

1 The face of Ebola: changing frequency of haemorrhage in the  
2 West African compared with Eastern-Central African outbreaks. A  
3 meta-analysis

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20

## 21 Abstract

22 **Background:** The West-African (WA) Zaire Ebolavirus disease (EVD) outbreak was characterized by  
23 an exceptionally high number of cases and deaths as compared with the Eastern-Central African  
24 (ECA) outbreaks. Despite the Zaire Ebolavirus being the most lethal for humans, case-fatality rate,  
25 close to 80% in ECA outbreaks, almost halved to 47% in Guinea-Liberia-Sierra Leone (WA). Such an  
26 improvement was due to the remarkable implementation of international humanitarian aids.  
27 Some studies also suggested that the long human-to-human transmission cycle occurred in WA,  
28 gave rise to human adaptation and consequent immune escape. Haemorrhage, the main feature  
29 in seriously infected EVD patients, is due to the immune system that triggers the infected  
30 endothelial cells which expose the spike-like glycoprotein (GP) of the virion on their surface. If the  
31 human adaptation hypothesis holds true, the proportion of EVD patients with haemorrhage in the  
32 WA outbreak should be lower than in the ECA outbreaks due to immune escape. Therefore, the  
33 aim of this meta-analysis was to compare the relative frequencies of three typical haemorrhagic  
34 symptoms (conjunctival –CB, nasal –NB, gingival –GB- bleedings) in the ECA and WA outbreaks.

35 **Methods:** Literature searches were performed through PubMed and Scopus using generic  
36 keywords; surveys including at least ten patients reporting CB, NB, GB relative frequencies were  
37 extracted and split into ECA and WA. The meta-analytical methods chosen were based on the  
38 levels of between-study heterogeneity and publication bias. Pooled CB, NB, GB relative  
39 frequencies in ECA and WA were estimated and compared. Subgroup analysis including only  
40 studies on Zaire Ebolavirus also was performed. **Results.** Fifteen studies (10 ECA, 5 WA) were  
41 located with 4,867 (CB), 3,859 (NB), 4,278 (GB) EVD patients overall. GB pooled relative frequency  
42 was 45.3% (95% confidence interval -95CI, 34.7-56.1%) and 18.0% (95CI, 6.0-34.5%), in ECA and  
43 WA; NB was 10.6% (95CI, 5.7-16.8%) and 1.3% (1.0-1.8%); GB was 24.2% (95CI, 11.9-39.2%) and  
44 1.9% (95CI, 1.4-2.4%). Subgroup analysis confirmed these results.

45 **Conclusions:** During the WA outbreak the relative frequency of GB decreased by two thirds, while  
46 NB and GB almost disappeared, suggesting that the Zaire Ebolavirus human adaptation hypothesis  
47 is plausible.

48

49 **Keywords:** Ebolavirus disease, Case-fatality rate, Haemorrhage, Conjunctivitis, Epistaxis, Gingival  
50 bleeding

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52

## 53 Background

54

55 Ebola virus disease (EVD) is a zoonosis with persistence of the virus in reservoir species. Indeed, at  
56 least three species of fruit bats (suborder *Megachiroptera*) are naturally infected: *Hypsignathus*  
57 *monstrosus* (hammer-headed fruit bat), *Epomops franqueti* (singing fruit bat) and *Myonycteris*  
58 *torquata* (little collared fruit bat). Asymptomatically-infected bats drop partially eaten fruit and  
59 masticated fruit pulp, contaminated by their saliva and blood, to the ground, where they are eaten  
60 by great apes and forest duikers (antelopes). Ebola virus may then be transmitted from these  
61 infected animals to humans. Humans and primates have been considered the end hosts and have  
62 not been regarded as reservoir species [1]. Ebola virus is transmitted between humans through  
63 direct contact with blood or blood-containing vomit, faeces and other bodily fluids (or tissues)  
64 from EVD patients in the acute stage of the disease. Conversely, subjects who reside in confined,  
65 shared and close spaces, but who have no direct physical contact with acute cases, do not develop  
66 EVD [2-4]. Transmission through air is unlikely [5,6], while sexual transmission through infected  
67 patients is possible [7]. EVD has an incubation period of 2-21 days and the main symptoms are  
68 fever, headache, fatigue, myalgia, diarrhoea, vomiting, abdominal pain and, typically, bleeding  
69 including conjunctival, nasal, gingival bleeding and bleeding from skin and injection sites [8,9].  
70 Recovery may be followed by muscle pains and prolonged virus carriage.

71 The first cases of EVD were reported in 1976 in southern South Sudan and in the northern  
72 Democratic Republic of the Congo [10,11]. Since 1976, human-to-human transmission has been  
73 reported in several Eastern-Central African outbreaks [12]. The latest EVD outbreak started in  
74 Guinea (West Africa), the index case was a 18-month-old boy died in December 2013, but the  
75 outbreak was not detected until March 2014, thus leaving thousands of people becoming infected.  
76 In this largest outbreak ever seen, the case number was 100-1000 times greater than in previous  
77 outbreaks [13]. Assuming that the EVD attack rate did not change significantly from previous  
78 outbreaks [11,14-16] and that in the early stages of the West African outbreak the Ebola virus did  
79 not undergo important genetic changes affecting transmission rate [17-19], such an increase was  
80 probably explained by the exceptional human crowding due to both the urban setting and the  
81 generally higher population density in West Africa than in Eastern-Central Africa [20], which led to  
82 a large increase in the number of exposed individuals. In addition to human crowding there were  
83 other social determinants explaining the West-African outbreak, which, according to the World

84 Health Organization (WHO), were: (1) Extremely damaged Public Health infrastructures in Guinea,  
85 Liberia and Sierra Leone, which only recently emerged from years of civil war; (2) High population  
86 mobility across uncontrolled borders, another consequence of war; (3) Severe shortages of trained  
87 healthcare workers: before the 2013-2015 outbreak there were 1-2 doctors every 100,000  
88 individuals; (4) Cultural beliefs and behavioral practices, such as the adherence to ancestral  
89 funeral and burial practices. The WHO has estimated that 60-80% EVD cases were attributable to  
90 these practices; (5) Reliance on traditional healers: as the outbreak began, the high EVD mortality  
91 in healthcare facilities inducing local people to think that hospitals were the places where  
92 contagion and death actually occur; (6) Community resistance: many people refused to believe  
93 that EVD was real because they and their ancestors had been living in the same environment and  
94 had never developed EVD before; (7) Public Health messages that were intended to promote  
95 protective behaviors proved to have the opposite effect, because they emphasized that the  
96 disease was extremely serious and there was no vaccine or treatment; (8) Other endemic  
97 infectious diseases, such as cholera and malaria, which can be difficult to distinguish from EVD in  
98 the early stages; (9) Ebolavirus is endemic in West Africa; and (10) Spread by international air  
99 travel [21].

