

1 **Title: The relative roles of maternal survival and inter-personal violence as selection pressures on**
2 **the persistence of Neanderthal hypercoagulability alleles in modern Europeans**

3 **Running title:** Do Neanderthal *SELP* alleles protect against maternal peri-partum blood loss?

4

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12

13 **Abstract**

14 Background

15 Simoni *et al* (2016) reported variation in the frequency of Neanderthal alleles found in modern
16 humans and argued that they may have provided an evolutionary advantage. One such allele is SNP
17 rs3917862, associated with hypercoagulability. rs3917862 can be deleterious but can also help
18 prevent blood loss. We investigated two possible selective pressure hypotheses for rs3917862
19 surviving to higher frequencies: deaths from interpersonal violent trauma and childbirth.

20 Results

21 Mortality data from modern hunter-gatherers models the living conditions and causes of death of
22 humans and Neanderthals at the point of admixture. National census data indicates a positive
23 correlation between presence of rs3917862 and decreased maternal mortality ratios. When
24 maternal mortality ratio is modelled using GDP, births attended by skilled assistants and the
25 presence of rs3917862, women are 0.1% more likely to die in childbirth in populations lacking
26 rs3917862. Deaths due to violence show no correlation with rs3917862.

27 Conclusion

28 These findings challenge the idea that Neanderthal admixture has negatively impacted the overall
29 health of modern humans. Maternal survival may have acted as a selective pressure for the
30 persistence of hypercoagulability alleles in modern Europeans. Understanding the role of
31 hypercoagulability in childbirth, and the role of rs3917862, could help to reduce maternal mortality
32 ratios.

33 **Keywords**

34 Ancient DNA; maternal mortality; hypercoagulation; Neanderthal; adaptive introgression

35 **Abbreviations**

36	AICc	Akaike information criterion with correction for small sample sizes
37	F5	Coagulation Factor V
38	GDP	Gross domestic product
39	IVMR	Interpersonal violence per 100,000 deaths per year
40	MMR	Maternal mortality ratio
41	Protein C	Autoprothrombin IIA

42 SBA Skilled birth assistance

43

44 **Word count:** 4593/8000

45 **Introduction**

46 The successful sequencing of the Neanderthal genome in 2010 revealed hitherto unsuspected levels
47 of gene flow from Neanderthals into modern humans, providing researchers with the chance to
48 examine the role this genetic admixture has had on modern human evolution (Green et al. 2010;
49 Prüfer et al. 2014). Interbreeding between Neanderthals and anatomically modern humans has
50 resulted in the genomes of modern humans (specifically non- Africans) containing between 1 and 4%
51 Neanderthal DNA (Green et al. 2010; Wall et al. 2013; Prüfer et al. 2014). The overlap between
52 anatomically modern humans and Neanderthals living outside Africa is estimated to have occurred
53 approximately 45,000-41,000 years ago and lasted for between 2,600 and 5,400 years (Higham et al.
54 2014) although the overlap may have been for as much as 30,000 years with multiple pulses of
55 admixture (Villanea & Schraiber 2019).

56 These interbreeding events have left a legacy of Neanderthal DNA which contributes to phenotypic
57 variation in modern human populations. Simonti (Simonti et al. 2016) reported the contribution of
58 common Neanderthal variants to over 1000 electronic health record-derived phenotypes in
59 approximately 28,000 adults of European ancestry. A non-coding SNP of Neanderthal origin,
60 rs3917862, located in an intron of P-selectin (*SELP*) on chromosome 1, was significantly associated
61 with hypercoagulability. It had functional genomic marks suggestive of regulatory function and was
62 significantly associated with the increased expression of *SELP* and *F5* in arteries which in turn
63 increases risk of diseases related to hypercoagulability such as deep vein thrombosis and embolism
64 (Simonti et al. 2016). The frequency of rs3917862 varies from 8.6% in CEU in Utah, 6.5% in Finnish in
65 Finland, 3.2% in Columbians from Medellin, Columbia, and 1.2% in Bengali from Bangladesh, to 0.1%
66 in Han Chinese in Beijing, and absent in African populations included in the 1000 Genomes Project
67 (Pybus et al. 2014; Gibbs et al. 2015). However, high-resolution frequency data for this allele on a
68 country-by-country basis is not currently available.

69 *SELP* codes for a cell adhesion protein, which is expressed on the endothelial cells and platelets that
70 recruits leukocytes to injuries during inflammation (Simonti et al. 2016). *F5* codes for factor V, a
71 coagulation cofactor which circulates in plasma and combines with factor X to activate prothrombin
72 (Rallapalli et al. 2014). Rallapalli (Rallapalli et al. 2014) showed that the 11 coagulation factors in
73 modern humans have undergone positive selection. Simonti concludes that there must have some
74 advantage to the Neanderthal haplotype and that hypercoagulability would have been beneficial to
75 early anatomically modern humans moving out of Africa (Simonti et al. 2016).

76 It is significant that rs3917862 has persisted given that hypercoagulability has been shown to have
77 negative health impacts, potentially detrimentally affecting reproductive fitness. For example,
78 hypercoagulability leads to increased risk of clot formation. Clot formation in turn can lead to venous
79 thromboembolism – pulmonary embolism and deep vein thrombosis (Heit 2007). Malfunction of F5,
80 a mutation called Factor V Leiden, increases thrombophilia and is found in approximately 20% of
81 cases of venous thromboembolism (Rosendaal & Reitsma 2009). This mutation increases
82 thrombophilia by causing amino acid substitution at one of the activated protein C cleavage sites of
83 Factor V, rendering it resistant to activated protein C inactivation, and so leads to a reduction in the
84 natural anticoagulant system, and enhances thrombin production (De Stefano & Leone 1995;
85 Hooper & De Staercke 2002; Martinelli et al. 2010; Kyrle et al. 2010). Hypercoagulability, often used
86 as a synonym for thrombophilia, increases the risk of venous thromboembolism. Increased risk of
87 venous thromboembolism as a result of rs3917862 would, in turn, decrease the reproductive fitness
88 carriers of this SNP, as it affects humans before and during their reproductive years.

