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# 1 Ruminant-dense environments increase risk of Shiga toxin-producing

## 2 Escherichia coli independently of ruminant contact

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- 21

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## 22 ABSTRACT

23 Cattle and other domestic ruminants are the primary reservoirs of O157 and non-O157 Shiga 24 toxin-producing *Escherichia coli* (STEC). Living in areas with high ruminant density has been 25 associated with excess risk of infection, which could be due to both direct ruminant contact and 26 residual environmental risk, but the role of each is unclear. We investigated whether there is any 27 meaningful risk to individuals living in ruminant-dense areas if they do not have direct contact 28 with ruminants. Using a Bayesian spatial framework, we investigated the association between the 29 density of ruminants on feedlots and STEC incidence in Minnesota from 2010 to 2019, stratified 30 by serogroup and season, and adjusting for direct ruminant contact. For every additional head of 31 cattle or sheep per 10 acres, the incidence of O157 STEC infection increased by 30% (IRR 1.30; 32 95% CrI 1.18, 1.42) or 135% (IRR 2.35; 95% CrI 1.14, 4.20), respectively, during the summer 33 months. Sheep density was also associated with O157 STEC risk during winter (IRR 4.28; 95% 34 CrI 1.40, 8.92). The risk of non-O157 STEC infection was only elevated in areas with goat 35 operations during summer (IRR 19.6; 95% CrI 1.69, 78.8). STEC risk associated with ruminant 36 density was independent of direct ruminant contact across serogroups and seasons. Our findings 37 demonstrate that living in a ruminant-dense area increases an individual's risk of O157 and non-38 O157 STEC infection even without direct ruminant contact, indicating that prevention efforts 39 need to extend to community strategies for averting indirect transmission from local ruminant 40 populations.

41

42 IMPORTANCE

43 STEC are zoonotic enteric bacteria responsible for 2.5 million illnesses each year. Infections in
44 young children can be especially devastating, causing hemolytic uremic syndrome (HUS), a

45 debilitating and sometimes fatal form of acute kidney injury. STEC's primary reservoirs are 46 cattle and other domestic ruminants, and transmission can occur through food, water, animal 47 contact, and person-to-person. Living near ruminants poses a significant risk of STEC infection; 48 however, the proportion of that risk due to direct ruminant contact or other routes of transmission 49 is unknown. Our research demonstrates that direct ruminant contact is a substantial risk 50 irrespective of location, and that individuals living in ruminant-rich regions are at high risk of 51 STEC infection regardless of whether they come into contact with ruminants. These findings 52 indicate a need for multi-pronged prevention efforts that emphasize control of contamination in 53 the environments surrounding ruminant populations, in addition to biosafety precautions when 54 contacting ruminants directly.

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## 56 **INTRODUCTION**

Shiga toxin-producing *Escherichia coli* (STEC) are estimated to cause 2.5 million illnesses each
year (1), including 265,000 in the United States (2). *E. coli* O157:H7 remains the single most
common serotype in the U.S., but reported infections with non-O157 STEC surpassed infections
with the O157 serogroup in 2013 (3).

61

62 Cattle are considered STEC's primary reservoir, but STEC have been isolated from a wide 63 variety of species, and small ruminants such as sheep and goats have also been recognized as 64 important reservoirs (4). STEC can be transmitted from their animal reservoirs to human 65 populations through food, direct animal contact, and contaminated environments, including 66 water. While the largest STEC outbreaks are predominantly due to nationally or internationally 67 distributed food products, the majority of STEC cases are sporadic (5).

68

69 The role of local reservoirs in an individual's risk of STEC infection is unclear. Reported cases 70 are disproportionately from rural populations and individuals with animal contact. In Scotland, 71 an estimated 26% of O157 STEC infections were of environmental origin due to livestock in 72 rural areas (6). In Minnesota, 22% of O157 and 16% of non-O157 reported STEC infections had 73 an animal agriculture exposure, and large proportions specifically had contact with ruminants or 74 ruminant environments (7). Concordantly, multiple studies have identified local cattle density as 75 a risk factor for STEC infection (8-13), and the density of small ruminants has also been 76 associated with increased risk of STEC infection (6, 10, 14). One study examining the effect of 77 cattle, sheep, and goat density on STEC risk found that only goat density was associated with 78 incidence (14), suggesting that some of the studies identifying cattle density as a risk factor could

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79 be confounded by small ruminant density. Additionally, the majority of studies on ruminant 80 density have focused exclusively on the O157 serogroup. One study that examined multiple 81 STEC serogroups found differences in the presence and magnitude of the effect of cattle density 82 on incidence (11), supporting arguments that reservoirs and transmission for O157 and non-83 O157 STEC should not be assumed to be the same (15). 84 85 The existing studies on livestock density have been conducted almost exclusively in Europe. Geographic variation in STEC strains has been well-established (16, 17), and there is evidence 86 87 for genetic separation of strains from different hosts, driven by geography (18), suggesting the 88 potential for host differences between Europe and North America. Most importantly, current 89 studies of ruminant density have not separated the effects of direct animal contact and residual 90 environmental risk. It is unknown whether the risk associated with ruminant density is primarily 91 due to direct animal contact, a known risk factor (19–22), or whether simply living in an area 92 with ruminants significantly increases an individual's risk of STEC infection even in the absence

93 of direct contact.

94

95 Here, we undertake one of the first investigations of the role of ruminant density on O157 and 96 non-O157 STEC incidence in North America. We include cattle, sheep, and goat density in our 97 analysis to examine the effect of and simultaneously adjust for the collocation of multiple 98 species. In the first study to combine ruminant density and direct animal contact, we use direct 99 contact exposure information reported by cases to adjust for direct ruminant contact, isolating the 910 effect of ruminant density. We find that cattle and sheep density are associated with O157 92 incidence, goats are strongly associated with non-O157 incidence, and the risk posed by

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- 102 ruminant density is independent of direct ruminant contact. Our findings demonstrate that living
- 103 in a ruminant-dense area is sufficient to increase an individual's risk of O157 and non-O157
- 104 STEC infection even without direct ruminant contact, indicating that prevention efforts need to
- 105 extend to community strategies for averting indirect transmission from local ruminant
- 106 populations.
- 107

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## 108 MATERIALS & METHODS

## 109 STUDY POPULATION

110 Laboratory-confirmed (23), symptomatic STEC cases reported to the Minnesota Department of 111 Health (MDH) from 2010 through 2019 with complete data for home zip code were included in 112 the analysis. MDH conducts statewide, active surveillance for STEC. Individuals with STEC 113 were asked if they lived on, worked on, or visited a farm and if they had visited a public animal 114 venue such as a petting zoo, fair, or educational exhibit. Those responding yes were asked about 115 the presence of and direct contact with different animals. Reported contact with cattle, sheep, or 116 goats during the 7 days prior to illness were combined into a single direct ruminant contact 117 variable. We defined presence of cattle, sheep, or goats without direct contact, which constituted 118 exposure to the ruminant environment, as indirect ruminant contact. To stratify by season, 119 November through April were defined as "winter" and May through October as "summer". 120 Missing data for age, sex, direct and indirect ruminant contact, and serogroup were imputed 121 using the MICE package in R (24, 25). We imputed 40 datasets with 5 iterations each. Because 122 we used a Bayesian approach for the primary analysis, we summarized the imputed data in a 123 single dataset using the mode of each imputed variable for each individual. This study was ruled 124 exempt by the University of Minnesota IRB.

125

126 The 2010 census was used to obtain age/sex stratified population counts at the zip code

127 tabulation area (ZCTA) level (26). Age categories were collapsed to 0-4, 5-9, 10-49, and  $\geq$ 50

128 years to reflect important age populations for STEC infection. Sex was classified as male or

129 female, resulting in eight total age/sex categories.

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131 Minnesota ZCTA boundaries were obtained from the Minnesota Geospatial Commons (27). The

132 data contained boundaries for 887 ZCTAs fully or partially in the state of Minnesota. Of these,

133 six ZCTAs specific to a single business or business block were not contained in the 2010 Census

134 data and were dropped, yielding 881 ZCTAs for final analysis.

