Animal contact and paediatric acute febrile illness in Greater Accra Region, Ghana

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SUMMARY

Objective: To examine the association between animal contact (primarily dogs and cats) and non-malarial fever, as well as with secondary symptoms of headache, nausea, vomiting, and cough, in 687 children in Greater Accra Region, Ghana.

Design: Cross-sectional study of acute febrile illness among children aged 1-15 years old between October 2016 and August 2017.

Setting: Ledzokuku-Krowor Municipal Assembly (LEKMA) Hospital, Teshie, Greater Accra Region.

Participants: The study included children with acute fever, defined as a measured temperature of greater than 37.5°C, occurring less than seven days before the hospital visit, and afebrile children as controls.

Main outcome measures: Measured fever, self-reported fever, and secondary symptoms, each adjusting for patient household characteristics.

Results: Animal contact was neither associated with measured fever (OR = 1.04, 95% CI 0.73-1.49) nor with self-reported fever (OR = 0.97, 95% CI 0.68-1.39). Animal contact was associated with headache (OR = 3.26, 95% CI 2.23-4.77, P < .01) and nausea (OR = 3.05, 95% CI 1.99-4.68, P < .01), but not with vomiting or cough. Additional models that used alternate inclusion criteria to define non-malarial fever yielded similar results. Several bacterial zoonoses that could plausibly have been transmitted by dogs and cats were diagnosed in the study population.

Conclusion: These findings suggest the need for future studies to evaluate animal contact as a risk factor for bacterial zoonoses that may serve as an etiological driver of acute febrile illness.

Keywords: fever, acute febrile illness, bacterial zoonosis, pet infections **Funding:** no external funding

INTRODUCTION

Acute febrile illness, a rapid onset of fever occurring for less than 7 days, causes a large burden of disease globally.^{1,2} Febrile illness is among the most common reasons for people in low-resource areas to seek healthcare.^{3,4} In sub-Saharan Africa, fever is a common symptom of many important sources of morbidity and mortality.² Annually, over six million children die due to preventable or treatable illnesses, many of which are febrile, and approximately 50% are in resource-limited settings.⁵

Undifferentiated fever is the main clinical symptom of many diseases of global importance, including malaria, bacterial diseases, bacterial zoonoses and many viral infections.³ But the diagnosis of patients with febrile illness is challenging due to both a lack of specific presentation and limited diagnostics.^{2,6}

Except for malaria diagnostics, laboratory tests for many febrile diseases are often complex, costly, and not widely available in low-resource areas.^{1,3} While malaria remains a major cause of fever in tropical sub-Saharan Africa, the incidence of malaria has been declining since 2003,² and recent studies have demonstrated that a significant proportion of febrile patients suspected of malaria are actually suffering from other infections.^{2,4}

Efforts to better understand the causes of febrile illness have identified bacterial zoonoses as common, under-recognized causes of fever in Africa.^{2,4} In a study of 870 febrile children and adults in Northern Tanzania, bacterial zoonoses were identified among 26.2% of clinic admissions.⁴ Household animals, such as dogs and cats, are known to transmit bacterial zoonoses such as leptospirosis, brucellosis, and Q fever, as well as common infections such as *staphylococci*, rickettsia, and *Streptococcus pneumoniae*.^{2,7-10}

Both pets and feral animals can become vectors of bacterial zoonoses in settings where inadequate water, sanitation, and hygiene (WASH) services expose freeranging animals to waste and pathogens in the environment. Exposed animals may then carry pathogens into human living spaces and create subsequent infection pathways similar to faecal-oral contamination from food, water, and fomites,¹¹ and physical contact between humans and outdoor-living animals. Children's outdoor play spaces may present additional transmission pathways if they overlap with animal activity spaces. These types of animal exposures are typical of the urban ecology of low-income communities with insufficient WASH services¹², particularly concerning rodents¹³, but a few studies have examined the potential of such public health burden in the African context associated with dogs and cats.

Understanding the full range of aetiologies of acute febrile illness has important implications for prevention and treatment. This study examined the association between animal contact (primarily dogs and cats) and fever, as well as secondary symptoms of headache, nausea, vomiting, and cough, in paediatric acute febrile illness patients in Teshie, a large town in the Greater Accra Region, Ghana. Our goal was to assess whether animal contact might be a risk factor for febrile illness in this setting and whether bacterial and other zoonoses warrant additional exploration as a source of undifferentiated fever.