89 However, hypercoagulability is not the only risk factor for venous thromboembolism. Old age,
90 sedentism, obesity, air travel, oral contraceptives, surgery and cancer are all aspects of modern life
91 that are also risk factors for venous thromboembolism (Rosendaal 2005; Prandoni et al. 2005; Heit
92 2007; Rosendaal & Reitsma 2009; Martinelli et al. 2010; Kyrle et al. 2010; Yang et al. 2012).
93 Therefore, while hypercoagulability is a risk factor for contracting venous thromboembolism,
94 modern life increases the occurrence of other predisposing factors, suggesting that
95 hypercoagulability today is much more likely to contribute to the development of deep vein
96 thrombosis or pulmonary embolism than it would have some 50,000 years ago. In situations where
97 people were injured frequently (or otherwise dying from blood loss) hypercoagulability would be
98 beneficial, as it would reduce the risk of death due to blood loss. It is these circumstances that are
99 considered here, in order to determine which selective pressure could result in the persistence of
100 introgressed Neanderthal hypercoagulability alleles in modern humans.

101 In order to understand the circumstances that would have resulted in hypercoagulability being
102 advantageous, we must understand the living conditions of Neanderthals and anatomically modern
103 humans approximately 50,000 years ago. Anthropologists have studied this in two ways: by
104 observing the fossil record and through modelling (using modern hunter-gatherers as a model of
105 likely living conditions before the advent of agriculture). Berger and Trinkaus (Berger & Trinkaus
106 1995) suggested that the frequency of trauma on Neanderthal skeletons is abnormally high and
107 suggested that the trauma patterns on Neanderthal skeletons are similar to those seen in North
108 American rodeo performers, as the injuries are predominantly to the upper body, which they argue
109 might be evidence that Neanderthals came into close contact with large ungulates, presumably as a

110 result of close-quarter hunting. Trinkaus (Trinkaus 2012) later builds on this argument with the
111 theory that Middle Palaeolithic spears were heavy and well suited for defence against carnivores and
112 for ambush hunting. However, he also purports that the incidence of impact fractures on projectiles
113 is suggestive of hunting at a distance (Trinkaus 2012). He offers two alternatives to hunting as the
114 cause of these patterns of injury: interpersonal violence and foraging. Trinkaus (Trinkaus 2012)
115 suggests that a number of the incidences of trauma show distinct signs of interpersonal violence: for
116 example Zollikofer found evidence in the St Cesaire Neanderthal that the skull had been struck with
117 a sharp implement, presumably during an act of interpersonal violence (Zollikofer et al. 2002).
118 However, Underdown (Underdown 2006) argues that while Neanderthal trauma was common it
119 was not abnormally high for hunter-gatherer populations and that patterns of trauma on
120 Neanderthal skeletons are not related to hunting or interpersonal violence, suggesting instead that
121 the trauma profile is most similar to that of the largely vegetarian Holocene Australian Aborigine
122 population, and that the evidence of trauma is a result of accidents. Trinkaus (Trinkaus 2012) does
123 suggest that a highly mobile foraging lifestyle could explain the predominantly upper body
124 distribution of injuries, as injuries to the lower limbs would have prevented mobility and resulted in
125 the injured being left behind in locations where remains would not have fossilised. Beier (Beier et al.
126 2018) analysed cranial trauma in 114 Neanderthal and 90 *Homo sapiens* skulls from the Upper
127 Palaeolithic and showed that while males had a higher incidence of trauma than females there was
128 no significant difference between the species. Importantly their work shows that it is not species
129 that causes increased vulnerability to trauma but rather the risks posed by hunter-gathering in the
130 Eurasian Palaeolithic.

131 Modern data sets from hunter-gatherers provide an insight into Neanderthal trauma and hominin
132 mortality profiles. Marlowe (Marlowe 2005) suggests that there are several difficulties in using
133 hunter-gatherer data in modelling hominin lifestyles before the advent of agriculture. Firstly,
134 modern hunter-gatherers are likely to be a biased sample, as they exist only in marginal habitats not
135 sought by agriculturalists. Secondly, it is highly unlikely that any of these forager groups exist entirely
136 isolated from interaction with their agricultural neighbours, and so they are not living as our pre-
137 agricultural ancestors were. However, Marlowe also makes the point that the ethnography of
138 modern hunter-gatherers is the only way we can observe human behaviour in the absence of
139 agriculture and so is the best proxy for the study of human behavioural and cultural evolution
140 (Marlowe 2010). Once again, anthropologists are divided on what the hunter-gatherer evidence tells
141 us about violence as a selective pressure. Hill and colleagues (HILL et al. 2007) use data from the
142 Hiwi of Venezuela to suggest that high adult mortality rates in Neanderthal and early modern human
143 populations are best explained by high levels of conspecific violence, as the Hiwi have lower rates of

144 infectious disease mortality, which more closely resembles the mortality profile of our Palaeolithic
145 ancestors. If this is the case, then such high levels of interpersonal violence may have been a
146 selective pressure for the persistence of hypercoagulability alleles. However, Wrangham and
147 colleagues (Wrangham et al. 2006) suggest that mortality rates due to violence in forager and
148 subsistence farming societies are highly variable. Indeed, they found that subsistence farming
149 societies had higher rates of death due to interpersonal violence than hunter-gatherer societies
150 (Wrangham et al. 2006). By this measure, violence would not be a consistent selective pressure.
151 While anthropologists are largely divided on the presence and nature of violent and accidental
152 deaths in Neanderthals and early modern humans, it is worth considering as a selective pressure for
153 the persistence of hypercoagulability alleles in living Europeans.