135

136 FOOD PRODUCTION ANIMALS

137 Data on food animal production facilities in Minnesota are reported to the Minnesota Pollution 138 Control Agency and were obtained from the Minnesota Geospatial Commons (January 27, 2023) 139 (28). Per state law, the Agency defines and registers all operations capable of holding  $\geq$ 50 animal 140 units, or  $\geq 10$  animal units within shorelands as "feedlots". Data included latitude/longitude, 141 registration information, and animal counts and types for 25,062 feedlot facilities with animals. 142 Data were updated daily and contained only the most current registration information for each 143 feedlot, with 4.7% of feedlots having new registrations or permit issuances, 94.7% having updated registrations, and 0.5% with missing registration types. Of all new registrations, 99% 144 145 had registration periods of 4 years or more. Based on this information, feedlots with new 146 registration dates during the period 2010-2019 or updated registrations within 4 years of 2019 147 (2020-2023) were retained for analysis (n=24,410). Animal types considered for analysis were 148 cattle (including dairy and beef cattle, bulls, and veal calves), swine, goats, and sheep. Latitude 149 and longitude coordinates of each feedlot were used to identify their ZCTA, and the total number 150 of animals in each ZCTA was calculated. Animal densities were calculated by dividing the total 151 number of animals by the area in acres of each ZCTA and multiplying by 10 to yield the number 152 of animals per 10 acres.

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#### 154 FOODNET POPULATION SURVEY

155 While contact with ruminants was ascertained directly from STEC cases, ruminant contact in the 156 rest of the population was unknown but was needed to establish an expected number of 157 individuals with ruminant contact. We determined the number of individuals per ZCTA expected 158 to have ruminant contact using the most recent CDC FoodNet Population Survey, which was 159 administered in 2018-2019 (29). The survey ascertained information on acute diarrheal illness 160 and associated risk factors, including whether an individual had contact with cattle, sheep, or 161 goats during the past 7 days. Demographics on each survey respondent were also collected, 162 including ZCTA and county of residence, age, sex, and month of response. As not all ZCTAs 163 were represented in the survey, we conducted subsequent analyses at the county level. We 164 mapped ZCTAs to counties using the Census ZCTA to County Relationship File (30) by 165 assigning the largest Minnesota county by area that overlapped the zip code. Age, sex, and 166 month of response were completed for all respondents. Individuals with missing county of 167 residence or animal contact were excluded from the analysis. 168 169 Three sampling frames (landline phone, cell phone, and address-based (ABS)) and two data 170 collection modes (computer-assisted telephone (CATI) and web) were used for data collection. 171 To account for systematic differences in responses and estimates across the sampling frames and 172 data collection modes, a collapsed variable was created with three categories: landline/CATI,

173 cellphone/CATI, and ABS/Web.

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#### 175 MODELING THE EXPECTED POPULATION WITH ANIMAL CONTACT

176 We used the FoodNet Population Survey data in a Bayesian spatial model to estimate the 177 probability of contact with cattle, goats, or sheep across the state. For the 87 counties in the state 178 of Minnesota, boundaries were obtained from the Minnesota Geospatial Commons (February 22, 179 2023) (31). The recommended analysis approach for the FoodNet Population Survey data uses 180 weighted generalized estimating equations (GEE) to fit a marginal model while controlling for 181 correlation across the three mode/frame categories. While GEE marginal estimation does not 182 translate to the Bayesian disease mapping framework, we incorporated mode/frame in the 183 analysis. The number of respondents in each county that reported contact with ruminants was 184 assumed to be binomially distributed with size equal to the total number of respondents in the 185 county. The probability of contact was modeled using a logit link function and covariates for age 186 group, season, and mode/frame category; sex was initially also included but removed to reduce 187 variability in the analysis. The estimates were collapsed across mode/frames by a weighted 188 average approach. The model also included a spatial random effect using an intrinsic Conditional 189 Autoregressive prior and an uncorrelated heterogeneity term to account for potential 190 overdispersion, which was modeled using an independent and identically distributed normal 191 prior. Independent, diffuse normal priors were used for coefficients of the fixed effects. The 192 model was fit using the R package NIMBLE (32, 33). Convergence was assessed on three 193 independent chains with different starting values using the Gelman-Rubin diagnostic (34), with 194 all parameters having values below 1.1.

195

196 Using the estimated county-level posterior mean probabilities, we calculated the expected

197 number of ruminant contacts in each ZCTA in each season by age and sex. To convert from the

198 county level to ZCTA level, all ZCTAs within a county were assumed to have the same expected 199 probability. Then, the expected probability of ruminant contacts in summer and winter were 200 multiplied by the ZCTA population in each age/sex group yielding an expected number of 201 contacts for both seasons by strata. Subtracting the expected number of contacts from the total 202 population, we also calculated the expected number of individuals with no ruminant contact. In 203 summary, we split the total population in each ZCTA into 16 strata based on the four age groups, 204 two sexes (although the estimated probability of contact was constant by sex), and expected 205 ruminant contact in the past 7 days.

206

207 MODELING STEC INCIDENCE BY RUMINANT DENSITY AND CONTACT

208 The outcome of interest was the number of STEC cases in each ZCTA during the period 2010-209 2019. As this is a count outcome, a Poisson model with a log link was used. Separate models 210 were fitted by season (summer: May-October, winter: November-April) and by STEC serogroup 211 (O157 vs. non-O157). Across all models, an offset for the total person-years at risk in each 212 age/sex/contact strata and ZCTA was used. Two groups of models were created, one without 213 individual-level direct ruminant contact and one incorporating a binary covariate for direct 214 ruminant contact for each case. In both models, cattle, sheep, and goat density were included as 215 covariates.

216

As before, a Bayesian disease mapping approach was used with spatially structured and
uncorrelated random effects. Independent, diffuse normal priors were used for coefficients of the
fixed effects. Computation was done using NIMBLE (32, 33) with convergence of three chains
assessed using the Gelman-Rubin diagnostic (34), with all parameters having values below 1.1.

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- From the models with direct contact included, we estimated the age-sex-contact-adjusted STEC
- rates per 100,000 individuals in each ZCTA. These adjusted rates were computed by combining
- the age-sex-contact-specific estimated STEC rates across age/sex/contact groups via direct
- standardization. Covariate effects were expressed as incidence rate ratios (IRRs) with 95%
- 225 Bayesian credible intervals (CrI). Incidence rates were mapped using cutoffs derived from the
- Healthy People 2020 goal for O157 STEC incidence of 0.6/100,000 and Healthy People 2030
- 227 goal for all STEC incidence of 3.7/100,000 (35, 36).
- 228

## 229 SENSITIVITY ANALYSIS

230 To assess the potential role of swine in STEC incidence in Minnesota, we repeated the primary

analysis including swine density. We also assessed whether incorporating indirect exposure to

ruminants (i.e., exposure to ruminant environments), in addition to direct contact, meaningfully

affected our results.

