and measures: Previously studied disease associations (hypertension, uterine  
53 leiomyoma, vitamin D deficiency, atopic eczema) and phenotypes defined in the PheWAS Catalog.

54 Results: Of the 972 people with excessive scarring, there was a higher proportion of females  
55 compared to the 229106 controls (65% versus 55%) and a lower proportion of white ethnicity (86%  
56 versus 95%). Associations were identified with hypertension, vitamin D deficiency and atopic eczema  
57 in models accounting for age, sex and ethnicity, and the association with atopic eczema (Odds Ratio  
58 (OR) 1.68,  $p < 0.001$ ) was remained statistically significant after accounting for additional potential  
59 confounders. Fully-adjusted analyses within ethnic groups revealed associations with hypertension in  
60 black participants (OR 2.05,  $p = 0.019$ ) and with vitamin D deficiency in Asian participants (OR 2.24,  
61  $p = 0.006$ ). The association with uterine leiomyoma was borderline significant in black women (OR  
62 1.93,  $p = 0.05$ ) whereas the association with atopic eczema was significant in white participants (OR  
63 1.68,  $p < 0.001$ ) and showed a similar trend in Asian (OR 2.17,  $p = 0.048$ ) and black participants (OR 1.89,  
64  $p = 0.13$ ). The PheWAS identified 110 significant associations across disease systems; of the non-  
65 dermatological, musculoskeletal disease and pain symptoms were prominent.

66 Conclusions and Relevance: We validated comorbidities of excessive scarring in an independent  
67 cohort with comprehensive coverage of health outcomes. We also documented additional phenome-  
68 wide associations that will serve as a reference for future studies to investigate common underlying  
69 pathophysiologic mechanisms.

70

## 71 **Introduction**

72 Keloids and hypertrophic scars are chronic disfiguring manifestations of excessive cutaneous wound  
73 healing considered prototypic of skin fibrosis.[1–3] Whether they are distinct entities or a quantitative  
74 extreme of normal wound scars remains an active area of debate.[4] Both conditions involve excess  
75 extracellular matrix deposition and raised scar tissue, but unique to keloids are horizontal expansion  
76 of scars and absence of clinical regression. Common to both are high symptom burden[5,6] and lack  
77 of universally effective treatment.[7,8]

78 There is increasingly a shift from single-disease focused research towards studying disease systems.[9]  
79 Understanding disease comorbidities has both biological and clinical benefits, such as highlighting  
80 novel mechanisms and offering opportunities for targeted and early clinical intervention. Previous  
81 studies of keloid and hypertrophic scar comorbidities have been limited to candidate diseases, based  
82 on speculated biological[10–15] or demographic similarities.[16,17]

83 In this study, we comprehensively assessed disease associations of keloids or hypertrophic scars  
84 (henceforth excessive scarring) in UK Biobank (UKB), a multi-centre population-based longitudinal  
85 observational study of >500,000 participants including >950 with a diagnosis of excessive scarring. We  
86 performed multivariable logistic regression analyses for previously-studied comorbidities across  
87 several ethnic groups and carried out a phenome-wide association study (PheWAS) over a wide range  
88 of systematically-defined diseases.

## 89 **Methods**

90 This project used UKB data under project number 15147. The UKB study was approved by the  
91 National Health Service National Research Ethics Service (11/NW/0382).

### 92 **Study population**

93 UKB is a large population-based prospective cohort study that recruited >500,000 participants (40-69  
94 years at recruitment) who attended 1 of 22 assessment centers across the UK from 2006-2010.[18]  
95 Volunteers provided written informed consent for their participation. Rich health-related information

96 is available, including regularly updated self-reported health conditions, lifestyle indicators,  
97 anthropometric and biological measurements. Longitudinal health record data is available, including  
98 hospital episode statistics and primary care data. Participants were also asked to report their ethnicity  
99 based on a set of predefined ethnic categories, further detailed in  
100 <https://biobank.ctsu.ox.ac.uk/crystal/field.cgi?id=21000>. We restricted our analyses to individuals  
101 with linked-primary care records (limiting the possibility of misclassifying participants as unaffected  
102 due to missing data).