METHODS

Children aged 1-15 years old at the outpatient clinics of the Ledzokuku-Krowor Municipal Assembly (LEKMA) Hospital in the Greater Accra Region in southern Ghana were recruited to a cross-sectional study of acute febrile illness between October 2016 and August 2017. LEKMA was a coastal district between the Accra Metropolitan Area and Tema with a 2010 population of 227,932 $^{\rm 14}$ and anchored by two large towns, Teshie and Nungua (it was split into two municipal districts in 2018). This region has experienced recent economic development, but despite most households having piped water connections, most of the district still lacks reliable WASH services in many areas due to water rationing.15 Household animal ownership-which is typically dogs and cats, but occasionally goats, fowl, or cattle-is similar to the rates observed in other Ga communities in Accra, such as Bukom and Shiabu.11

The study included children with acute fever, defined as a measured temperature of greater than 37.5°C, occurring less than seven days before the hospital visit, and afebrile children as controls. The clinician also recorded whether the fever was self-reported by the child or parent/guardian. Additional inclusion criteria were the parent or guardian's willingness to provide informed consent, with the willingness of the child if over 10 years old, residency in the study area for two months, and parent/guardian consent to allow blood and other samples to be collected. Children showing signs of severe disease, children with a chronic illness, and minors unaccompanied by a parent or guardian were excluded from the study.

Because malaria is not transmitted via household animals, we excluded children diagnosed with malaria from our analysis of animal contact. For robustness, we tested two separate criteria of "malaria-negative": children with a negative parasite test or malaria RDT, and those who did not receive a clinical diagnosis of malaria from the attending physician. As already noted, fever also had two possible definitions: fever as self-reported by participants and temperature measured at the clinic. To ensure that our results were stable across clinical definitions of malaria and fever, we repeated our analysis of animal contact for all four combinations of inclusion criteria and fever diagnosis, from most to least objective: (1) parasite negative (for malaria), measured fever; (2) parasite negative, self-reported fever; (3) clinical negative (for malaria), measured fever; (4) clinical negative, self-reported fever. This robustness was important given how, in some regions of Ghana, traditional words for fever and *malaria* have historically been interchangeable.¹⁶

The study team, comprised of a field worker and a nurse, identified children and measured their temperature, height and weight. The study information and procedures were explained to every parent or guardian with sufficient time to contemplate participation. The study team obtained written informed consent from all parents/guardians. The participants were then sent to the study clinician who physically examined them and enrolled them in the study. A questionnaire was administered to collect socio-demographic and additional clinical data, including signs and symptoms. It included an item assessing whether the child had contact with animals (alive or dead) in the prior two weeks. All children were tested for malaria using microscopy and rapid diagnostic test (RDT). Blood samples from children with negative malaria RDT and microscopy were randomly selected by using a random number generator to select sample IDs to be further analysed using a customised multi-pathogen, real-time PCR-based TagMan probe-array card (TAC; Applied Biosystems, Carlsbad, CA, USA), as described elsewhere.¹⁷ The TaqMan Array Card assessed 26 pathogens, including several bacterial zoonoses (Figure S1, Supplementary Material).

Additional blood and urine cultures were performed at the request of the clinician. Nasopharyngeal samples were col-

lected from participants showing signs of respiratory infection; however, laboratory analyses of these samples were not completed due to resource constraints.

The study team performed bivariate analyses of participant demographic characteristics using chi-square tests for categorical variables and F tests for the difference in means between groups for continuous variables. We fitted multivariable logistic regression models for each of the four combinations of fever and malaria definitions to analyse the association between animal contact and fever, adjusting for covariates. We controlled for several participant characteristics, including gender, parent education, parent occupation, ethnicity, and whether a drug was being taken for the current illness at the time of the visit. We then used chi-square tests to explore bivariate associations between the two definitions of fever and the most common secondary symptoms potentially related to animal-transmitted diseases (headache, nausea, vomiting and cough). We then performed a similar multivariable regression analysis of these symptoms. All analyses were performed using the software R version 3.6.2.

Ethical Approval

The original febrile illness study was approved by the Institutional Review Committee of Kintampo Health Research Centre (approval number: 0004854), Institutional Review Board of Noguchi Memorial Institute for Medical Research at the University of Ghana (study number: 099/15-16), and Ghana Health Service Ethical Review Committee (GHS-ERC number: 12/06/2016). Informed consent was sought from the caregivers of the participants. Confidentiality and anonymity were ensured throughout the process. This study, conducted on anonymised patient data, was deemed nonhuman subjects research by the Institutional Review Board at the University of Miami.

RESULTS

Our analytical sample sizes were reduced from 775 to 687 after excluding 88 children due to parasite-positive malaria, and from 775 to 593 after excluding 182 children receiving a clinical malaria diagnosis. Table 1 summarises patient characteristics according to each definition of fever; 59% self-reported fever, and 54% had a measured fever at the clinic.

Table 1 Descriptive characteristics of select patient and household characteristics showing associations with measured and self-reported fever (n = 687).