100 Genome surveillance studies revealed that while during the Eastern-Central African  
101 outbreaks the Ebolavirus evolution rate was  $0.5-8 \times 10^{-4}$  nucleotide substitutions per site per year  
102 [22], in West Africa such a rate raised to  $9-20 \times 10^{-4}$  nucleotide substitutions per site per year  
103 [17,22-25] showing an accelerated evolution. In addition, many nonsynonymous nucleotide  
104 substitutions occurred on the surface of the spike-like glycoprotein (GP; which plays an important  
105 role in the virus entry into cells) [26] and promoted immune evasion [27].

106 These data support the hypothesis that after a series of tragic coincidences due to  
107 environmental factors, which promoted the spread of the outbreak, the persistence of the human-  
108 to-human transmission cycle gave raise to genetic changes which caused the generation of several  
109 lineages and to human adaptation [27]. This hypothesis is corroborated by the facts that similar  
110 trends have been observed for other cell receptor binding enzymes, such as the neuraminidase of  
111 the influenza virus subtypes A H1N1 and A H3N2 [28] and that a similar intra-individual evolution  
112 occurs to the envelope glycoprotein of hepatitis C virus after the acute phase of the infection to  
113 escape from the host immune response and establish persistent (i.e., chronic) infection [29,30].

114 Another element in support of this hypothesis is the change in case-fatality rate. Indeed,  
115 Zaire Ebolavirus is the most lethal species with an estimated case-fatality rate of 76%, considerably

116 higher than the Sudan (55%) and the Bundibugyo (37%) species [31]. During the latest Zaire  
117 Ebola virus outbreak the case-fatality rate dropped to 47% (considering only confirmed cases and  
118 deaths in Guinea and Sierra Leone) [13]. Such a decrease in case-fatality rate is typical of human  
119 adaptation [28] and is in line with the theory that the majority of human pathogens arose only  
120 after the advent of agriculture and originate from animal pathogens: they were initially  
121 responsible for zoonoses and later adapted to the human hosts, due to co-habitation between  
122 animals and humans and to human crowding [32]. The reported decrease in EVD case-fatality rate  
123 during the West-African EVD outbreak is, however, principally due to the prompt implementation  
124 of control and therapeutic measures by local and international organizations from all over the  
125 world (see, [https://en.wikipedia.org/wiki/Responses\\_to\\_the\\_Ebola\\_virus\\_epidemic\\_in\\_West\\_Africa](https://en.wikipedia.org/wiki/Responses_to_the_Ebola_virus_epidemic_in_West_Africa) for  
126 review. Accessed August 24, 2015).

128 Haemorrhage is a serious EVD consequence, often responsible for patients' demise.  
129 Actually, during the first EVD outbreak in South Sudan, almost all (91%) of the fatal cases and only  
130 one half (48%) of the nonfatal cases showed some visible bleeding features [10]. This link between  
131 the onset of haemorrhage and death is due to the fact that Ebola virus primarily infects endothelial  
132 cells. GP is responsible for virus internalization and is then displayed on the external surface of the  
133 infected cells. Haemorrhage is, therefore, due the lysis of infected endothelial cells caused by the  
134 human immune system triggered by GP [33].

135 Therefore, given the association between death and haemorrhage in EVD patients, in order  
136 to evaluate whether the large decrease in case-fatality rate observed during the West African  
137 outbreak was exclusively due to humanitarian aids or could be explained by Ebola virus human  
138 adaptation, the current study sought to investigate the relative frequency of EVD patients with  
139 haemorrhagic features. More specifically, if the immune evasion hypothesis holds true, patients  
140 with haemorrhage should be less frequent during the West-African outbreak than during the  
141 Eastern-Central African outbreaks. If, alternatively, Ebola virus adaptation to humans did not occur  
142 in West Africa, the frequency of EVD patients with haemorrhage did not vary. Thus, this meta-  
143 analysis sought to estimate the pooled relative frequencies of conjunctival and orofacial (nasal,  
144 gingival) haemorrhagic symptoms observed during the earlier Eastern-Central African and the  
145 latest West African outbreaks.

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147

## 148 Methods

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150 A meta-analysis was performed to estimate the pooled relative frequency of the three main  
151 orofacial haemorrhagic symptoms/signs, that is, conjunctival bleeding/injection (and  
152 conjunctivitis), epistaxis and gingival bleeding, features chosen because they are easily detectable,  
153 although they are not exclusive of EVD. It was assumed that the other conditions and diseases  
154 associated with similar symptoms (e.g. gingivitis) did not change in frequency within the study  
155 populations throughout these years and, therefore, did not interfere with the present analysis.

156 Literature searches were performed through Scopus and PubMed without time and  
157 language restrictions. Inclusion criteria were surveys reporting conjunctival, nasal and/or gingival  
158 bleeding in samples of at least ten EVD patients. Exclusion criteria were all non-relevant studies  
159 (e.g., studies on animals, laboratory studies, reviews, etc.), case reports/samples of less than ten  
160 EVD patients, surveys which did not report bleeding features analytically.

161 Generic key words were used to minimize selection bias which, in meta-analyses, refers to  
162 undetected published studies. Selection bias was considered more pervasive than publication bias,  
163 which refers to unpublished studies, due to the recent explosion of scientific journals and  
164 publications [34]. Thus, key words used were “Ebola” (title, abstract, key words) AND “symptom”  
165 (all fields).

166 Titles and abstracts were used to exclude non-relevant studies. Full texts of remaining  
167 surveys were searched and read and only surveys which met the inclusion criteria were selected.  
168 The reviewers independently extracted conjunctival bleeding (EMV), epistaxis (LTM) and gingival  
169 bleeding (GAM) relative frequencies. Extracted data were then supervised and meta-analyzed by  
170 the two other authors (SP and CS).

171 The pooled relative frequencies of the three features were assessed. The meta-analytic  
172 method was chosen on the basis of the level of between-study heterogeneity, which was  
173 investigated with the  $I^2$  statistic [35]. For non-significant  $I^2$  values the meta-analysis was performed  
174 with the fixed-effects method, for significant  $I^2$  values the random-effects method was used.

175 Publication bias was formally investigated [36] with the test of Egger and colleagues [26], if  
176 the test was significant the number of studies was adjusted for publication bias using the trim and  
177 fill method and identifying the potentially missing studies with the  $R_0$  method [38,39].

178 Sensitivity analysis to study inclusion was made to investigate whether the pooled relative  
179 frequency estimates were influenced by a single study.

180 Surveys were then split according to the country where they have been performed into  
181 Eastern-Central African and West African outbreaks. Pooled relative frequency estimates of  
182 conjunctival nasal and gingival bleeding were assessed for these subgroups using the  
183 aforementioned methodology.

184 Studies on conjunctival bleeding were split into studies which reported bleeding or injected  
185 eyes (conjunctival bleeding) and studies which reported more generic conjunctivitis. Pooled  
186 relative frequency estimates were assessed for all studies together, for Eastern-Central African  
187 studies and for West African studies.