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# 235 **RESULTS**

236	There were 3,048 symptomatic STEC cases reported to MDH from 2010-2019. Of these, a valid
237	zip code was unavailable for two, resulting in 3,046 cases for analysis. STEC serogroup was
238	identified for 2,905 (95%) infections, including 1,225 (40%) O157 and 1,680 (55%) non-O157
239	STEC (Table 1). Overall, cases were more likely to be female (59%), and 608 (20%) were $<5$
240	years old. A slightly greater proportion of O157 than non-O157 STEC cases were reported
241	during the summer months (78% vs. 66%) and reported direct contact with ruminants (10% vs.
242	7.5%). Incidence of STEC differed substantially by ZCTA, with most areas falling at the
243	extremes with <0.6 cases per 100,000 or >7.4/100,000 (Figure 1).
244	
245	Animal density per 10 acres varied spatially across the state (Figure 2). Swine were the most
246	numerous animal we examined and cattle the most common ruminant (Appendix Table A1).
247	
248	The FoodNet Population Survey collected responses from 3,793 children and adults in
249	Minnesota. We were unable to determine the county of residence for 91 respondents (2.4%), of
250	which 81 had no ZCTA or county reported, and 10 had a ZCTA from outside of Minnesota
251	reported with no county. There were 24 respondents (0.6%) with no data on contact with cows,
252	goats, or sheep in the past 7 days. This left a sample of 3,678 for analysis from 534 of the 881
253	ZCTAs in Minnesota. Differences in contacts were observed by season, age, and sex (Appendix
254	Table A2). The probability of ruminant contact varied spatially and was greatest among 0-4-
255	year-olds during summer (Figure 3).
256	

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### 257 AGE AND SEX WERE ASSOCIATED WITH BOTH O157 AND NON-O157 STEC

258	Both age and sex were consistently associated with STEC incidence in Minnesota independent of
259	ruminant density. Regardless of serogroup, season, or adjustment for direct animal contact, 0-4-
260	year-olds were at the greatest risk of STEC infections (Table 2). During summer, the risk
261	decreased with each increasing age group. After adjusting for direct ruminant contact, 5-9-year-
262	olds had a 48% lower rate of O157 STEC (IRR 0.52; 95% CrI 0.41, 0.64) and 59% lower rate of
263	non-O157 STEC (IRR 0.41; 95% CrI 0.31, 0.52) during the summer than 0-4-year-olds.
264	Reductions in incidence were even more pronounced for 10-49 and $\geq$ 50-year-olds during the
265	summer (Table 2). During winter, incidence rates were also reduced for individuals ≥5-year-olds
266	across both serogroups, though without the same dose-response relationship as in summer.
267	
268	Individuals reporting female sex were at consistently greater risk of STEC infection, with only
269	slight variation between serogroups and seasons (Table 2). For O157 STEC, the IRR comparing
270	female cases to male ranged from 1.18 (95% CrI 1.04, 1.33) during summer to 1.67 (95% CrI
271	1.34, 2.07) during winter. Compared to O157 STEC, the relative increase in incidence of non-
272	O157 STEC among female cases was greater during the summer (IRR 1.54; 95% CrI 1.36, 1.73),
273	but did not increase as substantially during winter (IRR 1.64; 95% CrI 1.37, 1.96).
274	
275	Direct Ruminant Contact Contributes Substantially to STEC Risk

Direct contact with cattle, sheep, or goats in the 7 days before illness significantly increased the
risk of STEC infection across both serogroups and seasons. Among individuals reporting
ruminant contact, the observed incidence of O157 STEC infections per 100,000 was 70.9 in
summer and 24.5 in winter. For non-O157 STEC infections, incidence per 100,000 was 85.0 in

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summer and 53.5 in winter. Comparatively, among cases without ruminant contact, the observed

- 281 O157 STEC incidence rates per 100,000 were 13.6 in summer and 5.3 in winter and for non-
- 282 O157 STEC incidence were 13.2 in summer and 6.9 in winter.
- 283
- After adjustment for age, sex, and ruminant density, the effect of direct ruminant contact on
- 285 O157 STEC incidence was IRR 5.11 (95% CrI 4.47, 5.81) during the summer, and similar during
- 286 the winter (Table 2). Non-O157 risk increased 7.47-fold (IRR 7.47; 95% CrI 6.58, 8.42) with
- direct ruminant contact during the summer and 9.01-fold (IRR 9.01; 95% CrI 7.44, 10.83) with
- direct ruminant contact during the winter. In sensitivity analysis, combining individuals with
- direct ruminant contact and individuals with only indirect ruminant contact accentuated the
- 290 effects observed with only direct contact across all serogroups and seasons (Appendix Table A3)
- 291 For example, during summer, direct and indirect ruminant contact were associated with 7.54
- times (IRR 7.54; 95% CrI 6.61, 8.56) the risk of O157 STEC infection and 9.61 times (IRR 9.61;
- 293 95% CrI 8.47, 10.9) the risk of non-O157 STEC infection.
- 294

295 RUMINANT DENSITY WAS ASSOCIATED WITH STEC INCIDENCE INDEPENDENTLY OF DIRECT CONTACT

296 The risk of O157 STEC infection increased for individuals living in a ZCTA with a high density

297 of cattle during summer, and a high density of sheep year-round. (Table 2). During summer,

incidence increased 30% for every additional 10 head of cattle (IRR 1.30; 95% CrI 1.18, 1.42)

and 135% for every additional 10 sheep (IRR 2.35; 95% CrI 1.14, 4.20). The effect of sheep was

- 300 greater during winter, with an IRR of 4.28 (95% CrI 1.40, 8.92). Concordantly, O157 STEC rates
- 301 were greatest in the center and southwest of the state, where large concentrations of cattle and
- 302 sheep are found (Figure 4).

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304	The effects of cattle and sheep density on O157 STEC risk were unaffected by adjustment for
305	direct animal contact (Table 2). The elevated risk during summer persisted in ZCTAs in the
306	central and southwest regions even in the absence of direct animal contact (Figure 5).
307	Conversely, O157 STEC risk during winter was notably low in almost all ZCTAs among
308	individuals without direct ruminant contact, the ZCTA with the greatest density of sheep being
309	the single exception.
310	
311	Non-O157 STEC incidence was associated only with goat density and only during the summer,
312	with an IRR of 19.6 (95% CrI 1.69, 78.8) for each additional goat per 10 acres (Table 2).
313	Ruminant density was not associated with non-O157 STEC incidence during the winter. Non-
314	O157 STEC incidence was greatest in the center and southeast of the state (Figure 4), similar to
315	goat density, though not perfectly aligned (Figure 2). As for O157, the effects of ruminant
316	density on non-O157 STEC incidence did not meaningfully change after adjustment for direct
317	animal contact (Table 2). However, few ZCTAs remained at elevated risk of non-O157
318	infections in the absence of direct animal contact, though notably the ZCTAs with the highest
319	goat density were among these (Figure 5). Compared to O157 risk, non-O157 incidence was
320	elevated to a greater degree in a greater proportion of ZCTAs during winter.
321	
322	In our sensitivity analyses, combining individuals with direct ruminant contact with those with
323	only indirect contact made no meaningful different in risk associated with ruminant density
324	(Appendix Table A3), and swine density was not associated with either serogroup in either
325	season (Appendix Table A4).

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### 327 **DISCUSSION**

328 No prior study has examined the effect of ruminant density on STEC incidence while accounting 329 for direct ruminant contact. We found that the association between living in an area with a high 330 density of ruminants and incident STEC infection was independent of direct ruminant contact for 331 both O157 and non-O157 STEC, in both summer and winter. Our study also demonstrates 332 important differences in the associations of O157 and non-O157 STEC with agricultural animal 333 reservoirs. Cattle and sheep density were specifically associated with O157 STEC incidence in 334 summer and year-round, respectively, and goat density was associated with non-O157 incidence 335 in summer. The risk associated with ruminant density was largely unaffected by the inclusion of 336 direct or indirect ruminant contact, which significantly increased risk of both serogroups in both 337 seasons.