	Measured			
	Febrile	Afebrile	Febrile	Afebrile
Overall Rate	368 (54%)	319 (46%)	406 (59%)	281 (41%)
Age (years)	3.45 (2.83)***	5.42 (4.23)	3.47 (2.78)***	5.75 (4.33)
Gender				
Female	169 (46%)	164 (51%)	189 (47%)	144 (51%)
Male	199 (54%)	155 (49%)	217 (53%)	137 (49%)
Yes	103 (28%)	86 (27%)	111 (27%)	78 (28%)
No	265 (72%)	233 (73%)	295 (73%)	203 (72%)
Cat	58 (16%)	53 (17%)	64 (16%)	47 (17%)
Cat & Dog	1 (<1%)	0 (0%)	1 (<1%)	0 (0%)
Dog	41 (11%)	32 (10%)	43 (11%)	30 (11%)
Sheep/Goat	2 (<1%)	1 (<1%)	2 (<1%)	1 (<1%)
Unknown	1 (<1%)	0 (0%)	1 (<1%)	0 (0%)
None	24 (7%)	26 (8%)	26 (6%)	24 (9%)
Primary	137 (37%)	109 (34%)	151(37%)	95 (33%)
Secondary	122 (33%)	111 (35%)	136 (33%)	97 (34%)
Tertiary	85 (23%)	73 (23%)	93 (23%)	65 (23%)
Professional, skilled labour	92(25%)***	116 (36%)	106 (26%)***	102 (36%)
Trader, unskilled labour,	276 (75%)	203 (64%)	300 (74%)	179 (64%)
unemployed				
Ethnicity				
Ga Adangbe	143 (39%)	103 (32%)	153 (38%)	93 (33%)
Akan	119 (32%)	118 (37%)	135 (33%)	102 (36%)
Ewe	71 (19%)	75 (24%)	81 (20%)	65 (23%)
Northern	27 (7%)	19 (6%)	29 (7%)	17 (6%)
Unknown	8 (2%)	4 (1%)	8 (2%)	4 (1%)
None	104(28%)***	160 (50%)	114 (28%)***	150 (53%)
Analgesic	165 (45%)	113 (35%)	188 (46%)	90 (32%)
Other	99 (27%)	46 (14%)	104 (26%)	41 (15%)
Yes	13 (4%)	0 (0%)	13 (3%)	0 (0%)
No	354 (96%)	318 (100%)	391 (96%)	281 (100%)
Pipe borne water	12 (3%)	1 (<1%)	12 (3%)	1 (0.3%)
Sachet water	356 (97%)	318 (99%)	394 (97%)	280 (99%)

* *P* < .05, ** *P* < .01, *** *P* < .001

† Associations not measured due to lack of variation

Regardless of fever definition, parasite-negative, febrile children were significantly younger than afebrile children, more likely to have parents employed in lower-skilled jobs or unemployed, and more likely to have been taking a drug for their illness at the time of visit (all P < .001) (Table 1). We observed the same differences between febrile and afebrile children using the alternate inclusion criteria of being clinically diagnosed as negative for malaria (Table S1, Supplementary Material). Among parasite-negative children, 28% of febrile (measured) and 27% of afebrile children reported contact with animals 14 days prior to the clinic visit, with similar rates observed when defining fever as self-reported. Animal exposures, defined as contact with any animal (alive or dead) in the past two weeks, were most commonly cats and dogs, with two febrile children reporting contact with sheep or goats.

Table S1 (see Supplementary Material) summarizes the characteristics of patients based on the secondary inclusion criteria of no malaria clinical diagnosis. A total of 593 patients were included in this analysis, 59% of which reported fever and 54% had a measured fever at the clinic. Ninety-three children, or 27% of febrile children, and 88 children, or 28% of children with a measured fever, reported contact with animals in the 14 days prior to their clinic visit.

Table 2 summarises the four multivariable logistic regression models for each inclusion criterion. Among children with a parasite-negative malaria test, animal contact was neither associated with self-reported fever (OR = 0.97, 95% CI 0.68-1.39), nor with a measured fever (OR = 1.04, 95% CI 0.73 -1.49), with similar results for children diagnosed as clinically negative for malaria.

Table 2 Multivariable logistic regression models of the association between patient characteristics and fever, using two fever criteria (measured vs. self-reported) and two methods of malaria diagnosis (parasite-negative vs. clinical negative).

	Measured Fever & Parasite Negative	Self-Reported Fever & Parasite Negative	Measured Fever & Clinical Negative	Self-Reported Fever & Clinical Negative
Animal Contact				
No †				
Yes	1.04 (0.73-1.49)	0.97 (0.68-1.39)	0.99 (0.68-1.44)	0.89 (0.61-1.30)
Gender				
Male †				
Female	0.79 (0.58-1.08)	0.82 (0.59-1.13)	0.79 (0.56-1.10)	0.80 (0.57-1.13)
Parent Education				
None †				
Primary	1.64 (0.86-3.17)	1.72 (0.89-3.32)	1.17 (0.59-2.33)	1.39 (0.69-2.76)
Secondary	1.28 (0.67-2.46)	1.36 (0.70-2.61)	0.98 (0.49-1.95)	1.21 (0.61-2.41)
Tertiary	1.54 (0.77-3.13)	1.51 (0.74-3.06)	1.18 (0.56-2.49)	1.37 (0.65-2.87)
Parent Occupation				
Trader, unskilled labour,				
unemployed †				
Professional/skilled la-	0.53 (0.36-0.75)***	0.56 (0.39-0.83)**	0.54 (0.36-0.79)**	0.60 (0.41-0.89)*
bour				
Ethnicity				
Ga Adangbe †				
Akan	0.75 (0.52-1.09)	0.83 (0.57-1.22)	0.76 (0.51-1.14)	0.80 (0.53-1.21)
Ewe	0.75 (0.48-1.15)	0.83 (0.53-1.28)	0.74 (0.47-1.16)	0.78 (0.49-1.24)
Northern	1.04 (0.53-2.06)	1.01 (0.51-2.05)	1.12 (0.53-2.39)	0.91 (0.44-1.95)
Unknown	1.31 (0.39-5.15)	1.05 (0.31-4.14)	1.19 (0.34-4.79)	0.96 (0.27-3.83)
Drug for current illness				
None †				
Analgesic	2.19 (1.54-3.12)***	2.71 (1.90-3.89)***	2.07 (1.42-3.03)***	2.36 (1.62-3.47)***
Other	3.48 (2.25-5.45)***	3.52 (2.26-5.54)***	3.38 (2.12-5.46)***	3.15 (1.97-5.09)***