188 Subgroup analysis also was performed to investigate whether there were differences  
189 between Zaire Ebolavirus outbreaks. Therefore, since the West-African outbreak was exclusively  
190 due to the Zaire Ebolavirus species, surveys from Eastern-Central Africa reporting only Zaire  
191 Ebolavirus outbreaks were considered and the pooled relative frequency estimates of conjunctival  
192 nasal and gingival bleeding were assessed.

193 This article followed the MOOSE guidelines for reporting meta-analyses of observational  
194 studies [40]. The statistical analyses were performed using StatView 5.0.1 (SAS Institute Inc.,  
195 NC,US). The level of significance was set at 95%.

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## 198 Results

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200 The Scopus and PubMed searches, performed on October 15, 2015, provided 130 and 691 studies,  
201 respectively (**Figure**). Most of the studies did not fall into the inclusion criteria, thus leaving  
202 twenty-one eligible studies. Four of these were excluded because bleeding features were reported  
203 cumulatively [10,41-43], one because the relative frequency was not reported [44], another one  
204 because it was part of a larger sample [45], thus fifteen primary studies were available for meta-  
205 analysis [2,11,46-58]. All studies were performed during Zaire Ebolavirus outbreaks, excluding two  
206 studies performed during Sudan Ebolavirus [2,51] and one study during a Bundibugyo Ebolavirus  
207 [52] (**Table 1**) outbreak. Five studies were performed during the West African outbreak [54-58],  
208 the remainder during the Eastern-Central African outbreaks.

209 The number of EVD patients considered for conjunctival bleeding were 4,867 (677 from  
210 Eastern-Central Africa, 4,190 from West Africa), those considered for nasal bleeding were 3,859  
211 (522 from Eastern-Central Africa, 3,337 from West Africa), those considered for gingival bleeding

212 were 4,278 (972 from Eastern-Central Africa, 3,306 from West Africa) (**Appendix 1**). The relative  
213 frequencies of conjunctival nasal and gingival bleeding reported by the primary studies are  
214 displayed in **Table 2**. Conjunctival bleeding was reported by thirteen studies, epistaxis by ten  
215 studies, gingival bleeding by eleven studies.

216 Between-study heterogeneity was significantly high for all these features (**Table 3**).  
217 Therefore, the random-effects method was used. According to the test of Egger and colleagues,  
218 the degree of publication bias was not significantly high and no adjustment was performed (data  
219 not in Table). The Forest plots (**Appendix 2**) including all studies show that relative frequency of  
220 conjunctival bleeding apparently decreased from the oldest to the newest study (**Appendix 2a**),  
221 the same trends were observed for epistaxis (**Appendix 2b**) and gingival bleeding (**Appendix 2c**).  
222 The pooled relative frequency estimates showed that conjunctival bleeding/conjunctivitis was the  
223 most frequent feature and was reported in 34% (95% confidence interval, 23-45%) EVD patients,  
224 followed by gingival bleeding (21%; 95% confidence interval, 7-40%) and epistaxis (9%; 95%  
225 confidence interval 3-16%). Conjunctival bleeding/conjunctivitis was significantly more frequent  
226 than epistaxis, while the differences between the other features were not significant (**Table 3**).  
227 Sensitivity analysis to exclusion criteria corroborated these estimates, since none of the studies,  
228 excluded in turn, produced a statistically significant departure from the estimates obtained with all  
229 the primary studies included (**Appendix 3**). The difference between conjunctivitis and conjunctival  
230 bleeding/injection was non-significant: 36% (95% confidence interval, 21-52%) and 31% (95%  
231 confidence interval 21-42%).

232 Studies performed during the Eastern-Central African outbreaks were eight for conjunctival  
233 bleeding [**10,47-53**] and epistaxis [**2,46-48,50-53**], ten for gingival bleeding [**2,10,46-53**], while  
234 studies performed during the West African outbreak were five for conjunctival bleeding [**54-58**],  
235 two for epistaxis [**55,57**] and one for gingival bleeding [**57**]. The random-effects method was used  
236 in almost all cases. The Forest plots including only Eastern-Central African studies were regularly  
237 distributed across the pooled estimates (**Appendix 4**). Fairly irregular distributions were observed  
238 for West African studies, due to their limited number. Conjunctival nasal and gingival bleeding  
239 features were significantly more frequent in the Eastern-Central African studies (**Table 4**).  
240 Conjunctival bleeding pooled relative frequency was 45% in Eastern-Central Africa (95%  
241 confidence interval, 35-56%) and 18% in West Africa (95% confidence interval, 6-34%); epistaxis  
242 was 11% in Eastern-Central Africa (95% confidence interval, 6-17%) and only 1% in West Africa

243 (95% confidence interval, 1-2%); gingival bleeding was 24% in Eastern-Central Africa (95%  
244 confidence interval, 12-39%) and only 2% in West Africa (95% confidence interval, 1-2%).

245 Eastern-Central African studies focusing on Zaire Ebolavirus outbreaks were six for  
246 conjunctival bleeding [10,47-50,53], five for epistaxis [46-48,50,53], seven for gingival bleeding  
247 [10,46-50,53]. The random-effects method was used due to large between-study heterogeneity  
248 (Table 4). No adjustment for publication bias was performed because of regular distribution of the  
249 studies in the Forest plots (data not shown). Pooled relative frequencies of conjunctival, nasal and  
250 gingival bleeding were 45% (95% confidence interval, 33-58%), 9.0% (95% confidence interval, 3-  
251 17%), 28% (95% confidence interval, 12-47%), respectively. Differences with the West African  
252 outbreak were significant at 95% level.

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254

## 255 Discussion

256

257 One problem in assessing the relative frequency of EVD haemorrhagic signs is that features were  
258 reported by healthcare workers when the patients presented at the healthcare facilities and  
259 therefore, depend on the disease stage at presentation (which, in turn, depends on the level of  
260 the population awareness toward EVD) and on the reporting protocol adopted by the healthcare  
261 staff. Although there are no elements to support this hypothesis, it is reasonable to conjecture  
262 that if the general population was unaware of EVD features, infected subjects with epistaxis,  
263 gingival bleeding and/or conjunctival bleeding as main features were less likely to present to  
264 healthcare facilities than were subjects more obviously ill with high fever, vomiting and/or  
265 diarrhoea. Few of the healthcare workers would be trained specifically in ENT/Ophthalmology  
266 and/or Dental examinations, and facilities for these were likely uncommon. Thus, primary studies  
267 must yield some degree of reporting bias, which probably led to underestimate the true relative  
268 frequencies of the these bleeding features. However, if reporting bias was present in the African  
269 healthcare facilities, it was probably lower in the studies used for this meta-analysis, because  
270 authors often followed the WHO EVD case definition recommendations, which comprised gingival  
271 bleeding and conjunctivitis among the key features [59].