### 103 Ascertainment of disease status

104 Clinical code selection - Disease status was ascertained through the following data sources: self-  
105 reported (verbal interview), linked-hospital episode statistics [International Classification of Diseases  
106 (ICD), ninth and tenth revisions], cancer register (ICD9 and ICD10), primary care records (Read2 and  
107 Read3), and the Office of Population Censuses and Surveys Classification of Interventions and  
108 Procedures (OPCS4). Clinical codelists were manually curated by study authors (CYU, ANW) for  
109 excessive scarring (described below), vitamin D deficiency and atopic eczema (Supplementary Table  
110 2). For conditions with pre-existing manually-curated codelists (hypertension, uterine leiomyoma),  
111 Read2 and ICD10 clinical codelists were obtained from the CALIBER portal.[21] These clinical codelists  
112 (Supplementary Table 2) were minimally adapted and mapped to Read3 and ICD9 equivalents  
113 respectively, using UKB Resource 592 (<https://biobank.ndph.ox.ac.uk/ukb/refer.cgi?id=592>).

114 Excessive scarring status - A broad definition of “excessive scarring” was used, constituting a diagnosis  
115 of keloid or hypertrophic scar, which can be difficult to clinically differentiate[19,20] and are not well-  
116 distinguished by electronic health record (EHR) coding systems. Primary care data included diagnostic  
117 codes specific to either keloid or hypertrophic scar whereas those from linked-hospital episodes  
118 statistics could pertain to either scar type. The final codes selected were: 7014 (ICD9), L910 (ICD10),  
119 7L19A (Read2), M218. (Read2), M214. (Read2 and Read3), M2y11 (Read2), XaC07 (Read3), XaPxn  
120 (Read3) and X78TS (Read3; Supplementary Table 2). Within this group, we additionally identified  
121 individuals who had likely received scar-related treatment to define a more homogenous cohort  
122 with moderate-to-severe scarring (hereafter “treated excessive scarring”), defined by codes for scar  
123 excision/refashioning and triamcinolone treatment (Supplementary Table 3).

124 Comorbidities/outcomes - A systematic search for excessive scarring disease associations was  
125 performed on Medline using the following query: (case-control studies/ or cohort studies/ or (Risk  
126 Factors or (comorbidity or Comorbidity) or comorbidities or (Prevalence or prevalence) or association  
127 or predispose or risk).mp.) and (hypertrophic scar.mp. or Cicatrix, Hypertrophic/ or ((Keloid or keloid  
128 or keloids) not Acne Keloid).m\_titl.). This resulted in 708 references that were independently  
129 reviewed for relevance. From the 21 remaining references (Supplementary Table 1), disease  
130 associations selected for analysis were those studied in  $\geq 2$  independent reports, namely  
131 hypertension, uterine leiomyoma, vitamin D deficiency and atopic eczema.

132 PheWAS - We used the PheWAS Catalog with phecodes that represent a single phenotype and  
133 corresponding groupings.[22] The same data sources used for clinical code selection were mapped to  
134 ICD10 codes and subsequently mapped to 1518 phecodes using Phecode Map 1.2.

#### 135 Statistical analyses

136 Descriptive statistics are presented as frequencies with percentages for categorical variables and  
137 means with standard deviations for continuous variables. Pearson's Chi-squared test and Welch's t-  
138 test were used for comparisons between categorical and continuous variables, respectively. Statistical  
139 significance was set at  $p < 0.05$ .

140 Significant comorbidity associations were further examined using multivariable logistic regression  
141 analyses to estimate odds ratios (OR) and 95% confidence intervals (CIs). In each model, excessive  
142 scarring was the exposure variable, and the comorbidity was the outcome. As we were testing four  
143 primary comorbidities (hypertension, uterine leiomyoma, vitamin D deficiency, atopic eczema), a  
144 Bonferroni-adjusted p-value threshold ( $0.05/4=0.0125$ ) was used.

145 We noted 3403 individuals (1.2% of the study cohort) with missing data for at least one of age, sex,  
146 ethnicity, Townsend Deprivation Index (TDI), body mass index (BMI) or smoking status. Participants  
147 with missing data were more likely to be male and of non-white ethnicity, with higher TDI and BMI  
148 (Supplementary Table 4). There were no significant differences in missingness between participants  
149 with and without excessive scarring diagnoses(Supplementary Table 5). Nevertheless, for testing  
150 associations, the full cohort of 230078 participants was analysed and multiple imputation (20 imputed  
151 datasets) was used to account for missing data.[23]

152 To investigate the independent association with excessive scarring, a “minimal model” adjusting for  
153 age, sex (except uterine leiomyoma, which was restricted to female participants only, n=125771) and  
154 ethnicity, and a “full model” adjusting for the additional potential confounding covariates were fitted  
155 for each disease association. The additional covariates were: (1) hypertension: BMI, TDI, smoking  
156 status, diabetes, hyperlipidaemia[24] (2) uterine leiomyoma: BMI[25,26], TDI, smoking status (3)  
157 vitamin D deficiency: BMI, TDI, smoking status[27,28] and (4) atopic eczema: BMI, TDI, allergic  
158 rhinitis, asthma[29].