Odds ratios (OR) presented with 95% confidence intervals (95% CI). * P < .05, ** P < .01, *** P < .001

† Reference category

Having a parent's occupation categorised as professional/skilled labour was significantly associated with measured fever (OR = 0.53, 95% CI 0.36-0.75), as well as taking either an analgesic drug (OR = 2.19, 95% CI 1.54-3.12) or other types of drug (OR = 3.48, 95% CI 2.25-5.45) for the current illness. We observed similar results for all four multivariable models (Table 2).

Across the four columns of Table 2, which represent decreasing objectivity of diagnosis from left to right, two of the three significant measures—parent occupation and using 'other' drug as treatment—had the smallest effects in the least objective combination of self-reported fever and being clinically negative for malaria. Although the differences were modest, this underscores the reason for testing the four different combinations of fever and malaria-negative definitions, given how diagnostic capacity can vary throughout the country.

Table 3 summarises associations between measured fever and reported secondary symptoms. Among the 687 patients with a negative parasite test or RDT, 115 patients reported nausea, 171 headache, 196 vomiting and 274 cough. Measured fever was significantly associated with headache, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and cough, $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^2(1, N = 687) = 8.85$, P < .01 and $\chi^$ N = 687) = 5.48, P = .02, while self-reported fever was significantly associated with headache, $\chi^2(1, N = 687) =$ 5.29, P = .02 and vomiting, $\chi^2(1, N = 687) = 4.74$, p = .04. Nausea was significantly associated with headache, $\chi^2(1, N = 687) = 51.22, P < .001$, and vomiting, $\chi^2(1, N =$ (687) = 213.91, P < .001. Vomiting was significantly associated with cough, $\chi^2(1, N = 687) = 13.54, P < .001$.

Table 3 Chi-square matrix showing associations between symptoms.

	Measured Fever	Self-Reported Fe- ver	dł
Measured Fever		512.20***	- 68
Self-Reported Fever	512.20***		cg
Headache	8.85**	5.29*	Ri
Nausea	0.65	0.65	p
Vomiting	1.45	4.74*	id
Cough	5.48*	2.23	2
* P < .05, $** P < .01$, $*** P$	< .001		

Table 4 summarises the multivariable logistic regression models of headache, nausea, vomiting, and cough with animal contact, again controlling for covariates. There was an association between animal contact and headache for patients with a parasite-negative test result (OR = 3.26, 95% CI 2.23-4.77, P < .01) and for patients without a clinical malaria diagnosis (OR = 2.95, 95% CI 1.96-4.42, *P* < .01). Animal contact was also associated with nausea for patients with a parasite negative test results (OR = 3.05, 95% CI 1.99-4.68, P < .01) and for patients without a clinical diagnosis of malaria (OR = 2.72, 95% CI 1.71-4.32, P < .01). Animal contact was not associated with vomiting or cough.

Blood and urine tests yielded positive culture results for just seven febrile children with a parasite-negative test who reported animal contact (see Table 5). Three children had coagulase-negative staphylococci, and there was one case of Enterococcus spp, HIV, Staphylococcus aureus, and Streptococcus pneumoniae. Adtradathy, among cNaussen why emiting trepoval animal ntact, test results included dengue virus, Escherichia 15 129 group D streptococci, Psejujamonas jugueinosa, ckettsia, Staphylogogeus auneros, and Straptococcus **12.37***töħiae*, underscoring th**2**3**r4***t***5***t***67**.02**n**fections efflified in the ori<u>213</u>**9***t*^{***} transformed to the transformed parent study, ^{18,19}**13.54*****

Table 4 Multivariable logistic regression models of the association between patient characteristics and headache, nausea, vomiting and cough, using two inclusion criteria