272 Another issue with the present meta-analysis was selection bias, which usually refers to  
273 published studies that have not been located. In this analysis it may also refer to the inclusion of  
274 partially duplicate studies, which reported data from the same sets of patients and to the

275 exclusion of studies which were mistakenly considered duplicate and actually were original. This  
276 problem regards principally the West African outbreak, due to the exceptionally high number of  
277 published studies. In order to obtain the most reliable pooled estimates, the risk to exclude  
278 potentially original studies was preferred to the risk to reuse data from the same set of patients.  
279 Indeed, in the first case, the limited number of patients led to broader confidence intervals, while  
280 in the latter case, the inclusion of duplicate data may have raised the burden of the bias of the  
281 primary study, in the event that there was any.

282         The present analysis suggests that conjunctival bleeding is the most frequent EVD orofacial  
283 symptom. However, it is not possible to diagnose EVD solely on the basis of conjunctival bleeding  
284 even if fever, vomiting and diarrhoea are co-existing, because these features are frequently  
285 present in many infectious diseases common in Eastern-Central and West Africa, such as measles  
286 [60] and malaria [61]. Very interestingly, conjunctival bleeding/injection along with conjunctivitis,  
287 which were detected in almost one half EVD patients during the Eastern-Central African  
288 outbreaks, were reported in less than one every six patients during the West African outbreak. As  
289 for the two other investigated features, they were detected relatively frequently during Eastern-  
290 Central African outbreaks, that is, one every four patients (gingival bleeding) and one every ten  
291 patients (epistaxis), while they were almost non-existent during the West African outbreak, being  
292 detectable, at the best, in only one in every forty to fifty EVD patients. These data are  
293 corroborated by another survey which reported that there were only three subjects with  
294 unexplained bleeding in a sample of 103 EVD patients in Sierra Leone [58].

295         Such a huge decrease in ocular and orofacial EVD features, which occurred during the West  
296 African outbreak, parallels the significant decrease in case-fatality rate, which fell by one third,  
297 from 65% (95% confidence interval, 55-75%) during the Eastern-Central African outbreaks [31], to  
298 47.0% (95% confidence interval, 46.2-47.9%) in the West African outbreak –estimated using data  
299 from Guinea and Sierra Leone reported by the WHO [13]. In addition, during the West African  
300 outbreak, the case-fatality rate progressively decreased in rural areas starting from rates similar as  
301 those observed in previous EVD outbreaks. This decrease is in line with the hypothesis of the  
302 adaptation of Ebolavirus to the human host [27], and is not in contrast with the indisputable  
303 excellent results due to the implementation of control policies, such as the national strategy for  
304 the Rapid Isolation and Treatment of Ebola (RITE) in Liberia [62].

305         Another element supporting these results showing that haemorrhagic features drastically  
306 decreased during the West African outbreak, is that in 20% EVD patients the transmission route

307 was unexplained [4]. This suggests that in one fifth of transmissions, the signs and the symptoms  
308 of EVD subjects who acted as infection source were so slight that the infected person could not  
309 remember the transmission event.

310 An element of debate is the meaning of the doubled mutation rate of the Ebolavirus  
311 occurred during the West African outbreak. According to two studies performed on whole Zaire  
312 Ebolavirus genome and on GP glycoprotein sequence, there were no signs of positive selection,  
313 which would have definitively confirmed the human adaptation hypothesis [18,19]. Conversely,  
314 most studies considered as evident signs of selective pressure events such as the emergence of  
315 multiple, even intra-host, novel Ebolavirus lineages during the West African outbreak, the many  
316 non-synonymous RNA nucleotide changes leading to amino-acid changes, some of them at  
317 positions of high level of conservation across Eastern-Central African Ebolaviruses and the positive  
318 selection within the GP glycoprotein amino-acidic sequence, which was not reported during  
319 previous outbreaks [17,23-25,30,63].

320 These data suggest that, although the above-mentioned social determinants have been  
321 responsible for part of the reduced case-fatality reported during the West-African EVD outbreak,  
322 human adaptation, along with immune evasion, may have occurred. Indeed, in the classification of  
323 stages leading to endemic human pathogens from animal pathogens, Ebolavirus was previously  
324 classified at stage 3, that is, an animal pathogen that can undergo only a few cycles of secondary  
325 transmission between humans, so that occasional human outbreaks triggered by a primary  
326 infection soon die out. However, the West African outbreak yielded the characteristics of a stage-4  
327 pathogen, that is, an infectious disease that exists in animals and that infects humans cyclically by  
328 primary transmission from animal host, but that also undergoes long sequences of secondary  
329 transmission between humans, without the involvement of animal hosts. Examples of stage-4  
330 infectious diseases are typhus, yellow fever and influenza A [32]. According to this theory of the  
331 animal origin of human pathogens, transitions from stage 3 to stage 4 frequently occurred during  
332 human evolution. However, the major barrier of such a transition is that animal pathogens need  
333 long secondary transmission chains to evolve adaptation to the novel host. This condition was  
334 prevented in the case of EVD by the small size of human communities where the outbreaks  
335 occurred. Indeed, Eastern/Central African outbreaks took place in villages where the number of  
336 susceptible hosts rapidly ended up, because susceptible individuals died or became immune, as  
337 demonstrated by the high case-fatality rate –as high as 80-90%– and the high immunization rate –  
338 as high as 30%, in villages where EVD outbreaks occurred [2]. However, according to this theory,

339 some external factors, such as frequent blood exposures, presence of susceptible pools of  
340 immunosuppressed hosts (e.g., malnourished populations) and large population sizes, may  
341 intervene to protract human-to-human transmission chains, thus promoting the transition from  
342 stage 3 to stage 4 [32].

343 In line with this theory and accounting for the reported decreased proportion of EVD  
344 patients with haemorrhagic symptoms, it is possible to conjecture that the aforementioned West  
345 Africa-specific social determinants have been responsible for the initially high number of  
346 secondary (and undetected) transmissions between humans [21]. Thanks to this long human-to-  
347 human transmission cycle [10] the Ebolavirus had the opportunity to adapt to the human host.

348

349

## 350 Conclusions

351

352 In conclusion, along with the increased Ebolavirus mutation rate and the decreased case-fatality  
353 rate that occurred during the West African outbreak, this outbreak was also characterized by a  
354 significant decrease in the proportion of patients with bleeding features. Some of these features,  
355 that is, nasal and gingival bleeding almost disappeared. These data support the hypothesis of  
356 Ebolavirus adaptation to human host.

357

358

## 359 Abbreviations

360 EVD, Ebolavirus disease.

361

## 362 Competing Interests

363 The authors declare that they have no competing interests.

364

## 365 Authors' contribution

366 SP designed the study, performed the literature search, verified the extracted data from the  
367 primary studies and reconciled discrepancies between reviewers, performed and commented the  
368 meta-analysis, drafted, wrote and edited the manuscript. GAM performed the literature search,  
369 extracted the data regarding gingival bleeding, commented the results. EMV performed the  
370 literature search, extracted the data regarding conjunctival bleeding, commented the results. LTM

371 performed the literature search, extracted the data regarding nasal bleeding, commented the  
 372 results. CS verified the extracted data from the primary studies and reconciled discrepancies  
 373 between reviewers, commented the meta-analysis, wrote and edited the paper. All the Authors  
 374 read and approved the final version of the paper.