159 To assess whether the disease associations varied by ethnicity, separate fully-adjusted logistic  
160 regressions were fitted in the three largest ethnic groups. These comprised self-reported white  
161 participants (85.9%), black or black British participants (henceforth “black participants”, 6.4%) and  
162 Asian or Asian British participants (henceforth “Asian participants”, 5.2%), as described in Study  
163 Population. To assess whether any differences in OR across ethnicities could be attributable to  
164 chance, the analysis was repeated in the full cohort with an interaction term between excessive  
165 scarring status and ethnicity.

166 For the PheWAS, logistic regressions were performed to assess the association between excessive  
167 scarring and the prevalence of each phecode diagnosis, adjusting for age, sex, ethnicity, smoking  
168 status, BMI and TDI. A Bonferroni-adjusted p-value threshold of  $0.05/1518=3.3\times 10^{-5}$  was used.  
169 Phecodes with <200 cases or controls were excluded.

170 All statistical analyses were performed with R version 4.1.2. Specific R packages included mice,[23]  
171 gtsummary,[30] tidyverse, [31] flextable,[32] PheWAS,[33] targets,[34] ukbwranglr[35] and  
172 codemapper.[36]

## 173 **Results**

174 We analysed 230078 UK Biobank participants for whom linked GP data were available  
175 (Supplementary Figure 1). 972 participants had a record of excessive scarring (740 with a diagnostic  
176 code specific for keloid, 110 specific for hypertrophic scar, 177 for either keloid or hypertrophic scar;  
177 Supplementary Figure 2). The prevalence of excessive scarring for the three largest ethnic groups  
178 were: 1.1% (Asian), 2.4% (black) and 0.4% (white). Table 1 shows the baseline characteristics for the  
179 excessive scarring and comparator groups. In the excessive scar-affected group, there was a higher



180 proportion of females than in the unaffected group (65% versus 55%) and a lower proportion of  
181 participants with self-reported white ethnicity (86% versus 95%). Of the 972 participants with  
182 excessive scarring, 106 had a record of scar-related treatment. There were no notable differences in  
183 baseline characteristics between these participants and those without a record of scar-related  
184 treatment (Supplementary Table 6).

185 Previously studied associations with excessive scarring

186 All previously studied comorbidities (hypertension, uterine leiomyoma, vitamin D deficiency, atopic  
187 eczema) were more prevalent for individuals with excessive scarring (Table 1). Vitamin D deficiency  
188 and atopic eczema were around twice as common in the excessive scar cohort: 5.1% versus 2.4%  
189 (vitamin D deficiency) and 10% versus 5.9% (atopic eczema). The higher prevalence for uterine  
190 leiomyoma and hypertension was less marked: 15% versus 11% (leiomyoma) and 37% versus 34%  
191 (hypertension). We found no evidence for differences in prevalence of the primary comorbidities  
192 between treated excessive scarring cases and the rest of the excessive scarring group (Supplementary  
193 Table 6) but with fewer cases, no statistically significant associations were observed when considering  
194 the treated excessive scarring subgroup against the whole unaffected cohort (Supplementary Table  
195 7).

196 Each comorbidity was analysed as a disease outcome using two multivariable logistic regression  
197 models: minimally-adjusted (for age, sex and ethnicity except for uterine leiomyoma which was tested  
198 within females only adjusting for age and ethnicity) and fully-adjusted (with additional potential  
199 confounders for each comorbidity; Table 2). Statistically significant associations with excessive  
200 scarring were observed for hypertension and atopic eczema in the minimally-adjusted models, while  
201 the association with vitamin D deficiency fell short of Bonferroni-corrected significance (OR 1.42 (95%  
202 CI 1.05-1.93),  $p=0.02$ ). In fully-adjusted models, only the association with atopic eczema (OR 1.68  
203 (95% CI 1.36-2.07),  $p<0.001$ ) remained significant. Despite a positive effect-size estimate, there was  
204 no significant association between excessive scarring and uterine leiomyoma (OR 1.19 (95% CI 0.95-  
205 1.49),  $p=0.13$ ).