154 Our hypothesis is that maternal mortality would present a more consistent selective pressure for the
155 persistence of rs3917862. There are several pieces of evidence that suggest that the Neanderthal
156 *SELP* SNP was maintained in the gene pool because it has advantages for maternal survival. Firstly,
157 the main cause of death during childbirth in Africa and Asia is haemorrhaging (Ronsmans et al.
158 2006): indicating that in environments with limited healthcare blood loss is the most significant
159 factor in maternal mortality. This is evidence to suggest that hypercoagulability plays a protective
160 role against maternal haemorrhage during childbirth. Secondly, pregnancy is a risk factor for both
161 hypercoagulability and venous thromboembolism (Rosendaal 2005; Heit 2007; Martinelli et al. 2010;
162 Bourjeily et al. 2010). This hypercoagulability reduces the risk of excessive blood loss and death by
163 haemorrhaging during delivery (Rosendaal 2005; Heit 2007; Martinelli et al. 2010; Bourjeily et al.
164 2010). Further evidence to suggest that hypercoagulability caused by rs3917862 might have
165 persisted due to maternal mortality is that both *F5* and *SELP* are expressed in the placenta (Uhlén et
166 al. 2015). While *SELP* RNA is expressed in many different tissues in the body, *F5* RNA is only
167 expressed in the liver, gallbladder and placenta (Uhlén et al. 2015). Together this indicates that
168 maternal mortality is a possible selective pressure for the persistence of rs3917862.

169 **Materials and methods**

170 Data collection

171 The data collected fall into four different categories: comparative causes of death for five modern
172 hunter-gatherer groups, proportion of deaths caused by violence in fifteen hunter-gatherer
173 populations, maternal mortality ratios (MMR) in different populations in Tanzania, and modern
174 national census data.

175 Values for MMR, percentage of births attended by skilled birth assistants (SBA) and number of
176 deaths due to interpersonal violence per 100,000 deaths per year (IVMR) were taken from the GBD
177 2015 SDG Collaborators (2016). GDP per capita data came from the World Bank (2016) for 188
178 countries. The presence or absence of Neanderthal allele rs3917862 was estimated from data from
179 the 1000 Genomes Project which showed that the SNP was present in populations in Asia as well as
180 Europe (Green et al. 2010; Aken et al. 2016). Allele frequency data is only available for 1000
181 Genomes populations (Gibbs et al. 2015) and per-country allele frequency data is not currently
182 available.

183 Data analysis

184 Proportion of deaths from childbirth, interpersonal violence and accidents were expressed as
185 percentages to ensure the deaths between populations were comparable, and were corrected for
186 only females being at risk of maternal mortality. Hunter-gatherer mortality data (the percentage of
187 deaths caused by childbirth, interpersonal violence and accidents) for five hunter-gatherer
188 populations was visualised in bar plots using R (R Core Team 2013) (the San Ildefonso Agta (Early &
189 Headland 1998), the Hiwi (HILL et al. 2007), the Hadza (Blurton Jones et al. 2002), the Ache (Hill &
190 Hurtado 1996), and the Aka (Hewlett et al. 1986)). The number of deaths by interpersonal violence
191 per 100,000 per year, as calculated by Kelly (Kelly 2013) was plotted as bar charts in order to
192 compare the rates of violent deaths for fifteen hunter-gatherer populations (the Hadza (Marlowe
193 2010), the Andamanese (Keeley 1996), the Ju/'hoansi speaking !Kung (Lee 1979), the San Ildefonso
194 Agta (Early & Headland 1998), the Gidjingali aborigine population (Hiatt 1965), the Tiwi (Keeley
195 1996), the Yahgan (Cooper 1917), the Yurok (Keeley 1996), the Casigurian Agta (Headland 1989), the
196 Yulngu aborigine population (Keeley 1996), the Modoc (Keeley 1996), the Ache (Hill & Hurtado
197 1996), the Hiwi (HILL et al. 2007), the Piegan (Keeley 1996) and the Batek (Endicott & Endicott 2008)
198 in (Kelly 2013)).

199 The authors calculated the MMR value for the Hadza (Blurton Jones 2016), and compared this data
200 to the average MMR of Tanzania (Blurton Jones 2016), and the MMR values for the Haydom
201 Lutheran Hospital in the Mbulu highlands (Olsen et al. 2002) and the Kilimanjaro Christian Medical
202 Centre (Armon 1979).

203 The R package rworldmap (South 2011) was used to plot the MMR values and the rate of deaths due
204 to interpersonal violence (IVMR) for 184 countries. R was used to plot the variables MMR, IVMR,
205 SBA, GDP per capita and presence or absence of the allele rs3917862 against each other. Kendall's
206 Rank Correlation Tau was used to determine variable correlation. R package ppcor (Kim 2015) was
207 used to test the partial correlation of each pair of variables, with respect to the other variables, to

208 see whether any partial correlation was evident in the Kendall's Tau. R was then used to generate
209 linear and polynomial regressions to model nine different potential explanations of the impact of
210 SBA, allele presence and GDP on MMR. R package MuMIn (Bartón 2018) was used to calculate the
211 AICc value for each of the nine models, giving the Akaike information criterion, which gives a
212 measure of relative quality of a model. Adjusted AICc values were calculated by subtracting the
213 lowest AICc value from the others. Models were then ranked in order of most to least effective. The
214 best (most effective) model was visualised in three dimensions using the scatterplot3d package
215 (Ligges & Mächler 2003). SBA was plotted against MMR, colouring the points by the presence or
216 absence of rs3917862 in each population, and plotting the linear regression for those with the
217 Neanderthal allele and those without. A Mann-Whitney U test was conducted to determine whether
218 the two groups, those with the allele and those without, were significantly different in their central
219 tendency. The R script used to produce these figures and results can be found in **Supplementary**
220 **material**.

221 Results

222 The mortality profile data was used to compare the proportion of deaths due to
223 interpersonal violence, childbirth (for females) and accidents for the San Ildefonso Agta, the
224 Hiwi, the Hadza, the Ache and the Aka. **Table 1** and **supplementary figure 1** show that
225 deaths due to interpersonal violence account for a higher percentage of deaths overall than
226 either childbirth or accidents. However in the Agta, the Hadza and the Aka childbirth is
227 responsible for a higher proportion of female deaths than violence. Deaths due to violence
228 vary more widely between groups, while childbirth in this dataset is more consistent,
229 accounting for between 0-6.7% of deaths.