375

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378

#### 379 **References**

- 380 1. Gonzalez JP, Pourrut X, Le roy E. Ebolavirus and other filoviruses. In: Childs JE,  
 381 Mackenzie JS, Richt JA, editors. *Wildlife and Emerging Zoonotic Diseases: The  
 382 Biology, Circumstances and Consequences of Cross-Species Transmission.*  
 383 Heidelberg: Springer; 2007. p. 363-88.
- 384 2. Baron RC, McCormick JB, Zubeir OA. Ebola virus disease in southern Sudan: hospital  
 385 dissemination and intrafamilial spread. *Bull World Health Organ.* 1983;61(6):997-  
 386 1003.
- 387 3. Dowell SF, Mukunu R, Ksiazek TG, Khan AS, Rollin PE, Peters CJ. Transmission of  
 388 Ebola hemorrhagic fever: a study of risk factors in family members, Kikwit,  
 389 Democratic Republic of the Congo, 1995. *Commission de Lutte contre les Epidémies  
 390 à Kikwit. J Infect Dis.* 1999;179(Suppl 1):S87-91.
- 391 4. Faye O, Boëlle PY, Heleze E, Faye O, Loucoubar C, Magassouba N, et al. Chains of  
 392 transmission and control of Ebola virus disease in Conakry, Guinea, in 2014: an  
 393 observational study. *Lancet Infect Dis.* 2015;15(3):320-6.
- 394 5. Jaax N, Jahrling P, Geisbert T et al. Transmission of Ebola virus (Zaire strain) to  
 395 uninfected control monkeys in a biocontainment laboratory. *Lancet.*  
 396 1995;346(8991-8992):1669-71.
- 397 6. Alimonti J, Leung A, Jones S, Gren J, Qiu X, Fernando L, et al. Evaluation of  
 398 transmission risks associated with in vivo replication of several high containment  
 399 pathogens in a biosafety level 4 laboratory. *Sci Rep.* 2014;4,:5824.
- 400 7. Christie A, Davies-Wayne GJ, Cordier-Lasalle T, Blackley DJ, Laney AS, Williams DE,  
 401 et al. Possible sexual transmission of Ebola virus - Liberia, 2015. *MMWR Morb  
 402 Mortal Wkly Rep.* 2015;64(17):479-81.

- 403 8. Samaranayake L, Scully C, Nair RG, Petti S. Viral haemorrhagic fevers with emphasis  
404 on Ebola virus disease and oro-dental healthcare. *Oral Dis.* 2015;21(1):1-6.
- 405 9. Vingolo EM, Messano GA, Fragiotta S, Spadea L, Petti S. Ocular Manifestations of  
406 Ebola Virus Disease: An Ophthalmologist's Guide to Prevent Infection and Panic.  
407 *Biomed Res Int.* 2015:487073.
- 408 10. World Health Organization. Ebola haemorrhagic fever in Sudan, 1976. Report of a  
409 WHO/International Study Team. *Bull World Health Organ.* 1978;56:247-70.
- 410 11. World Health Organization. Ebola haemorrhagic fever in Zaire, 1976. *Bull World*  
411 *Health Organ.* 1978;56(2):271-93.
- 412 12. Centers for Disease Control and Prevention. Outbreaks Chronology: Ebola  
413 Hemorrhagic Fever. 2014 (last updated April 7, 2015).  
414 <http://www.cdc.gov/vhf/ebola/outbreaks/history/chronology.html#five>. Accessed  
415 6 May 2015.
- 416 13. World Health Organization. Ebola Situation Report – 28 October 2015.  
417 [http://apps.who.int/iris/bitstream/10665/191299/1/ebolaitrep\\_28Oct2015\\_eng.pdf?ua=1](http://apps.who.int/iris/bitstream/10665/191299/1/ebolaitrep_28Oct2015_eng.pdf?ua=1)  
418 . Accessed 30 October 2015.
- 419 14. Okware SI, Omaswa FG, Zaramba S, Opio A, Lutwama JJ, Kamugisha J, et al. An  
420 outbreak of Ebola in Uganda. *Trop Med Int Health.* 2002;7(12):1068-75.
- 421 15. Shuaib F, Gunnala R, Musa EO, Mahoney FJ, Oguntimehin O, Nguku PM, et al. Ebola  
422 virus disease outbreak - Nigeria, July-September 2014. *MMWR Morb Mortal Wkly*  
423 *Rep.* 2014;63(39):867-72.
- 424 16. Nyenswah T, Fallah M, Sieh S, Kollie K, Badio M, Gray A, et al. Controlling the last  
425 known cluster of Ebola virus disease - Liberia, January-February 2015. *MMWR Morb*  
426 *Mortal Wkly Rep.* 2015;64(18):500-4.
- 427 17. Gire SK, Goba A, Andersen KG, Sealfon RS, Park DJ, Kanneh L, et al. Genomic  
428 surveillance elucidates Ebola virus origin and transmission during the 2014  
429 outbreak. *Science.* 2014;345(6202):1369-72.
- 430 18. Olabode AS, Jiang X, Robertson DL, Lovell SC. Ebolavirus is evolving but not  
431 changing: No evidence for functional change in EBOV from 1976 to the 2014  
432 outbreak. *Virology.* 2015;482:202-7.