206 We then performed association testing within subgroups of participants defined by self-reported  
207 Asian (53 excessive scar affected, 4654 unaffected), black (61 affected, 2448 unaffected), and white  
208 (829 affected, 217502 unaffected) ethnicity (Table 3A). With the exception of atopic eczema, our

209 analysis suggested a divergence between ethnic groups in the prevalence of each comorbidity and its  
210 association with excessive scarring (Table 3B). The associations with hypertension and uterine  
211 leiomyoma were nominally significant in black participants [OR 2.05 (95% CI 1.13-3.72),  $p=0.019$   
212 (hypertension) and OR 1.93 (95% CI 1.00-3.71),  $p=0.05$  (uterine leiomyoma)] and not significant in  
213 Asian or white participants. Vitamin D deficiency was only significantly associated with excessive  
214 scarring in Asian participants (OR 2.24 (95% CI 1.26-3.97),  $p=0.006$ ). For atopic eczema, the  
215 association with excessive scarring was highly significant in white participants (OR 1.68 (95% CI 1.34-  
216 2.12),  $p<0.001$ ), nominally significant in Asian participants (OR 2.17 (95% CI 1.01-4.67),  $p=0.048$ ), and  
217 although not statistically significant in black participants, exhibited a similar trend (OR 1.89 (95% CI  
218 0.83-4.28),  $p=0.13$ ). Finally, to formally assess these differences we fitted a full logistic regression  
219 model incorporating an interaction between excessive scarring status and ethnicity, finding statistical  
220 evidence for ethnicity-specific effect sizes in the case of hypertension in black participants (relative to  
221 white participants,  $p=0.049$ ).

## 222 Discovery analysis

223 We screened 1518 phecodes across 17 disease groups, identifying 110 diseases significantly enriched  
224 among participants with excessive scarring (Figure 1, Table 4, Supplementary Table 8).

225 There was strongest evidence of association for several dermatological diseases, most prominently  
226 sebaceous cyst (OR 2.56,  $p=9.45\times 10^{-30}$ ), non-epithelial skin cancer (OR 2.89,  $p=2.03\times 10^{-25}$ ) and the  
227 umbrella phenotype “diseases of hair/hair follicles” (OR 2.3,  $p=1.50\times 10^{-22}$ ), as well as infections of  
228 skin/subcutaneous tissue, seborrheic keratosis, actinic keratosis, acne, and notably, atopic/contact  
229 dermatitis, all with  $OR>1.9$  and  $p<1.0\times 10^{-11}$ . Similarly strong evidence was observed for pain-related  
230 symptoms, particularly for joint pain (OR 1.84,  $p=1.87\times 10^{-20}$ ) but also back pain, cervicalgia,  
231 enthesopathies and mastodynia, all with  $OR>1.6$  and  $P<1.0\times 10^{-12}$ . Significant associations with the  
232 largest effect sizes were abnormal weight gain (OR 3.97,  $p=9.32\times 10^{-8}$ ) and heart valve replacement  
233 (OR 3.9,  $p=6.65\times 10^{-7}$ ).

234 Associations were identified with hypertension (OR 1.26,  $p=2.05\times 10^{-3}$ ) and vitamin D deficiency (OR  
235 1.47,  $p=1.34\times 10^{-2}$ ) as expected, but these did not meet Bonferroni-corrected significance. Other  
236 previously reported associations that did not meet our selection criteria for specific analysis were  
237 explored, including obesity, osteoporosis, skin cancers, pancreatic cancer, migraine and asthma

238 (Supplementary Table 1). Of these, statistically significant associations were observed for skin cancers  
239 (melanoma, OR 3.17,  $p=3.33 \times 10^{-11}$ ) and migraine (OR 1.53,  $p=6.29 \times 10^{-6}$ )(Supplementary Table 9).

## 240 **Discussion**

241 To date there have been few large-scale association studies for excessive scarring.[10–13,15] This  
242 study aimed to both validate previously studied associations (hypertension,[16,37,38] uterine  
243 leiomyoma,[13,17] vitamin D deficiency[14,15] and atopic eczema[11,12]) as well as scan for  
244 excessive scarring associations across the phenome. Our ethnicity-specific analysis represents the first  
245 comprehensive study of excessive scarring in white people, the most-represented ethnic group in  
246 UKB.

247 In the trans-ethnic UKB population, we replicated associations with three (hypertension, vitamin D  
248 deficiency and atopic eczema) of the four primary comorbidities previously studied two or more  
249 times. Only the associations with hypertension and atopic eczema were statistically significant after  
250 correcting for multiple tests. The association with hypertension was attenuated after adjusting for  
251 additional risk factors (BMI, TDI, smoking status, diabetes, hyperlipidaemia). This suggests the  
252 observed difference in hypertension prevalence is linked to differences in these risk factors between  
253 people with and without excessive scarring. However, in our cross-sectional analysis we are unable to  
254 determine the causal relationship between excessive scarring and increased hypertension risk factor  
255 burden.