338

339 High ruminant density can theoretically increase the risk of STEC through multiple mechanisms. 340 Direct animal contact is principal among these, as shown by our stratified maps of ZCTA 341 incidence. During summer, for the population with direct ruminant contact, we found that 342 incidence rates of O157 and non-O157 STEC infections in almost all ZCTAs were each 343 >3.7/100,000, the Healthy People 2030 target for all serogroups combined (35). Even during 344 winter, at least half the ZCTAs had incidence rates >2.4/100,000 among the population with 345 direct ruminant contact. This is consistent with the estimated 11% of STEC outbreaks caused by 346 direct animal contact in the U.S. from 2010-2017 (19) and previous studies showing that direct 347 contact with cattle, particularly, has been associated with increased STEC risk (20–22). 348

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349 We also assessed the role of indirect contact through exposure to ruminant environments. When 350 we examined the effect of any ruminant contact, direct or indirect, the risk of O157 and non-351 O157 STEC infection was greater (e.g., IRR 7.5 for the summer, O157 analysis) than when 352 incorporating only direct contact (e.g., IRR 5.1). This suggests substantial STEC transmission 353 occurring when an individual lives in, works in, or visits areas with ruminants, even without 354 coming into direct contact with the animals. Contamination in soil around animal pens and on 355 fences or other fomites can pose an ongoing risk of STEC exposure, as demonstrated by an O157 356 outbreak among children visiting a barn over a week after it had held cattle (37). Secondary 357 transmission can also occur from individuals infected with STEC acquired through direct 358 ruminant contact. Vigilance in hand hygiene among individuals exposed to ruminant 359 environments but without direct contact and thorough decontamination of spaces previously 360 inhabited by ruminants could help reduce the risk of STEC infections associated with indirect 361 transmission.

362

363 Direct or indirect contact is not the only way in which living in a ruminant-dense area can 364 increase STEC risk. Our analysis demonstrates the important role of residual environmental risk 365 from ruminants in a region, independent of direct or indirect ruminant contact. The insensitivity 366 of ruminant density risk to adjustment for ruminant contact is likely due to the only partial 367 overlap of ruminant-dense ZCTAs (Figure 2) and ZCTAs with substantial direct ruminant 368 contact (Figure 3). The risk of direct ruminant contact was driven by statewide patterns and not 369 disproportionately by ZCTAs with high ruminant densities, so it had minimal impact on the risks 370 associated with ruminant density. We found that ZCTAs with STEC infection risk >2.4/100,000

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in the absence of direct contact were almost exclusively those with a high density of ruminants(Figure 5).

373

374 The residual environmental risk implied by the associations with ruminant density is likely 375 composed of multiple transmission routes, including via locally produced ruminant food 376 products, person-to-person (e.g., in child care or household settings) from an individual who had 377 direct contact with a ruminant, and via environmental reservoirs such as water contaminated by ruminants. Even dust contaminated by STEC has been shown to drift out of ruminant 378 379 environments to contaminate neighboring areas (38). Transmission chains involving other animal 380 species as intermediaries and direct contact with them could also be involved. How each of these 381 pathways contributes to the risk of living in a region with high ruminant density is unknown and 382 an area of ongoing work. Although no previous study has attempted to disentangle ruminant 383 contact from residual environmental risk, in Scotland, 26% of cases were estimated to be 384 environmental in origin (6), which would include both direct and indirect animal contact and 385 residual environmental risk.

386

Cattle are the prototypic STEC reservoir, largely because of their role in maintaining and transmitting O157 STEC. Our finding that O157 is associated with cattle density is consistent with several previous studies, including assessments of both ecological (8–13), and direct contact (20–22) risk. Our estimate of the magnitude of the association, IRR 1.30 for each additional head of cattle per 10 acres, was almost identical to that estimated by Frank et al. in Germany for O157 (2.45 per 100 cattle/km<sup>2</sup>, or ~1.36 per 1 cattle/10 acres) (11). Friesema et al. also stratified by season and similarly found an association between cattle density and O157 STEC incidence only

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394	during the summer months (12). However, Mulder et al. observed an association of O157 STEC
395	infections and small ruminant but not cattle density, and when small ruminants were analyzed
396	separately, only goats, not sheep, were associated with O157 incidence (14). This inconsistency
397	could be due to differences in local STEC ecology between Minnesota and the Netherlands. With
398	a different array of STEC strains, hosts may also differ. Our findings demonstrate significant
399	residual environmental O157 risk associated with cattle density and indicate transmission
400	pathways from cattle to the public, including through food products and environmental
401	contamination, should to be identified at a local level for tailored public health prevention
402	measures.
403	
404	We found that infection with the O157 serogroup was also associated with sheep density, which
405	agrees with other studies that have found an association between O157 and both cattle and sheep
406	(6, 10). Moreover, genomic evidence suggests that O157 strains circulate interchangeably
407	between cattle and sheep (18), supporting the finding that both cattle and sheep density would
408	contribute to STEC incidence. Sheep also stood out as the only species to be associated with
409	STEC incidence during what we defined as the "winter" months, November-April. Among
410	individuals without ruminant contact during winter, the only ZCTA with >2.4/100,000 O157 and
411	non-O157 STEC cases was the ZCTA with the highest sheep density. This may be because
412	sheering and lambing often occur in March and April, involving greater person-sheep contact
413	than at other times of the year.
414	
415	Surprisingly, we found that non-O157 incidence was not associated with cattle density. Studies

416 from both Europe and North America have identified high levels of non-O157 STEC carriage

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417	among cattle (39-42), and living, working, or visiting a farm or public animal venue has been
418	associated with increased risk of non-O157 infection (43). However, only one other study,
419	conducted in Germany, has examined the risk of cattle density on non-O157 STEC incidence. In
420	that study, they found that cattle density was associated with increased incidence of O111, O103,
421	and O145, but not O26 (11). It is possible that if we had sufficient power to examine individual
422	non-O157 serogroups, we would have identified this type of heterogeneity; however, it is also
423	possible that adjusting for the presence of other animals, particularly sheep and goats, would
424	have nullified the associations they observed for cattle, as agricultural areas are likely to contain
425	multiple species. Geographic variation in reservoirs is also likely, and cattle may not be as
426	significant a source of non-O157 STEC infections in Minnesota as they were in Germany.
427	
428	Non-O157 STEC have been associated with visiting petting zoos (43, 44), where goats and sheep
429	are the two most common species available (45). At 19.5 after adjustment for direct animal
430	contact, the IRR we estimated for the association of goats and non-O157 STEC incidence was
431	the highest we observed. The ZCTA with the highest density of goats had 0.44 goats per 10
432	acres, and the IRR quantifies the increase in risk for the addition of 1 goat per 10 acres. Thus,
433	
	even in the highest density ZCTA, the addition of 1 goat/10 acres would be equivalent to more
434	even in the highest density ZCTA, the addition of 1 goat/10 acres would be equivalent to more than tripling the number of goats. If goats pose a risk of non-O157 STEC infection, such a
434 435	even in the highest density ZCTA, the addition of 1 goat/10 acres would be equivalent to more than tripling the number of goats. If goats pose a risk of non-O157 STEC infection, such a relatively large increase can be understood to have the outsized impact on risk that we observed.
434 435 436	<ul> <li>even in the highest density ZCTA, the addition of 1 goat/10 acres would be equivalent to more</li> <li>than tripling the number of goats. If goats pose a risk of non-O157 STEC infection, such a</li> <li>relatively large increase can be understood to have the outsized impact on risk that we observed.</li> <li>However, the high IRR we estimated is also not unprecedented. In one case-control study,</li> </ul>
434 435 436 437	<ul> <li>even in the highest density ZCTA, the addition of 1 goat/10 acres would be equivalent to more</li> <li>than tripling the number of goats. If goats pose a risk of non-O157 STEC infection, such a</li> <li>relatively large increase can be understood to have the outsized impact on risk that we observed.</li> <li>However, the high IRR we estimated is also not unprecedented. In one case-control study,</li> <li>contact with goats was associated with a 21-fold increase in the odds of non-O157 STEC</li> </ul>

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popularity in agritourism suggests that additional prevention measures may be warranted atpublic animal contact venues.

441

Although STEC are most commonly found among ruminant reservoirs, outbreaks in Canada have been linked to pork products (46, 47), and at least one study has found an increased risk of non-O157 STEC infection associated with living on a farm with swine (43). After accounting for the presence of ruminant reservoirs, however, our study did not detect an association between STEC incidence and swine density. While swine are competent STEC hosts, they do not appear to serve as an important source of residual environmental risk for STEC infections in Minnesota.