- 433 19. Giovanetti M, Grifoni A, Lo Presti A, Cella E, Montesano C, Zehender G, et al. Amino  
434 acid mutations in Ebola virus glycoprotein of the 2014 epidemic. *J Med Virol.*  
435 2015;87(6):893-8.
- 436 20. Towers S, Patterson-Lomba O, Castillo-Chavez C. Temporal Variations in the  
437 Effective Reproduction Number of the 2014 West Africa Ebola Outbreak. *PLOS*  
438 *Currents Outbreaks.* 2014 Sep 18 . Edition 1. doi:  
439 10.1371/currents.outbreaks.9e4c4294ec8ce1adad283172b16bc908.
- 440 21. World Health Organization. One year into the Ebola epidemic: a deadly, tenacious  
441 and unforgiving virus. January 2015. [http://www.who.int/csr/disease/ebola/one-](http://www.who.int/csr/disease/ebola/one-year-report/ebola-report-1-year.pdf?ua=1)  
442 [year-report/ebola-report-1-year.pdf?ua=1](http://www.who.int/csr/disease/ebola/one-year-report/ebola-report-1-year.pdf?ua=1). Accessed April 10, 2015.
- 443 22. Carroll SA, Towner JS, Sealy TK, McMullan LK, Khristova ML, Burt FJ, et al. Molecular  
444 evolution of viruses of the family Filoviridae based on 97 whole-genome sequences.  
445 *J Virol.* 2013;87(5):2608-16.
- 446 23. Carroll MW, Matthews DA, Hiscox JA, Elmore MJ, Pollakis G, Rambaut A, et al.  
447 Temporal and spatial analysis of the 2014-2015 Ebola virus outbreak in West Africa.  
448 *Nature.* 2015;524(7563):97-101.
- 449 24. Park DJ, Dudas G, Wohl S, Goba A, Whitmer SL, Andersen KG, et al. Ebola Virus  
450 Epidemiology, Transmission, and Evolution during Seven Months in Sierra Leone.  
451 *Cell.* 2015;161(7):1516-26.
- 452 25. Simon-Loriere E, Faye O, Faye O, Koivogui L, Magassouba N, Keita S, Distinct  
453 lineages of Ebola virus in Guinea during the 2014 West African epidemic. *Nature.*  
454 2015;524(7563):102-4.
- 455 26. Nanbo A, Imai M, Watanabe S, Noda T, Takahashi K, Neumann G, et al. Ebolavirus is  
456 internalized into host cells via macropinocytosis in a viral glycoprotein-dependent  
457 manner. *PLoS Pathog.* 2010;6(9):e1001121.
- 458 27. Attar N. Viral evolution: Keeping a watchful eye on Ebola. *Nat Rev Microbiol.*  
459 2015;13(8):457.
- 460 28. Phillips JC. Similarity is not enough: Tipping points of Ebola Zaire mortalities. *Physica*  
461 *A.* 2015;427:277-81.
- 462 29. Booth JC, Kumar U, Webster D, Monjardino J, Thomas HC. Comparison of the rate  
463 of sequence variation in the hypervariable region of E2/NS1 region of hepatitis C

- 464 virus in normal and hypogammaglobulinemic patients. *Hepatology*. 1998;27(1):223-  
465 7.
- 466 30. Liu SQ, Deng CL, Yuan ZM, Rayner S, Zhang B. Identifying the pattern of molecular  
467 evolution for Zaire ebolavirus in the 2014 outbreak in West Africa. *Infect Genet*  
468 *Evol*. 2015;32:51-9.
- 469 31. Lefebvre A, Fiet C, Belpois-Duchamp C, Tiv M, Astruc K, Aho Glélé LS. Case fatality  
470 rates of Ebola virus diseases: a meta-analysis of World Health Organization data.  
471 *Med Mal Infect*. 2014;44(9):412-6.
- 472 32. Wolfe ND, Dunavan CP, Diamond J. Origins of major human infectious diseases.  
473 *Nature*. 2007;447(7142):279-83.
- 474 33. Ansari AA. Clinical features and pathobiology of Ebolavirus infection. *J*  
475 *Autoimmun*. 2014;55:1-9.
- 476 34. Corrêa-Faria P, Petti S. Are overweight/obese children at risk of traumatic dental  
477 injuries? A meta-analysis of observational studies. *Dent Traumatol*. 2015;31(4):274-  
478 82.
- 479 35. Higgins JP, Thompson SG. Quantifying heterogeneity in a meta-analysis. *Stat Med*.  
480 2002;21(11):1539-58.
- 481 36. Petti S. Over two hundred million injuries to anterior teeth attributable to large  
482 overjet: a meta-analysis. *Dent Traumatol*. 2015;31(1):1-8.
- 483 37. Egger M, Davey Smith G, Schneider M, Minder C. Bias in meta-analysis detected by  
484 a simple, graphical test. *BMJ*. 1997;315(7109):629-34.
- 485 38. Duval S, Tweedie R. Trim and fill: a simple funnel-plot-based method of testing and  
486 adjusting for publication bias in meta-analysis. *Biometrics*. 2000;56(2):455-63.
- 487 39. Sutton AJ, Duval SJ, Tweedie RL, Abrams KR, Jones DR. Empirical assessment of  
488 effect of publication bias on meta-analyses. *BMJ*. 2000;320(7249):1574-7.
- 489 40. Stroup DF, Berlin JA, Morton SC, Olkin I, Williamson GD, Rennie D, et al. Meta-  
490 analysis of observational studies in epidemiology: a proposal for reporting. Meta-  
491 analysis of Observational Studies in Epidemiology (MOOSE) group. *JAMA*.  
492 2000;283:2008-12.
- 493 41. World Health Organization. Outbreak of Ebola haemorrhagic fever in Yambio, south  
494 Sudan, April – June 2004 *Wkly Epidemiol Rec*. 2005;80(43):370-5.

- 495 42. MacNeil A, Farnon EC, Wamala J, Okware S, Cannon DL, Reed Z, et al. Proportion of  
496 deaths and clinical features in Bundibugyo Ebola virus infection, Uganda. *Emerg*  
497 *Infect Dis.* 2010;16(12):1969-72.
- 498 43. Barry M, Traoré FA, Sako FB, Kpamy DO, Bah EI, Poncin M, et al. Ebola outbreak in  
499 Conakry, Guinea: epidemiological, clinical, and outcome features. *Med Mal Infect.*  
500 2014;44(11-12):491-4.
- 501 44. Chertow DS, Kleine C, Edwards JK, Scaini R, Giuliani R, Sprecher A. Ebola virus  
502 disease in West Africa--clinical manifestations and management. *N Engl J Med.*  
503 2014;371(22):2054-7.
- 504 45. WHO Ebola Response Team. Ebola virus disease in West Africa--the first 9 months  
505 of the epidemic and forward projections. *N Engl J Med.* 2014;371(16):1481-95.
- 506 46. Sureau PH. Firsthand clinical observations of hemorrhagic manifestations in Ebola  
507 hemorrhagic fever in Zaire. *Rev Infect Dis.* 1989;11(Suppl. 4):S790-3.
- 508 47. Bwaka MA, Bonnet MJ, Calain P, Colebunders R, De Roo A, Guimard Y, et al. Ebola  
509 hemorrhagic fever in Kikwit, Democratic Republic of the Congo: clinical  
510 observations in 103 patients. *J Infect Dis.* 1999;179(Suppl 1):S1-7.
- 511 48. Georges AJ, Leroy EM, Renaut AA, Benissan CT, Nabias RJ, Ngoc MT, et al. Ebola  
512 hemorrhagic fever outbreaks in Gabon, 1994-1997: epidemiologic and health  
513 control issues. *J Infect Dis.* 1999;179(Suppl 1):S65-75.
- 514 49. Khan AS, Tshioko FK, Heymann DL, Le Guenno B, Nabeth P, Kerstiëns B, et al. The  
515 reemergence of Ebola hemorrhagic fever, Democratic Republic of the Congo, 1995.  
516 Commission de Lutte contre les Epidémies à Kikwit. *J Infect Dis.* 1999 ;179(Suppl  
517 1):S76-86.
- 518 50. Ndambi R, Akamituna P, Bonnet MJ, Tukadila AM, Muyembe-Tamfum JJ,  
519 Colebunders R. Epidemiologic and clinical aspects of the Ebola virus epidemic in  
520 Mosango, Democratic Republic of the Congo, 1995. *J Infect Dis.* 1999;179 (Suppl  
521 1):S8-10.
- 522 51. Mupere E, Kaducu OF, Yoti Z. Ebola haemorrhagic fever among hospitalised children  
523 and adolescents in northern Uganda: epidemiologic and clinical observations. *Afr*  
524 *Health Sci.* 2001;1(2):60-5.
- 525 52. Roddy P, Howard N, Van Kerkhove MD, Lutwama J, Wamala J, Yoti Z, et al. Clinical  
526 manifestations and case management of Ebola haemorrhagic fever caused by a