230 We further examined the rate of deaths due to interpersonal violence using the homicide
231 rate of fifteen hunter-gatherer groups. **Table 2** shows that the rate of deaths due to
232 interpersonal violence varies considerably between the different groups, with the lowest
233 rate 1 death per 100,000 people per year (the Batek), and the highest over 1000 deaths due
234 to homicide per 100,000 people per year.

235 We next considered the impact of hunter-gatherer lifestyle on death in childbirth,
236 comparing the maternal mortality ratios of four populations in Tanzania: the Hadza, the
237 Tanzanian national average, and two Tanzanian hospitals. **Table 3** shows that maternal
238 mortality is highest among the Hadza, which is higher than the national average. The two
239 hospitals show MMR values much lower than the national average.

240 Considering the hypothesised role of Neanderthal variant rs3917862 in survival due to
241 bloodloss-related trauma, we examined whether deaths due to interpersonal violence and
242 childbirth corresponded with the presence of rs3917862 outside Africa. **Figure 1** shows the
243 MMR values for 184 countries, and the rate of deaths due to interpersonal violence for the
244 same 184 countries. MMR correlates with the absence of the SNP in Africa, as the highest
245 mortality rates are within Africa and in East Asia. Deaths due to interpersonal violence, in
246 contrast, are highest in Russia, South America and parts of Africa.

247 **Table 4** shows the result of Kendall's Tau. We tested the correlation between MMR and SBA,
248 GDP per capita and presence/absence of the Neanderthal allele, and deaths due to
249 interpersonal violence and SBA, GDP per capita and presence/absence of the Neanderthal
250 allele. The correlations were significant for MMR and SBA, MMR and GDP per capita, and
251 MMR and the presence/absence of the Neanderthal allele. For deaths due to interpersonal
252 violence, correlations with GDP per capita and SBA were significant, however deaths due to
253 interpersonal violence did not significantly correlate with presence/absence of the
254 Neanderthal allele. To determine whether any of these correlations were a result of
255 correlation with other unaccounted-for variables, rather than as a result of direct
256 correlation, we ran partial correlation tests, again using Kendall's Tau. **Table 5** shows the
257 Tau and p-values for each pair of variables, controlling for each other variable in turn.
258 Correlations were still significant for all pairs of variables with MMR, however, deaths due
259 to interpersonal violence no longer correlated with skilled birth assistance when controlling
260 for GDP per capita. Therefore deaths due to interpersonal violence correlated only with GDP
261 per capita. As deaths due to interpersonal violence did not significantly correlate with
262 presence/absence of rs3917862, we did not model this relationship further.

263 To examine the correlation between MMR and the presence of rs3917862 further, we
264 modelled MMR using SBA, GDP, presence/absence of rs3917862 and log(GDP) as predictors.
265 **Table 6** shows the regression coefficients and p-value for each coefficient for all of the
266 models. **Table 7** shows the adjusted R^2 value for each of the nine models, the AICc values for
267 each model, the Δ AICc and finally the models are ranked from best to worst, in order to
268 determine which model best explains the variation in MMR. For this data the best model is:

$$269 \quad MMR = 970.52 - 522SBA - 104.37allele - 34.47\log(GDP)$$

270 which has an R^2 value of 0.7277, and uses SBA, presence/ absence of rs3917862 (*allele*) and the log
271 of GDP per capita to explain the variation in MMR. This equation suggests that those who come from

272 a population with the Neanderthal allele rs3917862 have 104.37/100,000 or approximately 0.1%
273 lower MMR, although this does not imply that the difference is caused by the presence of rs3917862

274 We visualised the best model using a three-dimensional scatterplot, in order to better understand
275 the relationship between the different predictors. **Figure 2** shows that MMR is generally lower, and
276 SBA generally higher for populations with the Neanderthal allele as compared to those without,
277 while GDP has little visible correlation. As GDP did not visibly cluster with higher or lower rates of
278 maternal mortality in our data, we plotted a two-dimensional scatter plot of SBA and MMR. **Figure 3**
279 shows that there is a difference in MMR between those populations that have the Neanderthal
280 variant and those who do not. Indeed, when not accounting for log(GDP) it appears a though there is
281 a difference of 125.3 deaths in childbirth per 100,000 live births between countries that have
282 rs3917862 represented and those who do not. In order to further investigate this difference, we
283 subdivided the data into the nations that have the Neanderthal SNP and those who do not. We
284 tested the difference in the central tendency of these two subsets using a Mann-Whitney Wilcoxon
285 test, in order to determine whether these groups were significantly different. The test returned a p-
286 value of <2.2e-16, which would suggest that the countries with rs3917862 are statistically
287 significantly different in the central tendency of the distribution of MMR values.

288 Discussion

289 Intronic Neanderthal SNP Rs391782 appears to have a regulatory function, resulting in increased
290 expression of *SELP* and *F5*, which in turn results in a hypercoagulability phenotype. Using modern
291 hunter-gatherer data, we examined what selective pressures pre-agricultural lifestyles would have
292 exerted to potentially cause the persistence of the hypercoagulability phenotype. The heightened
293 risk of maternal morbidity and mortality is well established (Ortner 1998) and this effect is increased
294 in the developing world where access to medical care can be severely reduced. In high income
295 countries the risk of maternal death is 1 in 3,3300 while in low income it is 1 in 41 (WHO 2015).
296 Prehistoric human populations would similarly have faced a much greater risk of death from child
297 birth than those in the modern developed world. Death during pregnancy can occur from a wide
298 range of complications such as infection, eclampsia, obstructed labour and haemorrhage but it is
299 largely impossible to measure the relative risk of these in archaeological populations (Ortner 2003).
300 Yet in modern data haemorrhage is the leading cause of maternal mortality accounting for 27% of
301 deaths (Say et al. 2014). However, pregnancy and life threatening complications arising from it are
302 extremely hard to detect in the archaeological record via skeletal evidence. Palaeopathologists have
303 suggested that changes to the pubic bone adjacent to the pubic symphysis in the form of pitting and
304 erosion are diagnostic of parturition (e.g. (Stewart 1968) and (Gilbert & McKern 1973)) but there