256 Subgroup analyses revealed possible ethnic variations in comorbidity risk, particularly for  
257 hypertension in black participants, in whom we found a significantly larger effect size than in white  
258 participants. Although our relatively small subgroup sample sizes make robust interpretation  
259 challenging, given the well-established disproportionate burden of diseases within the respective  
260 ethnic groups,[25,39–41] we propose that there may be ethnicity-specific risk determinants that are  
261 shared by these pathologies. For example, vascular dysfunction is thought to contribute to the severe  
262 hypertension and hypertensive heart failure specifically affecting the black population.[42,43]  
263 Abnormal endothelial function and microvascular architecture are also observed in keloids.[47]  
264 Replication of these findings in a larger cohort may make a case for the early identification of  
265 cardiovascular disease in black individuals with excessive scarring.

266 The only association that showed nominally significant evidence for association in multiple ethnic  
267 groups was atopic eczema. Interestingly, this has previously been reported in Taiwanese[12] and  
268 Korean[11] populations, and we now observe this association across all three of our broadly defined  
269 ethnic groupings; however, regional variations in disease prevalence and associations may still  
270 emerge. Atopic eczema skin is more likely to be excoriated and scars may thus be an epiphenomenon.  
271 Nevertheless, our result adds epidemiological support to the hypothesis that the Th2 inflammatory  
272 axis contributes to keloid pathogenesis.[48]

273 Previous non-genetic applications of PheWAS's[54–57] have been based only on ICD diagnostic codes.  
274 This would have excluded a large proportion of our cases who were only identified through primary  
275 care codes. Through our comprehensive strategy to maximize identification of people with excessive  
276 scarring, we highlighted numerous significant disease associations, potentially indicating an increased  
277 risk of poorer health outcomes.

278 The frequent female genitourinary disease associations are consistent with suggestions that sex  
279 hormones may play a role in keloid pathophysiology.[58–60] Potentially, this could be explained by  
280 the over-representation of female participants within the excessive scar-affected group; however, our  
281 models adjusted for sex. The highly significant associations with dermatological conditions and  
282 neoplasms may represent true predispositions or reflect ascertainment bias (i.e. if a patient presents  
283 with a dermatological condition or is reviewed post-surgically, a scar-related diagnosis is more likely  
284 to be recorded).

285 Of the dermatological associations, diseases of the pilosebaceous unit (sebaceous cyst, diseases of  
286 hair/hair follicles, acne) support the 'sebum hypothesis'[61] which is based on high sebaceous gland  
287 density observed in keloid-prone skin[62] and sebum being intrinsically pro-inflammatory.[63] The  
288 association of keloid with skin cancer has been previously reported.[10] Although plausible reasons  
289 have been proposed including similar bioenergetics (reliance on glycolysis)[64,65] and signalling  
290 pathways including TGF $\beta$ /Smad[66–68] and Wnt/beta-Catenin,[69–72] this finding is interpreted  
291 cautiously, again considering the risk of ascertainment bias.

292 The associations with musculoskeletal disorders (enthesopathy, pain in joint, back pain, cervicalgia)  
293 may support the observation of chondrogenic misdifferentiation in keloids[73] and the shared  
294 significance of TGF $\beta$  in joint pathologies.[74,75] Interestingly, associations with pain symptoms

295 spanned disease categories (non-specific chest pain, irritable bowel syndrome, mastodynia, acute  
296 pain, headache syndromes, pain in joint, back pain, cervicalgia). Pain is known to debilitate some  
297 keloids sufferers; [76] whether there is shared underlying biopsychosocial dysfunction with other pain  
298 entities or whether they may be mutually reinforcing is speculative. Nonetheless, chronic pain  
299 represents a major global burden of disease[77] and proactive identification of these conditions may  
300 aid patient counseling and treatment decisions.

301 Finally, whether an individual whose skin scars excessively is at risk of excessive internal scarring  
302 remains unanswered. In our study, the association of peritoneal adhesions with excessive scarring  
303 carried an OR of 3.68 ( $p=1.19 \times 10^{-4}$ ), which is intriguing but this is based only on 617 cases of  
304 peritoneal adhesions, nine of whom had excessive scarring.

## 305 **Limitations**

306 Although this study utilized a large biobank cohort, a relatively small sample size of excessive scarring  
307 cases were identified ( $n=972$ ). This is particularly relevant when attempting to dissect differences  
308 between ethnic groups, as there is less statistical power to detect significant associations in groups  
309 with lower sample numbers. However, it is reassuring that disease prevalence in our dataset is in  
310 keeping with currently available epidemiological studies.[59] As most participants report white  
311 ethnicity, the main findings, particularly from the PheWAS, may not be generalizable to other ethnic  
312 groups for whom excessive scarring is a more prominent issue.