	Headache (Par- asite Negative)	Headache (Clinical Nega- tive)	Nausea (Para- site Negative)	Nausea (Clinical Nega- tive)	Vomit (Parasite Neg- ative)	Vomit (Clinical Nega- tive	Cough (Parasite Negative)	Cough (Clinical Neg- ative)
Animal Contact								
No †								
Yes	3.26 (2.23- 4.77)***	2.95(1.96- 4.42)***	3.05 (1.99- 4.68)***	2.72 (1.71- 4.32)***	0.97(0.66-1.41)	0.88(0.581.32)	0.86 (0.60- 1.21)	0.82(0.56-1.19)
Gender								
Male †								
Female	1.09 (0.76-1.58)	1.09 (0.74-1.61)	1.04 (0.68-1.59)	1.09 (0.69-1.71)	1.27(0.90-1.78)	1.18(0.82-1.69)	1.08 (0.79- 1.47)	1.17 (0.84- 1.63)
Parent Education								
None †								
Primary	0.65 (0.33-1.36)	0.76 (0.36-1.67)	0.54 (0.23-1.18)	0.54 (0.25-1.23)	1.21(0.62-2.47)	0.97 (0.48-2.02)	1.02 (0.54- 1.93)	1.07 (0.55- 2.10)
Secondary	0.59 (0.29-1.23)	0.73 (0.35-1.61)	0.39 (0.18-0.87)*	0.38(0.170.89)*	0.79(0.40-1.64)	0.75 (0.37-1.57)	0.76 (0.41- 1.45)	0.80 (0.41- 1.58)
Tertiary	0.75 (0.35-1.64)	1.00 (0.45-2.31)	0.57 (0.25-1.33)	0.57 (0.24-1.38)	0.94(0.45-2.01)	0.81 (0.38-1.79)	0.71 (0.36- 1.42)	0.74 (0.36- 1.54)
Parent Occupation							,	,
Professional, skilled labour	1.16 (0.77-1.75)	1.13 (0.73-1.75)	0.93 (0.57-1.48)	0.78 (0.46-1.32)	1.16(0.79-1.69)	1.08 (0.71-1.62)	1.13 (0.79- 1.61)	1.12 (0.76- 1.65)
Ethnicity								
Ga Adangbe †								
Akan	1.22 (0.79-1.88)	1.31 (0.83-2.08)	0.93 (0.56-1.52)	0.92 (0.54-1.58)	1.09(0.73-1.64)	1.09(0.711.68)	1.34(0.93- 1.95)	1.28 (0.86- 1.91)
Ewe	0.99 (0.59-1.65)	1.05 (0.61-1.78)	0.98 (0.55-1.71)	1.05 (0.57-1.91)	0.99(0.62-1.58)	1.04(0.631.69)	1.04 (0.68- 1.59)	1.06 (0.67- 1.68)

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Northern	1.00 (0.44-2.16)	1.18 (0.49-2.68)	0.57 (0.18-1.49)	0.73 (0.23-1.94)	1.34(0.65-2.64)	1.19(0.542.52)	1.07 (0.54- 2.05)	0.99 (0. 2.04)	.47-
Unknown	1.56 (0.29-6.37)	1.77 (0.34-7.51)	0.43 (0.02-2.69)	0.46 (0.02-3.06)	0.59(0.09-2.38)	0.27(0.011.52)	1.29 (0.37- 4.21)	1.39 (0. 4.84)	.38-
Drug for current illn	ess								
None †									
Analgesic	0.96 (0.63-1.46)	0.94 (0.59-1.47)	0.57 (0.34-0.94)*	0.57(0.320.98)*	1.03(0.69-1.52)	1.03(0.681.58)	1.41 (0.99- 2.01)	1.43 (0. 2.11)	.98-
Other	1.70 (1.05-2.73)*	1.63 (0.98-2.71)	1.56 (0.93-2.60)	1.93(1.113.34)*	1.89(1.212.94*	1.90(1.183.07*	2.02 (1.33- 3.09)**	2.17 (1. 3.42)***	.38-

Odds ratios (OR) presented with 95% confidence intervals (95% CI).

* *P* < .05, ** *P* < .01, *** *P* < .001

† Reference category

DISCUSSION

This study found that animal contact was neither associated with self-reported nor measured fever in malarianegative children. Animal contact was associated with the secondary symptoms of headache and nausea, common yet non-specific symptoms of possible animal-transmitted infections, including leptospirosis, Q fever, and Campylobacteriosis.^{20,21} These findings, although inconclusive, may be attributed to the vast aetiology of fever, one or more undiagnosed zoonoses that present with headache and nausea but without fever, or the lack of context around the type of animal contact reported. But these associations beckon further study of the potential burden of bacterial zoonoses on undifferentiated paediatric fevers.

In the context of rising zoonosis infections globally particularly in the wake of the COVID-19 pandemic— Ghana may still benefit from a richer understanding of the prevalence and common symptoms of endemic zoonoses. Our results demonstrating borderline weaker associations between social determinants of health and the least objective forms of diagnosis (self-reported fever and clinical malaria) suggest that diagnostic technology continues to reproduce disparities in treatment.

While this study focused primarily on contact with dogs and cats, future studies might also consider contact with livestock such as goats and cattle, particularly in peri-urban and urban environments where higher population density can facilitate transmission, with a more detailed characterisation of animal contact, and a broader, more specific set of symptoms.