449 While similar patterns of STEC risk by age and sex have been observed before, the literature is 450 inconsistent in the age groups used for analysis, and an effect of sex has only been observed in 451 some studies (12, 48–52). We found that 0-4 year olds were at greatest risk of both O157 and 452 non-O157 STEC, independent of ruminant density and contact, which is consistent with previous 453 studies showing 0-4 or 1-4 year olds at greatest risk of infection (12, 49, 53). Elevated risk in 454 young children may be from greater exposure to STEC, reporting bias due to increased severity 455 and lower thresholds for healthcare-seeking and diagnostic testing seen in this age group, or 456 naïve immune systems. The dose-response relationship between greater age and lower STEC risk 457 during the summer is suggestive of higher thresholds for healthcare-seeking and diagnostic 458 testing as age increases or the acquisition of immunity over time, which has been reported 459 previously in studies of farmworkers and their families (54, 55). We also found that during 460 summer, independent of ruminant density and contact, the relative increase in non-O157 STEC 461 risk for cases reporting female sex was greater than the increase in O157 STEC, and the highest

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462 IRRs across serogroups were observed during winter. This likely indicates a relative increase in

463 transmission routes more common among individuals of female sex during the winter, and that

- 464 such transmission routes are more important for non-O157 than O157 STEC.
- 465

466 The ZCTAs with the greatest STEC risk were overwhelmingly rural, consistent with previous

467 work (6, 10, 13). Understanding the source of STEC infections in rural areas, including direct

468 and indirect animal contact, secondary transmission, well water, and environmental

469 contamination, is central to prevention efforts. Health care access can be more difficult in rural

470 areas, particularly with hospital closures (56, 57), which may put STEC cases in rural areas at

471 greater risk of severe outcomes such as HUS. With O157 STEC still the largest cause of HUS,

- 472 prevention in rural areas is a priority.
- 473

474 Our study was limited by insufficient sample size to analyze individual non-O157 serogroups. 475 While there is likely some heterogeneity in reservoirs between serogroups, a previous study 476 found the effect of ruminant density to be mostly consistent across non-O157 serogroups (11). 477 The Minnesota Pollution Control Agency's definition of a feedlot excluded operations with <50 478 animals (or <10 animals within shorelands), so our analysis does not include small farms. We 479 believe the impact of this is minimal, as small farms are likely to be located in the same regions 480 as larger operations. To incorporate individual-level direct ruminant contact, we had to determine 481 what portion of the population is exposed to ruminants. The FoodNet Population Survey we used 482 to do this asked about contact with cows, sheep, or goats in a single question, prohibiting us from 483 assessing the effect of direct contact with individual species. The FoodNet Population Survey 484 was also limited in its sample size, leaving the populations in 347 of 881 ZCTAs unsurveyed,

which required us to determine the expected number of individuals with ruminant contact at the
county level. Consequently, ZCTA-level expected contact counts were not as specific as they
would have been if all ZCTAs had been covered by the survey. Finally, unrecognized factors
may have confounded these results. To minimize this possibility, we adjusted for age and sex,
jointly modeled all species in our analysis, and accounted for two types of spatial correlation.

491 CONCLUSION

492 Our results indicate a need to identify and mitigate transmission routes from local cattle, sheep, 493 and goat reservoir populations. For the first time, our study demonstrates that the risk posed by 494 living in an area with high ruminant density does not operate solely through direct contact with 495 ruminants or even exposure to ruminant environments. Thus, more work is needed to identify 496 prevention measures for local transmission occurring through food, person-to-person contact 497 with individuals who encounter ruminants, and contamination of neighboring areas including 498 water bodies and produce fields. At the same time, both direct and indirect ruminant contact 499 dramatically increase an individual's risk of O157 and non-O157 STEC at all times of year, 500 emphasizing the importance of reinforcing handwashing and other best practices for contacting 501 ruminants.

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- 510 CW: Data curation, Formal analysis, Methodology, Software, Validation, Visualization, Writing
- 511 original draft; WF: Data curation, Writing original draft; KS: Conceptualization,
- 512 Investigation, Resources, Writing review & editing; JR: Investigation, Resources, Writing –
- 513 review & editing; CAK: Investigation, Writing review & editing; GAMT: Conceptualization,
- 514 Funding acquisition, Methodology, Writing original draft.
- 515
- 516 DATA AVAILABILITY
- 517 Data on individuals with reported STEC infections is available upon request and with an
- 518 appropriate agreement from the Minnesota Department of Health. Analysis code is available
- 519 upon request from the corresponding author.
- 520

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# 700 **FIGURES**



701

## 702 FIGURE 1. Incidence rate of STEC infections in Minnesota, 2010-2019, by serogroup and

- season. Incidence is shown as cases per 100,000 population at the zip code tabulation area
- 704 (ZCTA) level.

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- 707 FIGURE 2. Density in animal operations across Minnesota, covering the period 2010-2019,
- 708 **by animal type.** Animal density is shown per 10 acres at the zip code tabulation area (ZCTA)
- rog level. ZCTAs without animal operations for a given animal type are shown in grey.
- 710

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711

712 FIGURE 3. Estimated probability of contact with ruminants in Minnesota, 2010-2019.

713 Probability was calculated from FoodNet Population Survey results from Minnesota with

714 complete data for county and/or zip code and ruminant contact (29). Ruminant contact was

715 defined as contact with a cow, sheep, or goat in the previous 7 days.

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717

- 718 FIGURE 4. Smoothed incidence rate estimates of Shiga toxin-producing E. coli infection by
- 719 season and serogroup in Minnesota, 2010-2019. Incidence rates were adjusted for age, sex,
- 720 and direct ruminant contact.

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- 723 FIGURE 5. Smoothed incidence rate estimates of Shiga toxin-producing E. coli by season,
- 724 serogroup, and direct ruminant contact in Minnesota, 2010-2019. Incidence rates were
- adjusted for age and sex.

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# 727 **TABLES**

# 728 TABLE 1. Characteristics of reported STEC cases, Minnesota 2010-2019, by serogroup. Unknown

values were imputed prior to analysis.

Characteristic	O157 N=1,225	Non-O157 N=1,680	Unclassified N=141	Overall N=3,046
Age (years)	26.77 (24.25)	27.62 (22.56)	30.52 (24.20)	27.41 (23.34)
Unknown	0	0	1 (0.7%)	1 (<0.1%)
Age group				
0-4	268 (22%)	313 (19%)	26 (19%)	607 (20%)
5-9	126 (10%)	96 (5.7%)	8 (5.7%)	230 (7.6%)
10-49	565 (46%)	946 (56%)	75 (54%)	1,586 (52%)
50+	266 (22%)	325 (19%)	31 (22%)	622 (20%)
Unknown	0	0	1 (0.7%)	1 (<0.1%)
Gender				
Female	670 (55%)	1,030 (61%)	84 (60%)	1,784 (59%)
Male	554 (45%)	648 (39%)	56 (40%)	1,258 (41%)
Unknown	1 (0.1%)	2 (0.1%)	1 (0.7%)	4 (0.1%)
Contact with ruminants				
Yes	124 (10%)	126 (7.5%)	6 (4.3%)	256 (8.4%)
No	705 (58%)	925 (55%)	6 (23%)	1,662 (55%)
Unknown	396 (32%)	629 (37%)	103 (73%)	1,128 (37%)
Season				
Summer	955 (78%)	1,117 (66%)	103 (73%)	2,175 (71%)
Winter	270 (22%)	563 (34%)	38 (27%)	871 (29%)

## 731 TABLE 2. Posterior means and 95% credible intervals for incidence rate ratios of STEC infection

by season.