313 From our PheWAS, there are two further discussion points. The lack of significant associations with  
314 other fibrotic/scarring comorbidities (e.g. lung or liver fibrosis) supports previous reports[13,78]  
315 although low case numbers may mean our investigations were insufficiently powered to detect them.  
316 Secondly, it was striking that all significant associations were positive (i.e. increased prevalence of  
317 comorbidities in people with excessive scarring), potentially a result of coverage bias whereby  
318 participants with more complete coverage of linked health data may be more likely to have a record  
319 of excessive scar diagnosis as well as a diagnosis of any other comorbidity. This may mean that the  
320 effect sizes are overestimated, however, the relative order of the associations remains informative.  
321 We do not draw causal conclusions from our findings; rather, we inform on co-existing relationships  
322 between diseases that may not have previously been appreciated.

323 Our inclusion of keloids and hypertrophic scars in the definition of excessive scarring results in a study  
324 population that may be heterogeneous with respect to disease severity (and potentially  
325 pathophysiology). It might be expected that more severe cases of excessive scarring should be  
326 associated with a higher burden of comorbidity. We undertook to formally analyse this by  
327 distinguishing a subset of treated excessive scar cases. This attempt to refine the study population to  
328 those with moderate-to-severe keloid scarring yielded only 106 individuals, limiting our power to  
329 detect statistically significant disease associations and conduct further analyses. This highlights how  
330 data availability and information or misclassification bias are key challenges of phenotyping based on  
331 EHR. Using currently available clinical codes within UKB, we were only able to establish excessive  
332 scarring cases based on potentially subjective clinical assessment without clinicopathological  
333 correlation. Nonetheless, our results add to what is already known in the literature, as evidenced by  
334 the detection of both previously reported and novel associations.

## 335 **Conclusions**

336 We report a comprehensive observational analysis of UKB participants with excessive scarring, using a  
337 robust modelling approach with adjustments for a variety of disease confounders and associated risk  
338 factors. Previously reported disease associations for excessive scarring (hypertension, vitamin D  
339 deficiency and atopic eczema) were replicated, but only the association with atopic eczema showed a  
340 similar trend across the three major ethnic subgroups (Asian, black and white participants). Our  
341 PheWAS implicates a range of unreported associations for reference when studying the  
342 pathophysiology of excessive scarring and may prove valuable in studying the associated disease  
343 areas.

344

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348

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557

558 **Figure Legend**

559 **Figure 1:** Multivariate logistic regressions to estimate the effect of excessive scarring status on the risk  
560 of each phecode diagnosis, adjusting for age, sex, ethnicity, smoking status, body mass index and  
561 Townsend Deprivation Index. Dots represent phecodes and colours represent systemic categories.  
562 Statistical significance was set at  $p < 0.05/1518 = 3.3 \times 10^{-5}$  (based on the number of phecodes tested).

**Table 1: Baseline characteristics for the scar and comparator groups.**

Characteristic	Study Participants <sup>1</sup>	Keloid or Hypertrophic Scar Affected <sup>1</sup>	Keloid or Hypertrophic Scar Unaffected <sup>1</sup>	p-value <sup>2</sup>
<b>Participant numbers</b>	230,078	972	229,106	
<b>Age</b>	64 (8)	63 (8)	64 (8)	<b>&lt;0.001</b>
<b>Sex</b>				<b>&lt;0.001</b>
Female	125,771 (55%)	633 (65%)	125,138 (55%)	
Male	104,307 (45%)	339 (35%)	103,968 (45%)	
<b>Ethnic Background</b>				<b>&lt;0.001</b>
Asian or Asian British	4,707 (2.1%)	53 (5.5%)	4,654 (2.0%)	
Black or Black British	2,509 (1.1%)	61 (6.3%)	2,448 (1.1%)	
Chinese	600 (0.3%)	7 (0.7%)	593 (0.3%)	
Mixed	1,167 (0.5%)	7 (0.7%)	1,160 (0.5%)	
Other ethnic group	1,686 (0.7%)	10 (1.0%)	1,676 (0.7%)	
White	218,331 (95%)	829 (86%)	217,502 (95%)	
<b>Townsend Deprivation Index</b>	-1.33 (3.03)	-1.29 (3.12)	-1.33 (3.03)	0.7
<b>Ever Smoked</b>	135,946 (59%)	526 (54%)	135,420 (59%)	<b>0.001</b>
<b>Body Mass Index</b>	27.5 (4.8)	27.9 (5.2)	27.5 (4.8)	0.051
<b>Hypertension</b>	79,034 (34%)	362 (37%)	78,672 (34%)	0.062
<b>Uterine leiomyoma<sup>3</sup></b>	14,080 (11%)	92 (15%)	13,988 (11%)	<b>0.009</b>
<b>Vitamin D deficiency</b>	5,547 (2.4%)	50 (5.1%)	5,497 (2.4%)	<b>&lt;0.001</b>
<b>Eczema, atopic</b>	13,501 (5.9%)	99 (10%)	13,402 (5.8%)	<b>&lt;0.001</b>