Household animals are increasingly recognised as reservoirs for neglected infections. For example, it was originally thought that ownership of cats or dogs did not increase the risk of Rickettsia,²² but more recent studies have found pets to be an emerging reservoir after all.⁸²³⁻²⁵ Dog ownership has been established as a risk factor for Campylobacter, recognised as one of the most common bacterial infections associated with pet ownership.^{21,26,27} Coagulase-negative *staphylococci* have been found in

clinically healthy dogs,28 and have the ability to transfer between pets and their owners, contributing to an increased risk of staphylococci infections.7 A case of identical Staphylococcus intermedius strains found in a woman and her dog highlighted the risk of transmission of bacterial infections from household pets to humans.²⁹ The infections detected in the present study, except HIV, could plausibly have been transmitted by animal contact, among diverse potential origins.7,9,30,31 The HIV case was a ten-year-old boy, and we retained this case in the analysis because it was plausible that the observed symptoms were due to some other undetected infection, not HIV. This suggests that more research is needed to evaluate animal contact as a risk for bacterial zoonoses, which may serve as an etiological driver of acute febrile illness in sub-Saharan Africa.

This study was limited by its cross-sectional design and limited data resolution about the patients'-built environment, outdoor activities, and nature or frequency of animal contact. It is plausible that feral animals pose different risks than pets or that specific transmission pathways shape outcomes more than the mere presence of animals. Additionally, the household data were self-reported by the child's guardian during recruitment at the clinic, which may contribute to selection and recall biases. The lack of available blood tests and nasopharyngeal results for every child meant that a majority of the fever cases remained unidentified. In a related study, efforts to model paediatric febrile illness using these data found that the blood culture test results were not strongly associated with differential diagnosis,³² hence our focus on symptomatology.

CONCLUSION

Improving our understanding of the aetiology of acute febrile illness in sub-Saharan Africa remains an important issue for improving diagnosis, patient care, and proper prescription of medications. Bacterial zoonoses, and other common infections transmitted by household animals, transmit many pathogens that cause human fever and may contribute to paediatric morbidity from undifferentiated acute febrile illness. Future studies should assess the nature of animal contact as a risk factor for fever, the socio-environmental factors that increase pathogen transmission risk from animals, and the types of diagnostic tests that may help identify and treat these infections in a clinical setting.

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REFERENCES

- 1. Kaboré B, Post A, Lompo P, et al. Aetiology of acute febrile illness in children in a high malaria transmission area in West Africa. *Clinical Microbiology and Infection.* 2020.
- 2. Maze MJ, Bassat Q, Feasey NA, Mandomando I, Musicha P, Crump JA. The epidemiology of febrile illness in sub-Saharan Africa: implications for diagnosis and management. *Clinical Microbiology and Infection.* 2018;24(8):808-814.
- 3. Crump JA, Kirk MD. Estimating the Burden of Febrile Illnesses. *PLOS Neglected Tropical Diseases*. 2015;9(12):e0004040.
- 4. Crump JA, Morrissey AB, Nicholson WL, et al. Etiology of severe non-malaria febrile illness in Northern Tanzania: a prospective cohort study. *PLOS Neglected Tropical Diseases*. 2013;7(7):e2324.
- 5. Wang H, Liddell CA, Coates MM, et al. Global, regional, and national levels of neonatal, infant, and under-5 mortality during 1990-2013: a systematic analysis for the Global Burden of Disease Study 2013. *The Lancet*. 2014;384(9947):957-979.
- Dalrymple U, Cameron E, Bhatt S, Weiss DJ, Gupta S, Gething PW. Quantifying the contribution of Plasmodium falciparum malaria to febrile illness amongst African children. *Elife*. 2017;6:e29198.
- Gómez-Sanz E, Ceballos S, Ruiz-Ripa L, Zarazaga M, Torres C. Clonally diverse methicillin and multidrug resistant Coagulase Negative Staphylococci are ubiquitous and pose transfer ability between pets and their owners. *Frontiers in microbiology*. 2019;10(485).
- 8. Hii SF, Kopp SR, Abdad MY, et al. Molecular evidence supports the role of dogs as potential reservoirs for rickettsia felis. *Vector-Borne and Zoonotic Diseases*. 2011;11(8):1007-1012.
- 9. van der Linden M, Al-Lahham A, Nicklas W, Reinert RR. Molecular characterization of pneumococcal isolates from pets and laboratory animals. *PLOS ONE*. 2009;4(12):e8286.
- Tsai K-H, Yen T-Y, Wu W-J, Carvalho R, Raoult D, Fournier P-E. Investigation of ctenocephalides felis on domestic dogs and rickettsia felis infection

in the Democratic Republic of Sao Tome and Principe. *Zoonoses and Public Health*. 2020;67(8):892-902.