305 remains little definitive evidence to support this association (Ortner 2003). More recent analysis in
306 Denmark (Poulsen et al. 2001) and Norway (Turner-Walker et al. 2001; Mays et al. 2006) linked
307 significant decrease in bone mineral density in young Danish mediaeval female skeletons with stress
308 caused by pregnancy and lactation. They argue that this increase in physiological stress would have
309 increased the risk of mortality. However, Agarwal (Agarwal et al. 2018) highlights that the while
310 skeletal change could be attributed to pregnancy, patterns of geographical variation must be
311 considered and in very northerly latitudes the impact of reproductive bone loss could be greatly
312 compounded by lack of vitamin D due to low levels of UV exposure. Ultimately pregnancy is
313 extremely difficult to identify in the archaeological record yet the pressures faced by populations in
314 the past are likely to be similar, if not much greater, than those faced by women in the developing
315 world. The data presented here were insufficient to prove any single selective pressure was
316 responsible for the persistence of rs3917862; however, we show that deaths due to interpersonal
317 violence were more frequent, but much less consistent, than deaths as a result of childbirth. This
318 suggests that childbirth-related trauma and deaths are an evolutionarily more consistent selection
319 pressure which could drive selection for hypercoagulability phenotypes, and examines a problem
320 that is very difficult to assess in the archaeological record alone.

321 National census data for death due to interpersonal violence and childbirth was used to examine
322 whether the Neanderthal allele rs3917862 was associated with either cause of mortality. Deaths due
323 to interpersonal violence did not correlate with the presence or absence of the Neanderthal SNP,
324 suggesting that deaths due to interpersonal violence are not an adequate hypothesis to explain the
325 persistence of this hypercoagulability allele. Death in childbirth, by contrast, did correlate with the
326 presence or absence of rs3917862, even when taking into consideration that both correlate with
327 GDP and with percentage of births attended by skilled birth assistants. In modelling the causes of
328 MMR, the best model was an additive model including three variables, GDP, percentage of births
329 attended by skilled birth assistants and presence or absence of the Neanderthal SNP. Even with
330 three variables, the presence or absence of the allele is significant in best explaining variation in
331 maternal mortality in our model. Populations with rs3917862 experienced 0.1% fewer deaths during
332 childbirth than populations without the allele. Together with the fact that haemorrhaging is still the
333 biggest cause of death in Africa and Asia (Ronsmans et al. 2006), *SELP* and *F5* are expressed in the
334 placenta and that pregnancy is already associated with increased coagulation in order to prepare the
335 body for childbirth, it remains possible that rs3917862 has persisted as a result of selection for
336 adaptations which reduces deaths due to blood loss during childbirth. However, other factors not
337 captured in the model may also correlate with the presence of rs3917862 and rs3917862 may be
338 acting as a proxy for other factors.

339 Limitations of this study

340 While these findings are undoubtedly valuable in generating new hypotheses to understand the
341 persistence of hypercoagulability alleles in modern Europeans, there are a number of caveats that
342 must be considered. Firstly, country-level data for the frequency of rs3917862, which ranges from 0
343 0.1-8.5% outside Africa for 1000 Genomes populations, is only available for a limited number of
344 countries and is taken from sometimes modest population sample sizes. Any study of this kind must
345 consider the issue of the binary nature of the presence or absence of the allele which was recorded
346 as a compromise to deal with missing data. Here we have assumed that, when there is no direct
347 value available on ensemble.org or SNPdb, the Neanderthal allele is present outside Africa, and
348 absent in Africa, as is typical in almost all Neanderthal admixture studies (Green et al. 2010; Prüfer et
349 al. 2014; Sankararaman et al. 2016). However, it is possible that this binary is representative of some
350 other difference between Africa and outside Africa other than GDP and skilled birth assistance that
351 has not been taken into account. Secondly, the census data set is at national level, and does not
352 allow further break down of the data to ethnic or linguistic groups, which means that estimations of
353 presence/absence of rs3917862 may be further affected by within-country, between-group
354 variation. For example, within China only Han Chinese are known to carry rs3917862, but this was
355 recorded in this study as the entirety of China having the presence of the allele (Aken et al. 2016).
356 Thirdly, the mortality data subset gives no indication of what the specific cause of death was, other
357 than the categories of interpersonal violence or death in childbirth. This increases the noise of the
358 dataset: for example, we do not know the proportion of deaths in each group caused by
359 haemorrhaging, infection, hypertension or other direct or indirect causes. The measure of births
360 attended by skilled birth assistants is also likely to be inflated, as this is census data, and births are
361 more likely to be registered if they are attended by a skilled birth assistant. With these caveats in
362 mind, the presence of Neanderthal allele rs3917862 is still associated with increased maternal
363 survival during childbirth, and childbirth-related blood loss and trauma/mortality and would have
364 acted as a far stronger selective pressure than that presented by stochastic incidences of
365 interpersonal violence. The problems inherent in identifying pregnancy related changes to the
366 skeleton or skeletal markers of obstetric complications has created a false picture of how selection
367 would have operated on populations in the past. The method adopted here allows us to actively
368 generate new hypotheses that ameliorate the issues presented by the skeletal data.

369

370 Ultimately our results indicate a new hypothesis for the persistence of hypercoagulability alleles and
371 thus point to areas for further research; and we provide tentative evidence to suggest that maternal

372 survival could have acted as a selection pressure for the persistence of hypercoagulability alleles in
373 modern Europeans. Large population studies or commercial providers with extensive genotype and
374 phenotype data such as ALSPAC (Fraser et al. 2013), the UKBiobank (Sudlow et al. 2015) or 23andme
375 (Day et al. 2018) may be productive in examining the relationship between hypercoagulability alleles
376 and birth outcomes in women with linked phenotype and genotype data, or studies specifically
377 addressing the genetics of blood related traits, such as INTERVAL (Di Angelantonio et al. 2017).