<sup>1</sup>Mean (standard deviation); n (%)

<sup>2</sup>Welch Two Sample t-test; Pearson's Chi-squared test; Statistical significance is declared at P < 0.05 (bold)

<sup>3</sup>Only female participants (N total = 125,771, N affected = 633, N unaffected = 125,138) considered for uterine leiomyoma

565 **Table 2: Associations between excessive scarring and selected**  
 566 **comorbidities.**

Outcome	Minimal Model			Full Model		
	OR	95% CI	p-value	OR	95% CI	p-value
Hypertension	1.24	1.08, 1.43	<b>0.002</b>	1.11	0.96, 1.30	0.2
Uterine leiomyoma	1.20	0.96, 1.51	0.11	1.19	0.95, 1.49	0.13
Vitamin D deficiency	1.42	1.05, 1.93	0.022	1.47	1.08, 1.99	0.013
Atopic eczema	1.78	1.44, 2.19	<b>&lt;0.001</b>	1.68	1.36, 2.07	<b>&lt;0.001</b>

OR = Odds Ratio, CI = Confidence Interval

Minimal model adjusts for age, sex (except uterine leiomyoma) and ethnicity for the entire imputed cohort. Full model adjusts for age, sex (except uterine leiomyoma), ethnicity and additional confounders for the entire imputed cohort (see Methods). Models for uterine leiomyoma are restricted to females only without adjusting for sex.

Statistical significance is declared at  $P < 0.05/4 = 0.0125$  (bold). Nominal statistical significance is italicized.

567

568 **Table 3A. Summary of comorbidities in cases and controls within the three**  
 569 **main UKB self-reported ethnic groups.**

Characteristic	White participants			Black participants			Asian participants		
	Control, N = 217,502	Case, N = 829	p- value <sup>1</sup>	Control, N = 2,448	Case, N = 61	p- value <sup>1</sup>	Control, N = 4,654	Case, N = 53	p- value <sup>1</sup>
<b>Hypertension</b>	74,175 (34%)	290 (35%)	0.6	1,056 (43%)	38 (62%)	<b>0.004</b>	1,995 (43%)	23 (43%)	>0.9
<b>Uterine leiomyoma<sup>2</sup></b>	12,843 (11%)	63 (12%)	0.6	458 (33%)	18 (47%)	0.083	291 (13%)	4 (14%)	>0.9
<b>Vitamin D deficiency</b>	3,866 (1.8%)	20 (2.4%)	0.2	263 (11%)	6 (9.8%)	>0.9	1,041 (22%)	20 (38%)	<b>0.013</b>
<b>Eczema, atopic</b>	12,621 (5.8%)	83 (10%)	<b>&lt;0.001</b>	144 (5.9%)	7 (11%)	0.12	323 (6.9%)	8 (15%)	<b>0.042</b>

<sup>1</sup> Pearson's Chi-squared test; Statistical significance is declared at P < 0.05 (bold).

<sup>2</sup> Only female participants considered for uterine leiomyoma [N control (white) = 119,053, N case (white) = 544; N control (black) = 1,404, N case (black) = 38; N control (Asian) = 2,157, N case (Asian) = 29].

570 **Table 3B. Fully adjusted model investigating associations between**  
 571 **excessive scarring and selected comorbidities within the three main UKB**  
 572 **self-reported ethnic groups.**

Outcome	White participants			Black participants			Asian participants		
	OR <sup>1</sup>	95% CI <sup>1</sup>	p- value <sup>2</sup>	OR <sup>1</sup>	95% CI <sup>1</sup>	p- value <sup>2</sup>	OR <sup>1</sup>	95% CI <sup>1</sup>	p- value <sup>2</sup>
Hypertension	1.08	0.92, 1.28	0.3	2.05	1.13, 3.72	<i>0.019</i>	0.95	0.48, 1.86	0.9
Uterine leiomyoma <sup>3</sup>	1.07	0.82, 1.39	0.6	1.93	1.00, 3.71	<i>0.050</i>	1.04	0.36, 3.01	>0.9
Vitamin D deficiency	1.32	0.85, 2.07	0.2	0.88	0.37, 2.08	0.8	2.24	1.26, 3.97	<b>0.006</b>
Atopic eczema	1.68	1.34, 2.12	<b>&lt;0.001</b>	1.89	0.83, 4.28	0.13	2.17	1.01, 4.67	<i>0.048</i>

<sup>1</sup> OR = Odds Ratio, CI = Confidence Interval

<sup>2</sup> Statistical significance is declared at P < 0.05/4 = 0.0125 (bold). Nominal/borderline significance is italicized.