- Robb K, Null C, Teunis P, Yakubu H, Armah G, Moe CL. Assessment of fecal exposure pathways in low-income urban neighborhoods in Accra, Ghana: rationale, design, methods, and key findings of the SaniPath Study. *The American Journal of Tropical Medicine and Hygiene*. 2017;97(4):1020-1032.
- 12. Adams EA, Price H, Stoler JB. Urban slums, drinking water, and health: Trends and lessons from sub-Saharan Africa. In: *Handbook of Global Urban Health*. Taylor and Francis; 2019:533-552.
- 13. Costa F, Carvalho-Pereira T, Begon M, Riley L, Childs J. Zoonotic and vector-borne diseases in urban slums: opportunities for intervention. *Trends in Parasitology*. 2017;33(9):660-662.
- 14. Ghana Statistical Service. 2010 Population & Housing Census Report: Disability in Ghana. Ghana Statistical Service; 2014.
- 15. Doe HW. Assessing the challenges of water supply in urban Ghana: The case of North Teshie, Kungliga Tekniska högskolan; 2007.
- Ahorlu CK, Dunyo SK, Afari EA, Koram KA, Nkrumah FK. Malaria-related beliefs and behaviour in southern Ghana: implications for treatment, prevention and control. *Tropical Medicine* & *International Health*. 1997;2(5):488-499.
- 17. Liu J, Ochieng C, Wiersma S, et al. Development of a TaqMan Array Card for acute-febrile-illness outbreak investigation and surveillance of emerging pathogens, including ebola virus. *Journal of Clinical Microbiology*. 2016;54(1):49-58.
- 18. Amoako N. Microbial Etiology of Acute Febrile Illness in Children Presenting to Hospitals in Ghana, University of Ghana; 2019.
- 19. Amoako N, Duodu S, Dennis FE, et al. Detection of Dengue Virus among Children with Suspected Malaria, Accra, Ghana. *Emerging Infectious Diseases*. 2018;24(8):1544-1547.
- 20. Chomel BB, Boulouis H-J, Maruyama S, Breitschwerdt EB. Bartonella spp. in pets and effect on human health. *Emerging Infectious Diseases*. 2006;12(3):389.
- 21. Rabinowitz PM, Gordon Z, Odofin L. Pet-related infections. *American Family Physician*. 2007;76(9):1314-1322.
- 22. Wiggers RJ, Stewart RS. Ownership of cats or dogs does not increase exposure to rickettsia typhi. *Texas Medicine*. 2002;98(6):56-57.
- 23. Mtshali K, Nakao R, Sugimoto C, Thekisoe O. Occurrence of Coxiella burnetii, Ehrlichia canis,

rickettsia species and naplasma phagocytophilumlike bacterium in ticks collected from dogs and cats in South Africa. *Journal of the South African Veterinary Association.* 2017;88:1-6.

- 24. Parola P. Rickettsia felis: from a rare disease in the USA to a common cause of fever in sub-Saharan Africa. *Clinical Microbiology and Infection*. 2011;17(7):996-1000.
- 25. Berrian AM, Martínez-López B, Quan V, et al. Risk factors for bacterial zoonotic pathogens in acutely febrile patients in Mpumalanga Province, South Africa. *Zoonoses and Public Health.* 2019;66(5):458-469.
- 26. Gras LM, Smid J, Wagenaar J, et al. Increased risk for Campylobacter jejuni and C. coli infection of pet origin in dog owners and evidence for genetic association between strains causing infection in humans and their pets. *Epidemiology and Infection*. 2013;141(12):2526-2535.
- 27. Mbindyo S. Pet ownership as a risk for human campylobacteriosis-a review. *Tanzania Veterinary Journal*. 2019;34(1):18-23.
- 28. Chah KF, Gómez-Sanz E, Nwanta JA, et al. Methicillin-resistant coagulase-negative staphylococci

from healthy dogs in Nsukka, Nigeria. *Brazilian Journal of Microbiology*. 2014;45(1):215-220.

- 29. Kempker R, Eaton M, Mangalat D, Kongphet-Tran T. Beware of the pet dog: a case of Staphylococcus intermedius infection. *The American Journal of the Medical Sciences*. 2009;338(5):425-427.
- Lee S, Hwang J, Kim J, et al. Biofilm production of coagulase-negative staphylococci isolated from rescued wild animals in the Republic of Korea. Acta Veterinaria Scandinavica. 2019;61(1):50-50.
- 31. Gharsa H, Slama KB, Gómez-Sanz E, et al. Molecular characterization of staphylococcus aureus from nasal samples of healthy farm animals and pets in Tunisia. *Vector-Borne and Zoonotic Diseases.* 2015;15(2):109-115.
- 32. Toh KB, Amoako N, Duodu S, et al. Decision support tool to predict causes of childhood febrile illness using a Bayesian model approach. *Unpublished manuscript*. 2020.

Supplementary Material

Table S1. Descriptive characteristics of select patient and household characteristics with significant bivariate associations with measured and self-reported fever, using alternate case inclusion criteria of cases without a clinical diagnosis of malaria (n = 593), rather than a parasite-negative test result.