378 **Conclusion**

379 These findings add depth to our understanding of the impact of Neanderthal admixture on modern
380 human health and highlight the under-appreciated importance of the risks (and thus selective
381 pressures) associated with childbirth in human evolution. Understanding the selective pressures that
382 may have resulted in the persistence of rs3917862 in modern humans can help improve modern
383 healthcare, and draw further attention to haemorrhaging as a still-significant risk for people in Africa
384 and Asia, while hypertension and venous thromboembolism are a risk for women world-wide.

385 **Quick summary**

386 We hypothesise that Neanderthal *SELP* SNP rs3917862 is more likely to have been maintained in
387 human populations to prevent maternal death due to peri-partum blood loss, than to prevent blood
388 loss due to trauma and interpersonal violence, and present statistical models of the association
389 between the SNP and these causes of death.

390 **Acknowledgments**

391 The authors would like to thank Robert Attenborough for help with sources of hunter-gatherer data,
392 Katherine Boyle for helpful discussion on Palaeolithic environmental conditions, and Enrico Crema
393 for advice on data collection.

394 **Declaration of interest statement**

395 The authors declare no relevant conflicts of interest.

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609 Table 1: percentages of deaths attributable to childbirth (female only), violence and accident in five
610 hunter-gatherer populations

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Population	Continent	Percentage of female deaths in childbirth	Percentage of male deaths in violence	Percentage of female deaths in violence	Percentage of male deaths due to accident	Percentage of female deaths due to accident
San Idefonso Agta	Asia	9.9%	6.7%	0.6%	6.1%	0%
Hiwi	South America	7.9%	32.2%	28.6%	8.9%	11.1%
Hadza	Africa	14.7%	7.3%	2.9%	2.4%	0%
Ache	South America	1.8%	51.8%	60.5%	17.3%	7.4%
Aka	Africa	4.8%	0.5%	0%	6.5%	3.8%

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Table 2: The homicide rates of fifteen hunter-gatherer groups from five continents

Population	Continent	Homicide rate, per 100,000, per year
Hadza	Africa	6.6
Andamanese	Asia	20
Ju/'hoansi	Africa	42
San Ildefonso Agta	Asia	129
Gidjingali	Australia	148
Tiwi	Australia	160
Yaghan	South America	169
Yurok	North America	240
Casiguran Agta	Asia	326
Yolngu	Australia	330
Modoc	North America	450
Ache	South America	500
Hiwi	South America	1018
Piegan	North America	1000
Batek	Asia	1

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Table 3: Maternal mortality ratios from four Tanzanian sources: the Hadza hunter-gatherer population, the national average for Tanzania, and data for two hospitals in Tanzania.

Population	MMR (per 100,000 Live Births)
Hadza	1022
Tanzanian National Average	770
Haydom Lutheran Hospital	382
Kilimanjaro Christian Medical Centre	329

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623 Table 4: Results of Kendall's Tau, testing the correlation between MMR
 624 and SBA, GDP per capita and presence/absence of the Neanderthal
 625 allele, and deaths due to interpersonal violence and SBA, GDP per capita
 626 and presence/absence of the Neanderthal allele.
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	SBA		GDP per capita		Presence/absence of the Neanderthal allele	
	Tau	p-value	Tau	p-value	Tau	p-value
MMR	-0.6517	<0.001	-0.6433	<0.001	-0.5124	<0.001
IVMR	-0.1653	0.0018	-0.2262	<0.001	-0.0901	0.1342

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Table 5: Results of partial correlation tests, determining the Tau values and p-values for each pair of variables, controlling for each other variable in turn.

Variables tested	Tau	P-value
MMR & SBA controlling for GDP	-0.4303	<0.001
MMR & SBA controlling for allele	-0.5198	<0.001
MMR & allele controlling for SBA	-0.2529	<0.001
MMR & allele controlling for GDP	-0.3195	<0.001
MMR & GDP controlling for allele	-0.5312	<0.001
MMR & GDP controlling for SBA	-0.4056	<0.001
IVMR & SBA controlling for GDP	-0.0270	0.5768
IVMR & SBA controlling for allele	-0.1376	0.0057
IVMR & allele controlling for SBA	0.0063	0.8991
IVMR & allele controlling for GDP	0.0279	0.5746
IVMR & GDP controlling for allele	-0.2135	<0.001
IVMR & GDP controlling for SBA	-0.1648	<0.001

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635 Table 6: Regression coefficients and p-values for all possible models of simple and multiple
 636 regression equations for MMR

Explanatory Variable(s)	Regression coefficient							
	b0	p-value	b1	p-value	b2	p-value	b3	p-value
SBA	898.25	<0.001	-859.44	<0.001				
allele	343.32	<0.001	-280.31	<0.001				
GDP	209.74	<0.001	-0.0499	<0.001				
log(GDP)	1021.59	<0.001	-102.60	<0.001				
SBA+allele	829.05	<0.001	-674.11	<0.001	-131.57	<0.001		
SBA+allele+GDP	813.50	<0.001	-648.50	<0.001	-125.30	<0.001	-93420	0.0743
SBA*allele	823.54	<0.001	-666.47	<0.001	-117.20	0.198	-17.05	0.871
SBA+log(GDP)	1071.31	<0.001	-598.82	<0.001	-46.92	<0.001		
SBA+allele+log(GDP)	970.51	<0.001	-522.00	<0.001	-104.37	<0.001	-34.47	<0.001

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Table 7: Adjusted R^2 values and AICc values for the regression models