<sup>3</sup> Model for uterine leiomyoma is restricted to females only without adjusting for sex.

573



574 **Table 4: Phecodes significantly associated with keloid and hypertrophic**  
 575 **scar status.**

Group	Phecode description
Genitourinary (15)	<b>Mastodynia</b> ; Hypertrophy of breast (Gynecomastia); Inflammatory disease of breast; Cystitis; Frequency of urination and polyuria; Lump or mass in breast; Irregular menstrual cycle; Excessive or frequent menstruation; Ovarian cyst; Noninflammatory disorders of vagina; Noninflammatory disorders of vulva and perineum; Premenstrual tension syndromes; Urinary tract infection; Renal failure NOS; Inflammatory diseases of uterus, except cervix
Dermatologic (14)	<b>Sebaceous cyst; Diseases of hair and hair follicles; Atopic/contact dermatitis due to other or unspecified; Other local infections of skin and subcutaneous tissue; Seborrheic keratosis; Actinic keratosis; Acne</b> ; Rash and other nonspecific skin eruption; Other hypertrophic and atrophic conditions of skin; Hyperhidrosis; Disturbance of skin sensation; Carbuncle and furuncle; Pruritus and related conditions; Diseases of nail, NOS
Respiratory (12)	<b>Acute upper respiratory infections of multiple or unspecified sites; Cough; Acute sinusitis</b> ; Other diseases of respiratory system, NEC; Acute pharyngitis; Acute laryngitis and tracheitis; Influenza; Shortness of breath; Pneumonia; Pneumonia due to fungus (mycoses); Chronic sinusitis; Chronic pharyngitis and nasopharyngitis
Infectious diseases (11)	<b>Postoperative infection</b> ; Tuberculosis; Viral warts & HPV; Dermatophytosis of nail; Sexually transmitted infections (not HIV or hepatitis); Dermatophytosis; Candidiasis; Herpes simplex; Viral infection; Mycoses; Gram negative septicemia
Sense organs (11)	<b>Conjunctivitis, infectious</b> ; Dizziness and giddiness (Light-headedness and vertigo); Inflammation of eyelids; Eustachian tube disorders; Disorders of lacrimal system; Otagia; Otitis externa; Infection of the eye; Hearing loss; Conjunctivitis, noninfectious; Tinnitus
Neoplasms (8)	<b>Other non-epithelial cancer of skin</b> ; Melanomas of skin; Malignant neoplasm, other; Lipoma; Other benign neoplasm of connective and other soft tissue; Benign neoplasm of lip, oral cavity, and pharynx; Malignant neoplasm of female breast; Nevus, non-neoplastic
Symptoms (8)	<b>Malaise and fatigue; Back pain; Cervicalgia</b> ; Abdominal pain; Nausea and vomiting; Swelling of limb; Sciatica; Edema
Circulatory system (7)	Angina pectoris; Nonspecific chest pain; Other chronic ischemic heart disease, unspecified; Coronary atherosclerosis; Heart valve replaced; Endocarditis; Orthostatic hypotension
Musculoskeletal (7)	<b>Pain in joint; Peripheral enthesopathies and allied syndromes; Enthesopathy</b> ; Bursitis; Other and unspecified disc disorder; Other disorders of soft tissues; Juvenile osteochondrosis

<b>Group</b>	<b>Phecode description</b>
Digestive (6)	Hemorrhage of rectum and anus; Symptoms involving digestive system; Irritable Bowel Syndrome; Diseases of lips; Anal and rectal polyp; Dysphagia
Neurological (5)	Acute pain; Organic or persistent insomnia; Other headache syndromes; Other peripheral nerve disorders; Migraine
Injuries & poisonings (3)	Sprains and strains; Contusion; Certain early complications of trauma or procedure
Endocrine/metabolic (2)	Abnormal weight gain; Hypercholesterolemia
Mental disorders (1)	Adjustment reaction

Significant associations with excessive scarring ( $p < 0.05/1518 = 3.3 \times 10^{-05}$ ), grouped by category (number of phecodes per category shown in brackets). The 20 most significantly associated phecodes are in bold.