Summer						
	Without ruminant contact		With ruminant contact			
Variable	0157	Non-O157	0157	Non-O157		
Age						
0-4	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)		
5-9	$0.44 (0.35, 0.55)^{a}$	$0.33 (0.26, 0.43)^{a}$	$0.52 (0.41, 0.64)^{a}$	$0.41 (0.31, 0.52)^{a}$		
10-49	$0.24 (0.20, 0.27)^{a}$	$0.33 (0.28, 0.38)^{a}$	$0.23 (0.20, 0.27)^{a}$	$0.32 (0.28, 0.37)^{a}$		
50+	0.19 (0.16, 0.23) <sup>a</sup>	$0.15 (0.13, 0.19)^{a}$	0.16 (0.13, 0.19) <sup>a</sup>	$0.12 (0.10, 0.15)^{a}$		
Sex						
Male	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)		
Female	1.18 (1.04, 1.33) <sup>a</sup>	1.54 (1.36, 1.74) <sup>a</sup>	1.14 (1.01, 1.29) <sup>a</sup>	1.48 (1.31, 1.67) <sup>a</sup>		
Cattle per 10 acres	1.31 (1.19, 1.44) <sup>a</sup>	1.02 (0.90, 1.14)	1.30 (1.18, 1.42) <sup>a</sup>	0.98 (0.87, 1.10)		
Goats per 10 acres	3.23 (0.26, 13.69)	21.52 (1.91, 89.42) <sup>a</sup>	2.88 (0.23, 12.22)	19.61 (1.69, 78.84) <sup>a</sup>		
Sheep per 10 acres	2.35 (1.14, 4.17) <sup>a</sup>	2.14 (0.95, 3.97)	2.35 (1.14, 4.20) <sup>a</sup>	2.15 (0.96, 3.96)		
<b>Ruminant contact</b>						
No	-	-	1.00 (Ref)	1.00 (Ref)		
Yes	-	-	5.15 (4.51, 5.86) <sup>a</sup>	7.46 (6.57, 8.43) <sup>a</sup>		
Winter						
	Without ruminant contact		With ruminant contact			
Variable	0157	Non-O157	0157	Non-O157		
Age						
0-4	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)		
5-9	$0.50 (0.29, 0.80)^{a}$	$0.23 (0.13, 0.37)^{a}$	$0.55 (0.31, 0.88)^{a}$	$0.27 (0.15, 0.44)^{a}$		
10-49	$0.50 (0.36, 0.69)^{a}$	$0.53 (0.41, 0.68)^{a}$	$0.49 (0.35, 0.68)^{a}$	$0.52 (0.41, 0.67)^{a}$		
50						
30+	$0.46 (0.31, 0.64)^{a}$	$0.28 (0.21, 0.37)^{a}$	$0.40 (0.27, 0.56)^{a}$	$0.22 (0.16, 0.30)^{a}$		
Sex	0.46 (0.31, 0.64) <sup>a</sup>	0.28 (0.21, 0.37) <sup>a</sup>	0.40 (0.27, 0.56) <sup>a</sup>	0.22 (0.16, 0.30) <sup>a</sup>		
Sex Male	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref)	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref)	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref)	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref)		
Sex Male Female	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup>	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup>	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup>	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup>		
Sex Male Female Cattle per 10 acres	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.96 (0.77, 1.14)	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup> 1.03 (0.87, 1.20)	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup> 0.89 (0.73, 1.07)	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup> 0.98 (0.82, 1.15)		
Sex Male Female Cattle per 10 acres Goats per 10 acres	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.96 (0.77, 1.14) 21.04 (0.05, 152.55)	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup> 1.03 (0.87, 1.20) 4.37 (0.03, 28.37)	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup> 0.89 (0.73, 1.07) 21.62 (0.04, 154.68)	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup> 0.98 (0.82, 1.15) 3.98 (0.02, 26.09)		
SexMaleFemaleCattle per 10 acresGoats per 10 acresSheep per 10 acres	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.96 (0.77, 1.14) 21.04 (0.05, 152.55) 4.37 (1.49, 9.18) <sup>a</sup>	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup> 1.03 (0.87, 1.20) 4.37 (0.03, 28.37) 2.59 (0.84, 5.51)	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup> 0.89 (0.73, 1.07) 21.62 (0.04, 154.68) 4.28 (1.40, 8.92) <sup>a</sup>	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup> 0.98 (0.82, 1.15) 3.98 (0.02, 26.09) 2.58 (0.82, 5.58)		
Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres Ruminant contact	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.96 (0.77, 1.14) 21.04 (0.05, 152.55) 4.37 (1.49, 9.18) <sup>a</sup>	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup> 1.03 (0.87, 1.20) 4.37 (0.03, 28.37) 2.59 (0.84, 5.51)	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup> 0.89 (0.73, 1.07) 21.62 (0.04, 154.68) 4.28 (1.40, 8.92) <sup>a</sup>	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup> 0.98 (0.82, 1.15) 3.98 (0.02, 26.09) 2.58 (0.82, 5.58)		
Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres Ruminant contact No	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.96 (0.77, 1.14) 21.04 (0.05, 152.55) 4.37 (1.49, 9.18) <sup>a</sup>	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup> 1.03 (0.87, 1.20) 4.37 (0.03, 28.37) 2.59 (0.84, 5.51)	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup> 0.89 (0.73, 1.07) 21.62 (0.04, 154.68) 4.28 (1.40, 8.92) <sup>a</sup> 1.00 (Ref)	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup> 0.98 (0.82, 1.15) 3.98 (0.02, 26.09) 2.58 (0.82, 5.58) 1.00 (Ref)		
Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres Ruminant contact No Yes	0.46 (0.31, 0.64) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.96 (0.77, 1.14) 21.04 (0.05, 152.55) 4.37 (1.49, 9.18) <sup>a</sup> -	0.28 (0.21, 0.37) <sup>a</sup> 1.00 (Ref) 1.64 (1.38, 1.96) <sup>a</sup> 1.03 (0.87, 1.20) 4.37 (0.03, 28.37) 2.59 (0.84, 5.51) -	0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.64 (1.31, 2.03) <sup>a</sup> 0.89 (0.73, 1.07) 21.62 (0.04, 154.68) 4.28 (1.40, 8.92) <sup>a</sup> 1.00 (Ref) 4.84 (3.72, 6.17) <sup>a</sup>	0.22 (0.16, 0.30) <sup>a</sup> 1.00 (Ref) 1.59 (1.33, 1.89) <sup>a</sup> 0.98 (0.82, 1.15) 3.98 (0.02, 26.09) 2.58 (0.82, 5.58) 1.00 (Ref) 8.867.29, 10.67) <sup>a</sup>		

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# 734 APPENDIX TABLES

- 735 *TABLE A1.* Characteristics of reported STEC cases, Minnesota 2010-2019, by serogroup.
- 736 Unknown values were imputed prior to analysis.

Animal Type	Number of Feedlots	Number of Animals	Mean per Feedlot <sup>a</sup>
Cattle	18,441	2,967,940	160.9
Swine	5,869	10,611,608	1808.1
Goats	684	45,262	66.2
Sheep	1,347	206,509	153.3

<sup>a</sup> Mean animals per feedlot calculated among feedlots with >0 animals of the given type.

739 **TABLE A2.** Estimated percentages of individuals with contact with a cow, sheep, or goat in the

Season	Age	Male (%)	Female (%)	Both Sexes Combined (%)
Summer	0-4	14.40 (14.01, 14.81)	16.05 (8.58, 28.03)	15.29 (10.83, 21.14)
	4-9	7.49 (5.51, 10.10)	11.12 (5.74, 20.45)	9.05 (7.03, 11.57)
	10-	3.68 (2.46, 5.46)	12.14 (9.9, 14.81)	8.22 (6.44, 10.45)
	49			
	50+	3.74 (3.22, 4.36)	5.59 (3.87, 8.01)	4.69 (3.98, 5.52)
Winter	0-4	3.62 (2.23, 5.83)	6.79 (4.59, 9.93)	5.03 (3.23, 7.75)
	4-9	3.07 (2.33, 4.06)	10.10 (6.88, 14.6)	6.88 (4.92, 9.55)
	10-	4.35 (3.51, 5.39)	8.68 (6.64, 11.28)	6.38 (5.72, 7.12)
	49			
	50+	4.73 (3.94, 5.67)	3.54 (2.73, 4.58)	4.11 (3.54, 4.77)

past 7 days by season, age, and sex, calculated from the FoodNet Population Survey (29).