- 527 newly identified virus strain, Bundibugyo, Uganda, 2007-2008. *PLoS One*.  
528 2012;7(12):e52986.
- 529 53. Maganga GD, Kapetshi J, Berthet N, Kebela Ilunga B, Kabange F, Mbala Kingebeni P,  
530 et al. Ebola virus disease in the Democratic Republic of Congo. *N Engl J Med*.  
531 2014;371(22):2083-91.
- 532 54. Schieffelin JS, Shaffer JG, Goba A, Gbakie M, Gire SK, Colubri A, et al. Clinical illness  
533 and outcomes in patients with Ebola in Sierra Leone. *N Engl J Med*.  
534 2014;371(22):2092-100.
- 535 55. Bah EI, Lamah MC, Fletcher T, Jacob ST, Brett-Major DM, Sall AA, et al. Clinical  
536 presentation of patients with Ebola virus disease in Conakry, Guinea. *N Engl J Med*.  
537 2015;372(1):40-7.
- 538 56. Dallatomasina S, Crestani R, Sylvester Squire J, Declerk H, Caleo GM, Wolz A, et al.  
539 Ebola outbreak in rural West Africa: epidemiology, clinical features and outcomes.  
540 *Trop Med Int Health*. 2015;20(4):448-54.
- 541 57. WHO Ebola Response Team, Agua-Agum J, Ariyarajah A, Blake IM, Cori A, Donnelly  
542 CA, et al. Ebola virus disease among children in West Africa. *N Engl J Med*.  
543 2015;372(13):1274-7.
- 544 58. Yan T, Mu J, Qin E, Wang Y, Liu L, Wu D, et al. Clinical characteristics of 154 patients  
545 suspected of having Ebola virus disease in the Ebola holding center of Jui  
546 Government Hospital in Sierra Leone during the 2014 Ebola outbreak. *Eur J Clin  
547 Microbiol Infect Dis*. 2015 Jul 30. doi: 10.1007/s10096-015-2457-z
- 548 59. World Health Organization. Case definition recommendations for Ebola or Marburg  
549 virus diseases. Geneva: WHO; 2014.
- 550 60. Abdalla TM, Karsany MS, Ali AA. Correlation of measles and dengue infection in  
551 Kassala, Eastern Sudan. *J Med Virol*. 2015;87(1):76-8.
- 552 61. Menaca A, Pell C, Manda-Taylor L, Chatio S, Afrah NA, et al. Local illness concepts  
553 and their relevance for the prevention and control of malaria during pregnancy in  
554 Ghana, Kenya and Malawi: findings from a comparative qualitative study. *Malar J*.  
555 2013;12:257.
- 556 62. Kateh F, Nagbe T, Kieta A, Barskey A, Gasasira AN, Driscoll A, et al. Rapid response  
557 to Ebola outbreaks in remote areas - Liberia, July-November 2014. *MMWR Morb  
558 Mortal Wkly Rep*. 2015;64(7):188-92.

559 63. Tong YG, Shi WF, Di Liu, Qian J, Liang L, Bo XC, et al. Genetic diversity and  
560 evolutionary dynamics of Ebola virus in Sierra Leone. Nature 2015; doi:  
561 10.1038/nature14490.  
562  
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565 **Table 1.** Characteristics of the outbreaks described by the primary studies.

566

Study	Ebolavirus	Country	Survey year
WHO, 1978	Zaire	Democratic Republic of Congo	1976
Baron et al., 1983	Sudan	South Sudan	1979
Sureau, 1989	Zaire	Democratic Republic of Congo	1976
Bwaka et al., 1999	Zaire	Democratic Republic of Congo	1995
Georges et al., 1999	Zaire	Gabon	1996
Khan et al., 1999	Zaire	Democratic Republic of Congo	1995
Ndambi et al., 1999	Zaire	Democratic Republic of Congo	1995
Mupere et al., 2001	Sudan	Uganda	2000
Roddy et al., 2012	Bundibugyo	Uganda	2007
Maganga et al., 2014	Zaire	Democratic Republic of Congo	2014
Schieffelin et al., 2014	Zaire	Sierra Leone	2014
Bah et al., 2015	Zaire	Guinea	2014
Dallatomasina et al., 2015	Zaire	Sierra Leone	2014
WHO Ebola Response Team, 2015	Zaire	Guinea, Liberia, Sierra Leone	2014
Yan et al., 2015	Zaire	Sierra Leone	2014

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570 **Table 2.** Relative frequencies (95% confidence intervals between parentheses) of conjunctival,  
 571 nasal and gingival bleeding features reported by the primary studies.

572

Study	Conjunctival bleeding and conjunctivitis	Nasal bleeding	Gingival bleeding
WHO, 1978	55.0% (48.5-61.3%)		67.4% (61.4-73.3%)
Baron et al., 1983		21.9% (7.6-36.2%)	34.4% (17.9-50.8%)
Sureau, 1989		14.7% (10.5-19.0%)	22.3% (17.3-27.3%)
Bwaka et al., 1999	42.7% (33.0-52.8%)	1.9% (0.0-4.6%)	12.6% (6.2-19.0%)
Georges et al., 1999	53.3% (26.6-78.7%)	13.3% (0.0-30.5%)	40.0% (15.2-64.8%)
Khan et al., 1999	35.7% (29.2-42.6%)		22.1% (16.5-27.8%)
Ndambi et al., 1999	78.3% (56.3-92.5%)	4.3% (0.0-12.7%)	30.4% (11.6-49.2%)
Mupere et al., 2001	40.0% (19.1-63.9%)	10.0% (0.0-23.1%)	10.0% (0.0-23.1%)
Roddy et al., 2012	50.0% (29.9-70.1%)	7.7% (0.0-17.9%)	3.8% (0.0-11.2%)
Maganga et al., 2014	15.8% (6.0-31.3%)	10.5% (0.8-20.3%)	7.9% (0.0-16.5%)
Schieffelin et al., 2014	25.0% (13.2-40.3%)		
Bah et al., 2015	10.8% (3.0-25.4%)	5.4% (0.0-12.7%)	
Dallatomasina et al., 2015	2.0% (0.7-4.7%)		
WHO Ebola Response Team, 2015	26.0% (24.6-27.4%)	1.3% (0.9-1.7%)	1.9% (1.4-2.4%)
Yan et al., 2015	34.3% (25.4-44.0%)		

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575

576 **Table 3.** Pooled relative frequency estimates of conjunctival, nasal and gingival bleeding features.577 Between-study heterogeneity ( $I^2$  statistic, 95% confidence interval).