	Measur	ed	Self-Reported		
	Febrile	Afebrile	Febrile	Afebrile	
Overall Rate	320 (54%)	273 (46%)	349 (59%)	244 (41%)	
Age (years)	M = 3.29	M = 5.24	M = 3.24	M = 5.55	
	(SD = 2.64)***	(SD = 4.21)	(SD = 2.58)***	(SD = 4.31)	
Gender					
Female	151 (47%)	143 (52%)	166 (48%)	128 (52%)	
Male	169 (53%)	130 (48%)	183 (52%)	116 (48%)	
Animal Contact					
Yes	88 (28%)	75 (27%)	93 (27%)	70 (29%)	
No	232 (72%)	198 (73%)	256 (73%)	174 (71%)	
Type of Animal					
Cat	51 (16%)	46 (17%)	54 (15%)	43 (18%)	
Dog	35 (11%)	28 (10%)	37 (11%)	26 (11%)	
Sheep/Goat	1 (<1%)	1 (<1%)	1 (<1%)	1 (<1%)	
Unknown	1 (<1%)	0 (0%)	1 (<1%)	0 (0%)	
Parent Education					
None	24 (8%)	21 (8%)	24 (7%)	21 (9%)	
Primary	119 (37%)	99 (36%)	130 (37%)	88 (36%)	
Secondary	103 (32%)	91 (33%)	114 (33%)	80 (33%)	
Tertiary	74 (23%)	62 (22%)	81 (23%)	55 (23%)	
Parent Occupation					
Professional/skilled labour	80 (25%)**	99 (36%)	92 (26%)***	87 (36%)	
Trader/unskilled labour/unemployed	240 (75%)	174 (64%)	257 (74%)	157 (64%)	
Ethnicity					
Ga Adangbe	124 (39%)	89 (33%)	132 (38%)	80 (33%)	
Akan	103 (32%)	99 (36%)	115 (33%)	87 (36%)	
Ewe	63 (20%)	66 (24%)	71 (20%)	58 (24%)	
Northern	23 (7%)	15 (5%)	24 (7%)	15 (6%)	
Unknown	7 (2%)	4 (1%)	7 (2%)	4 (2%)	
Drug for current illness					
None	88 (28%)***	131 (48%)	97 (28%)***	122 (50%)	
Analgesic	143 (45%)	101 (37%)	160 (46%)	84 (34%)	
Other	89 (27%)	41 (15%)	92 (26%)	38 (16%)	
Owns car, motorbike or refrigerator †					
Yes	11 (3%)	0 (0%)	11 (3%)	0 (0%)	
No	308 (96%)	272 (100%)	336 (96%)	244 (100%)	
Water Source†					
Pipe borne water	10 (3%)	1 (<1%)	10 (3%)	1 (<1%)	
Sachet water	310 (97%)	272 (99%)	339 (97%)	243 (99%)	

*P < .05, **P < .01, ***P < .001

† Associations not measured due to lack of variation

M = mean; SD = standard deviation

LPo	ort R.	1	2	3	4	5	6	7	L
Chikungunya	Chikungunya	24	25	25	25	25	25	25	Ľ
CCHF	CCHF	23			너그	너그	너그		Ц
Dengue	Dengue (mod)	22 🗋 🗋	47		5	겁Ъ	\Box	디그	Ц
Bundibugyo	Sudian	21 🗋 🗋				5			Ц
Ebola Zaire	Nipah	20 4	\Box	\Box					
Hantavirus HTN	Hantavirus SEO	19 🗋 🗋	\Box	\Box	\Box		\Box		Ц
Hepatitis E	Hepatitis E	18 🗋 🗋	\Box	\Box	57	\Box	\Box	\Box	Б
Marburg	Marburg	17 🗋 🗋	\Box	\Box	\Box	\Box	\Box	\Box	Б
Rift Valley Fever	Rift Valley Fever	16 🗋 🗋	\Box	\Box	\Box	\Box	\Box		Б
Yellow Fever	Yellow Fever	15 🗋 🗋	\Box	\Box		\Box	\Box		Ц
O'nyong	West Nile	14 🗖 🗍	52	52	52	52	52	52	Б
MS2	MS2	13							
Brucella	Brucella	12	\Box	\Box	\Box	\Box	\Box	\Box	Ц
18S	Bartonella	11							
PhHV	PhHV	10							
Coxiella burnetii	Coxiella burnetii	9匚 그	57	\Box	5	57	\Box		Б
Leptospira	Leptospira	« <u>Г</u> Д		$\Box \Box$					Б
Rickettsia	Rickettsia	747	\Box	\Box	\Box	\Box	\Box	\Box	
Salmonella enterica	Salmonella enterica	6 []	5		5	5			Б
Salmonella Typhi	Salmonella Typhi	₅∟⊐	디그	\Box	5	디그	\Box	디그	Ц
Yersinia pestis	Yersinia Pestis	4 🗋 🗋	디그	\Box	디그	디그	\Box	디그	Ц
Plasmodium	Plasmodium	зСД	ςЪ	\Box	5	디그	\Box	디그	Ц
T. brucei	T. brucei	2	5			디그		디그	Ц
16S	Leishmania	1							

Figure S1. Configuration of TaqMan array card with list of pathogens that were tested in the parent febrile illness study.

Pathogen assay

Clinical Sample Control

Manufacture Positive Control