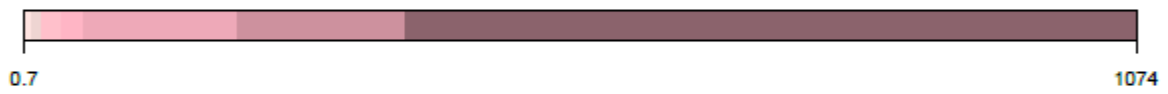
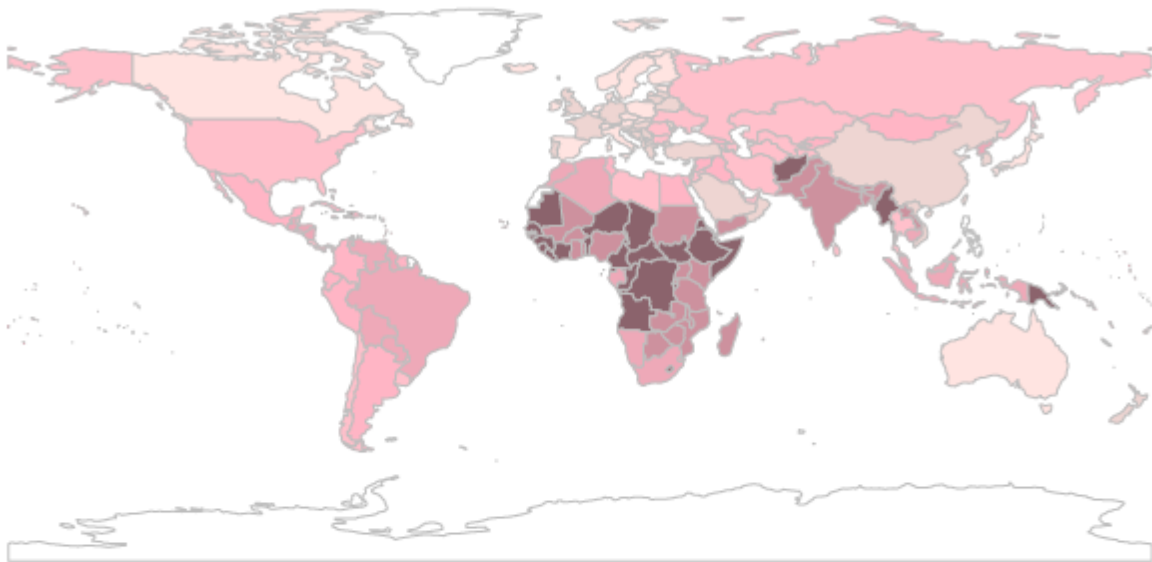
Model	Adjusted R^2 value	AICc	Δ AICc	Rank
MMR~SBA+allele+GDP	0.7054	2249.415	14.461	2
MMR~SBA+allele	0.7018	2250.566	15.612	3
MMR~SBA*allele	0.7002	2252.653	17.699	4
MMR~SBA	0.6373	2285.545	50.591	6
MMR~allele	0.4283	2369.253	134.299	8
MMR~GDP	0.1771	2436.276	201.322	9
MMR~log(GDP)	0.54	2329.247	94.293	7
MMR~SBA+log(GDP)	0.6913	2256.965	22.011	5
MMR~SBA+allele+log(GDP)	0.7277	2234.954	0	1

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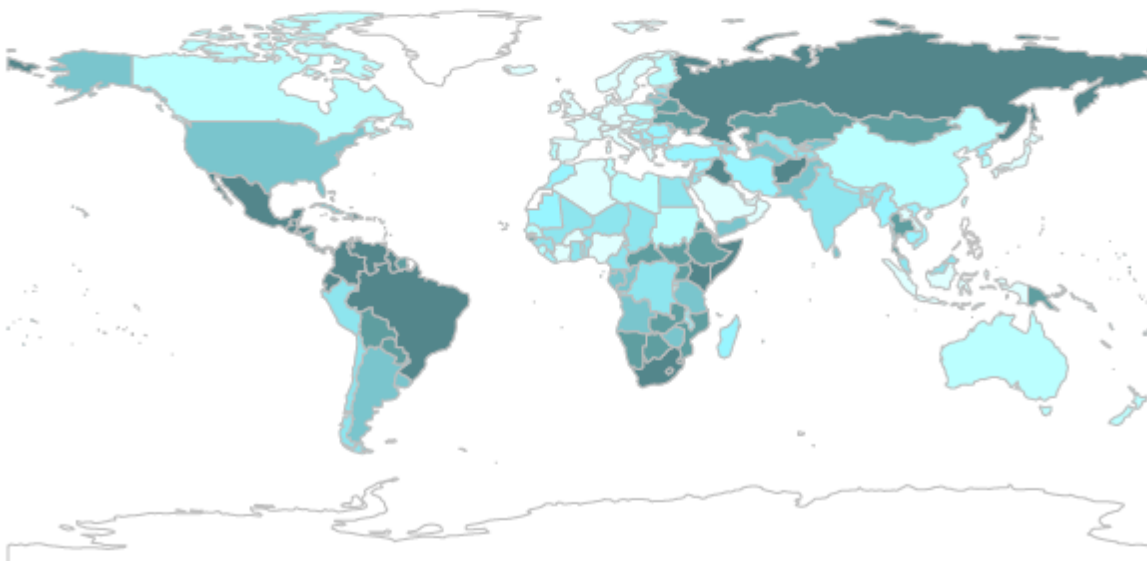
641 **Figures**

642 **Figure 1**

Maternal Mortality Ratio (Number of Deaths per 100,000 Live Births)