578

Bleeding	Pooled relative frequency	95% confidence interval	$I^2$ statistic
Conjunctival (all)	33.6%	23.5-44.6%	94.6-97.2%
Conjunctivitis only	35.9%	20.9-52.5%	96.4-98.3%
Conjunctival bleeding/injection	31.1%	21.3-41.8%	42.1-90.3%
Nasal	8.7%	3.5-16.0%	87.8-95.1%
Gingival	21.1%	7.0-40.1%	98.4-99.0%

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582 **Table 4.** Pooled relative frequency estimates of conjunctival, nasal and gingival bleeding features  
 583 in the Eastern-Central African outbreaks (all outbreaks, Zaire Ebolavirus outbreaks) and in the  
 584 West-African outbreak. Between-study heterogeneity ( $I^2$  statistic, 95% confidence interval).

585

	Pooled relative frequency	95% confidence interval	$I^2$ statistic
Conjunctival bleeding/injection and conjunctivitis			
Eastern-Central Africa (8 studies)	45.3%	34.7-56.1%	70.7-91.6%
Eastern-Central Africa (Zaire Ebolavirus, 6 studies)	45.4%	32.6-58.4%	77.9-94.2%
West Africa (5 studies)	18.0%	6.0-34.5%	95.6-98.3%
Conjunctival bleeding/injection only			
Eastern-Central Africa (3 studies)	38.2%	33.0-43.6%	0.0-95.5 <sup>a</sup>
West Africa (2 studies)	18.8%	11.0-28.8%	0.0-91.3 <sup>a</sup>
Conjunctivitis only			
Eastern-Central Africa (5 studies)	50.0%	31.0-68.2%	72.0-94.0%
West Africa (3 studies)	17.9%	3.0-41.5%	95.6-99.2%
Nasal bleeding			
Eastern-Central Africa (8 studies)	10.6%	5.7-16.8%	33.1-84.8%
Eastern-Central Africa (Zaire Ebolavirus, 5 studies)	9.0%	3.3-17.2%	47.7-90.9%
West Africa (2 studies)	1.3%	1.0-1.8%	0.0-92.8% <sup>a</sup>
Gingival bleeding			
Eastern-Central Africa (10 studies)	24.2%	11.9-39.2%	93.6-97.0%
Eastern-Central Africa (Zaire Ebolavirus, 7 studies)	27.9%	12.5-46.7%	95.1-97.9%
West Africa (1 study)	1.9%	1.4-2.4%	Not applicable <sup>b</sup>

586 <sup>a</sup> non-significant  $I^2$  statistic, fixed-effects meta-analytic method; <sup>b</sup> prevalence estimate reported by a single study

587 All the differences between pooled relative frequencies in Eastern-Central Africa and West Africa are statistically  
 588 significant at 95% level

589

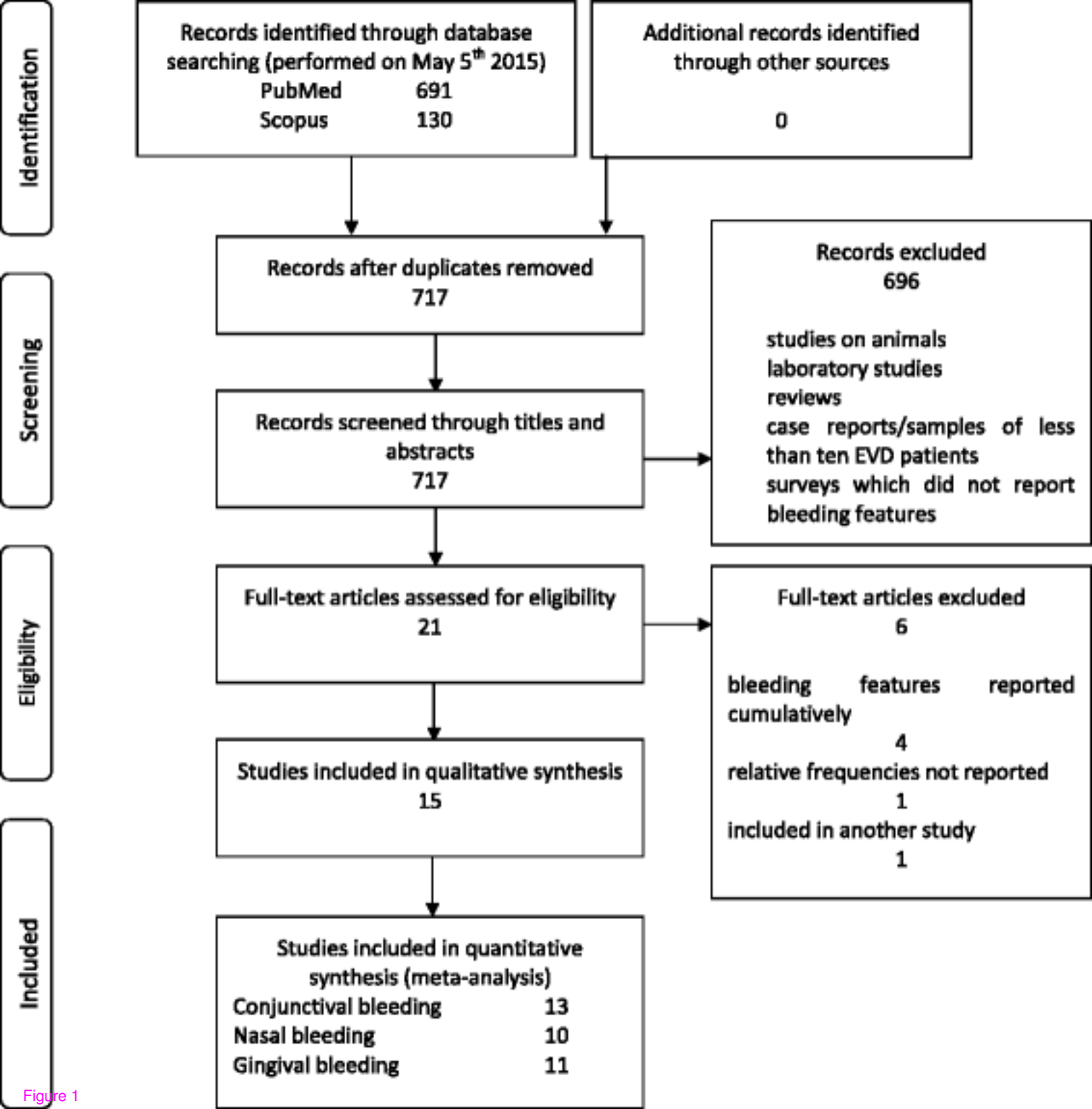


Figure 1

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