- 742 TABLE A3. Posterior means and 95% credible intervals for incidence rate ratios of STEC
- 743 infection, combining direct ruminant contact and any reported exposure to a ruminant

## 744 environment.

Summer						
	With direct ruminant contact		With direct + indirect ruminant			
	only		contact			
Variable	0157	Non-O157	0157	Non-O157		
Age						
0-4	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)		
5-9	$0.52 (0.41, 0.64)^{a}$	0.41 (0.31, 0.52) <sup>a</sup>	$0.55 (0.43, 0.68)^{a}$	$0.43 (0.33, 0.55)^{a}$		
10-49	$0.23(0.20, 0.27)^{a}$	$0.32 (0.28, 0.37)^{a}$	$0.23 (0.20, 0.27)^{a}$	$0.32 (0.28, 0.37)^{a}$		
50+	0.16 (0.13, 0.19) <sup>a</sup>	0.12 (0.10, 0.15) <sup>a</sup>	0.15 (0.12, 0.18) <sup>a</sup>	0.12 (0.09, 0.14) <sup>a</sup>		
Sex						
Male	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)		
Female	1.18 (1.04, 1.33) <sup>a</sup>	1.54 (1.36, 1.73) <sup>a</sup>	1.18 (1.04, 1.33) <sup>a</sup>	1.54 (1.36, 1.73) <sup>a</sup>		
Cattle per 10 acres	1.30 (1.18, 1.42) <sup>a</sup>	0.98 (0.87, 1.11)	1.29 (1.17, 1.41) <sup>a</sup>	0.98 (0.86, 1.10)		
Goats per 10 acres	2.84 (0.23, 11.96)	19.51 (1.66, 79.08) <sup>a</sup>	2.78 (0.22, 11.70)	19.32 (1.70, 78.21) <sup>a</sup>		
Sheep per 10 acres	2.36 (1.14, 4.17) <sup>a</sup>	2.15 (0.95, 4.00) <sup>a</sup>	2.35 (1.12, 4.15) <sup>a</sup>	2.14 (0.94, 3.99)		
Ruminant contact						
No	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)	1.00 (Ref)		
Yes	5.11 (4.47, 5.81) <sup>a</sup>	7.47 (6.58, 8.42) <sup>a</sup>	7.54 (6.61, 8.56) <sup>a</sup>	9.61 (8.47, 10.88) <sup>a</sup>		
Winter						
***						
	With direct run	minant contact	With direct + in	direct ruminant		
···inter	With direct run on	minant contact ly	With direct + in con	direct ruminant tact		
Variable	With direct run on O157	minant contact ly Non-O157	With direct + in con O157	direct ruminant tact Non-O157		
Variable Age	With direct run on O157	minant contact ly Non-O157	With direct + in con O157	ndirect ruminant tact Non-O157		
Variable Age 0-4	With direct run on O157 1.00 (Ref)	ninant contact ly Non-O157 1.00 (Ref)	With direct + in con 0157 1.00 (Ref)	Adirect ruminant tact Non-O157		
Variable Age 0-4 5-9	With direct run on 0157 1.00 (Ref) 0.55 (0.31, 0.87) <sup>a</sup>	minant contact ly Non-O157 1.00 (Ref) 0.27 (0.16, 0.43) <sup>a</sup>	With direct + in con 0157 1.00 (Ref) 0.56 (0.32, 0.90) <sup>a</sup>	Indirect ruminant           Non-O157           1.00 (Ref)           0.28 (0.16, 0.44) <sup>a</sup>		
Variable           Age           0-4           5-9           10-49	With direct run           01           0157           1.00 (Ref)           0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup>	minant contact           ly           Non-O157           1.00 (Ref)           0.27 (0.16, 0.43) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup>	With direct + in           con           0157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup>	Image: mail of the second system         Image: mail of the seco		
Variable           Age           0-4           5-9           10-49           50+	With direct run           01           0157           1.00 (Ref)           0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.40 (0.27, 0.56) <sup>a</sup>	Iminant contact           Instrument           1.00 (Ref)           0.27 (0.16, 0.43) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup>	With direct + in           con $0157$ 1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup>	Indirect ruminant           Non-O157           1.00 (Ref)           0.28 (0.16, 0.44) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup>		
Variable Age 0-4 5-9 10-49 50+ Sex	With direct run on 0157 1.00 (Ref) 0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.40 (0.27, 0.56) <sup>a</sup>	Iminant contact           Indext           1.00 (Ref)           0.27 (0.16, 0.43) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup>	With direct + in           con           0157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup>	Image: Non-O157           1.00 (Ref)           0.28 (0.16, 0.44) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup>		
Variable           Age           0-4           5-9           10-49           50+           Sex           Male	With direct run on O157 1.00 (Ref) 0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref)	Iminant contact           Indext           1.00 (Ref)           0.27 (0.16, 0.43) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup> 1.00 (Ref)	With direct + in           con           0157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup> 1.00 (Ref)	Image: main of the second system           Non-O157           1.00 (Ref)           0.28 (0.16, 0.44) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup> 1.00 (Ref)		
Variable           Age           0-4           5-9           10-49           50+           Sex           Male           Female	With direct run on O157 1.00 (Ref) 0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup>	$\begin{array}{c} \textbf{minant contact} \\ \textbf{ly} \\ \hline \textbf{Non-O157} \\ \hline \\ \hline 1.00 \ (\text{Ref}) \\ \hline 0.27 \ (0.16, \ 0.43)^a \\ \hline 0.52 \ (0.41, \ 0.66)^a \\ \hline 0.22 \ (0.16, \ 0.29)^a \\ \hline \\ \hline \\ \hline 1.00 \ (\text{Ref}) \\ \hline 1.64 \ (1.37, \ 1.96)^a \end{array}$	With direct + in           con           0157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup> 1.00 (Ref)           1.68 (1.34, 2.08) <sup>a</sup>	Indirect ruminant         Non-O157 $1.00 (Ref)$ $0.28 (0.16, 0.44)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (Ref)$ $1.00 (Ref)$ $1.64 (1.38, 1.95)^a$		
Variable Age 0-4 5-9 10-49 50+ Sex Male Female Cattle per 10 acres	With direct run           on           O157 $1.00 (Ref)$ $0.55 (0.31, 0.87)^a$ $0.50 (0.35, 0.68)^a$ $0.40 (0.27, 0.56)^a$ $1.00 (Ref)$ $1.67 (1.34, 2.07)^a$ $0.89 (0.73, 1.07)$	minant contact           ly           Non-O157 $1.00 (\text{Ref})$ $0.27 (0.16, 0.43)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (\text{Ref})$ $1.00 (\text{Ref})$ $0.22 (0.16, 0.29)^a$ $0.98 (0.82, 1.15)$	With direct + in           con           0157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup> 1.00 (Ref)           1.68 (1.34, 2.08) <sup>a</sup> 0.87 (0.71, 1.05)	Image: matrix of the system           Non-O157           1.00 (Ref)           0.28 (0.16, 0.44) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup> 1.00 (Ref)           1.64 (1.38, 1.95) <sup>a</sup> 0.98 (0.82, 1.15)		