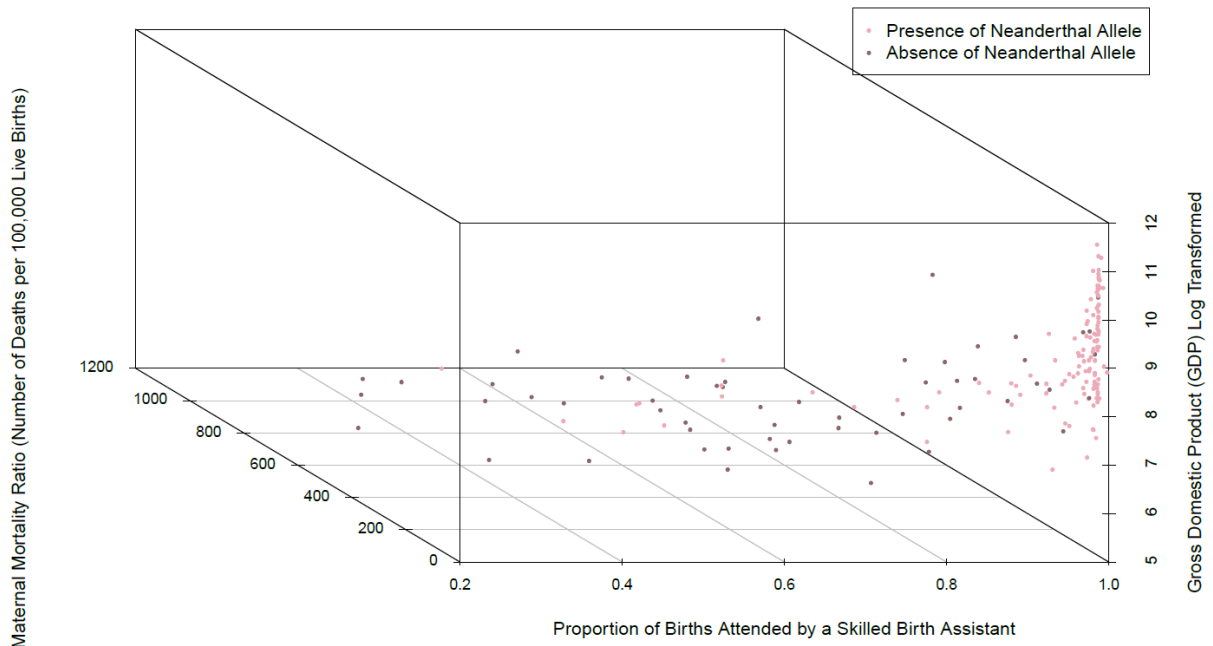


Rate of Deaths due to Interpersonal Violence (per 100,000)



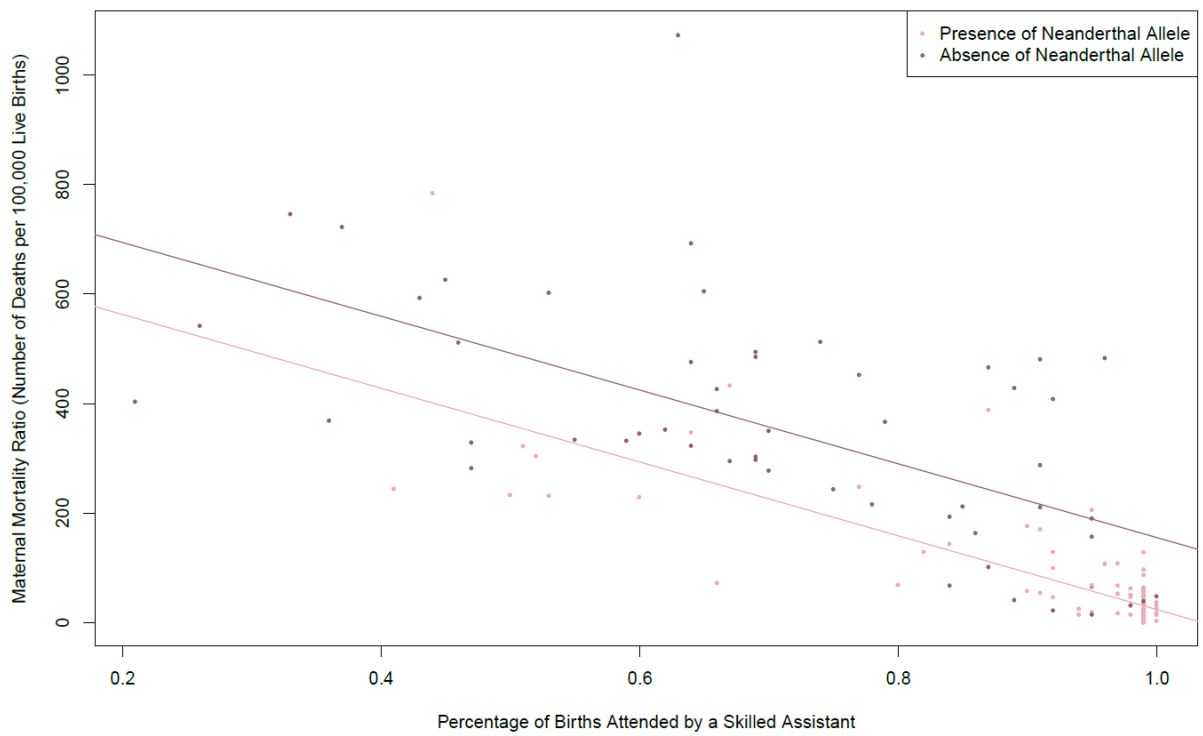
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644 Figure 2



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646 Figure 3



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650 Figure 1: World maps showing variation in maternal mortality ratios in 184 countries (top panel) and
651 variation in deaths due to interpersonal violence for the same 184 countries (lower panel). Data is
652 taken from the GBD 2015 SDG Collaborators (2016).

653 Figure 2: 3D scatter plot showing the best model for prediction of MMR variation. MMR is generally
654 lower, and SBA generally higher for populations with the Neanderthal SNP as compared to those
655 without, while GDP has little visible correlation with these variables.

656 Figure 3: Scatter plot showing the relationship between the percentage of births attended
657 by a skilled birth assistant and maternal mortality ratios. The central tendency of
658 populations with and without Neanderthal allele rs3917862 are shown. Countries with the
659 Neanderthal variant are statistically significantly different in the central tendency of the
660 distribution of MMR values.

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