Variable Age 0-4 5-9 10-49 50+ Sex Male Female Cattle per 10 acres Goats per 10 acres	With direct run           on           O157 $1.00 (Ref)$ $0.55 (0.31, 0.87)^a$ $0.50 (0.35, 0.68)^a$ $0.40 (0.27, 0.56)^a$ $1.00 (Ref)$ $1.67 (1.34, 2.07)^a$ $0.89 (0.73, 1.07)$ $20.03 (0.04, 137.85)$	minant contact           ly           Non-O157 $1.00 (\text{Ref})$ $0.27 (0.16, 0.43)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (\text{Ref})$ $1.64 (1.37, 1.96)^a$ $0.98 (0.82, 1.15)$ $3.79 (0.02, 24.90)$	With direct + in           con           0157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup> 1.00 (Ref)           1.68 (1.34, 2.08) <sup>a</sup> 0.87 (0.71, 1.05)           20.03 (0.04, 137.85) <sup>a</sup>	Image: different system           Non-O157           1.00 (Ref)           0.28 (0.16, 0.44) <sup>a</sup> 0.52 (0.41, 0.66) <sup>a</sup> 0.22 (0.16, 0.29) <sup>a</sup> 1.00 (Ref)           1.64 (1.38, 1.95) <sup>a</sup> 0.98 (0.82, 1.15)           3.86 (0.02, 24.62)		
Variable Age 0-4 5-9 10-49 50+ Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres	With direct run           on           O157 $1.00 (Ref)$ $0.55 (0.31, 0.87)^a$ $0.50 (0.35, 0.68)^a$ $0.40 (0.27, 0.56)^a$ $1.00 (Ref)$ $1.67 (1.34, 2.07)^a$ $0.89 (0.73, 1.07)$ $20.03 (0.04, 137.85)$ $4.29 (1.42, 8.92)^a$	Iminant contact           Iv $1.00 (\text{Ref})$ $0.27 (0.16, 0.43)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (\text{Ref})$ $1.00 (\text{Ref})$ $0.98 (0.82, 1.15)$ $3.79 (0.02, 24.90)$ $2.59 (0.84, 5.66)$	With direct + in           con           O157           1.00 (Ref)           0.56 (0.32, 0.90) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.38 (0.27, 0.54) <sup>a</sup> 1.00 (Ref)           1.68 (1.34, 2.08) <sup>a</sup> 0.87 (0.71, 1.05)           20.03 (0.04, 137.85) <sup>a</sup> 4.29 (1.42, 8.92) <sup>a</sup>	Indirect ruminant         Non-O157 $1.00 (Ref)$ $0.28 (0.16, 0.44)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (Ref)$ $1.00 (Ref)$ $0.98 (0.82, 1.15)$ $3.86 (0.02, 24.62)$ $2.59 (0.83, 5.62)$		
Variable Age 0-4 5-9 10-49 50+ Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres Ruminant contact	With direct run on O157 1.00 (Ref) 0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.89 (0.73, 1.07) 20.03 (0.04, 137.85) 4.29 (1.42, 8.92) <sup>a</sup>	Iminant contact           Ivestimation $1.00 (Ref)$ $0.27 (0.16, 0.43)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (Ref)$ $1.64 (1.37, 1.96)^a$ $0.98 (0.82, 1.15)$ $3.79 (0.02, 24.90)$ $2.59 (0.84, 5.66)$	With direct + in           con           O157           1.00 (Ref) $0.56 (0.32, 0.90)^a$ $0.50 (0.35, 0.68)^a$ $0.38 (0.27, 0.54)^a$ 1.00 (Ref)           1.68 (1.34, 2.08)^a $0.87 (0.71, 1.05)$ 20.03 (0.04, 137.85)^a           4.29 (1.42, 8.92)^a	Image: direct ruminant tact         Non-O157 $1.00 (Ref)$ $0.28 (0.16, 0.44)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (Ref)$ $1.64 (1.38, 1.95)^a$ $0.98 (0.82, 1.15)$ $3.86 (0.02, 24.62)$ $2.59 (0.83, 5.62)$		
Variable Age 0-4 5-9 10-49 50+ Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres Ruminant contact No	With direct run on O157 1.00 (Ref) 0.55 (0.31, 0.87) <sup>a</sup> 0.50 (0.35, 0.68) <sup>a</sup> 0.40 (0.27, 0.56) <sup>a</sup> 1.00 (Ref) 1.67 (1.34, 2.07) <sup>a</sup> 0.89 (0.73, 1.07) 20.03 (0.04, 137.85) 4.29 (1.42, 8.92) <sup>a</sup> 1.00 (Ref)	minant contact           ly           Non-O157 $1.00 (\text{Ref})$ $0.27 (0.16, 0.43)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (\text{Ref})$ $1.64 (1.37, 1.96)^a$ $0.98 (0.82, 1.15)$ $3.79 (0.02, 24.90)$ $2.59 (0.84, 5.66)$ $1.00 (\text{Ref})$	With direct + in con 0157 1.00 (Ref) $0.56 (0.32, 0.90)^{a}$ $0.50 (0.35, 0.68)^{a}$ $0.38 (0.27, 0.54)^{a}$ 1.00 (Ref) $1.68 (1.34, 2.08)^{a}$ 0.87 (0.71, 1.05) $20.03 (0.04, 137.85)^{a}$ $4.29 (1.42, 8.92)^{a}$ 1.00 (Ref)	Image: display state in the image.         Non-O157 $1.00 (Ref)$ $0.28 (0.16, 0.44)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (Ref)$ $1.64 (1.38, 1.95)^a$ $0.98 (0.82, 1.15)$ $3.86 (0.02, 24.62)$ $2.59 (0.83, 5.62)$ $1.00 (Ref)$		
Variable Age 0-4 5-9 10-49 50+ Sex Male Female Cattle per 10 acres Goats per 10 acres Sheep per 10 acres Ruminant contact No Yes	With direct run           on           O157 $1.00 (\text{Ref})$ $0.55 (0.31, 0.87)^a$ $0.50 (0.35, 0.68)^a$ $0.40 (0.27, 0.56)^a$ $1.00 (\text{Ref})$ $1.67 (1.34, 2.07)^a$ $0.89 (0.73, 1.07)$ $20.03 (0.04, 137.85)$ $4.29 (1.42, 8.92)^a$ $1.00 (\text{Ref})$ $4.29 (1.42, 8.92)^a$	Iminant contact           Iv $1.00$ (Ref) $0.27$ (0.16, 0.43) <sup>a</sup> $0.52$ (0.41, 0.66) <sup>a</sup> $0.22$ (0.16, 0.29) <sup>a</sup> $1.00$ (Ref) $1.64$ (1.37, 1.96) <sup>a</sup> $0.98$ (0.82, 1.15) $3.79$ (0.02, 24.90) $2.59$ (0.84, 5.66) $1.00$ (Ref) $1.00$ (Ref)	With direct + in con 0157 1.00 (Ref) $0.56 (0.32, 0.90)^a$ $0.50 (0.35, 0.68)^a$ $0.38 (0.27, 0.54)^a$ 1.00 (Ref) $1.68 (1.34, 2.08)^a$ 0.87 (0.71, 1.05) $20.03 (0.04, 137.85)^a$ $4.29 (1.42, 8.92)^a$ 1.00 (Ref) $6.66 (5.20, 8.40)^a$	Image: direct ruminant stact         Non-O157 $1.00 (Ref)$ $0.28 (0.16, 0.44)^a$ $0.52 (0.41, 0.66)^a$ $0.22 (0.16, 0.29)^a$ $1.00 (Ref)$ $1.64 (1.38, 1.95)^a$ $0.98 (0.82, 1.15)$ $3.86 (0.02, 24.62)$ $2.59 (0.83, 5.62)$ $1.00 (Ref)$ $1.00 (Ref)$		

# 746 TABLE A4. Posterior means and 95% credible intervals for incidence rate ratios of STEC

747 infection, including swine density.

		Without ruminant contacts		With ruminant contacts		
	Variable	0157	Non-O157	0157	Non-O157	
	Age					
	0-4	1.00 (ref)	1.00 (ref)	1.00 (ref)	1.00 (ref)	
	5-9	0.44 (0.35, 0.55)	0.33 (0.26, 0.42)	0.52 (0.41, 0.64)	0.41 (0.31, 0.52)	
	10-49	0.24 (0.20, 0.28)	0.33 (0.28, 0.38)	0.23 (0.20, 0.27)	0.32 (0.28, 0.37)	
	50+	0.19 (0.16, 0.23)	0.15 (0.13, 0.19)	0.16 (0.13, 0.19)	0.12 (0.10, 0.15)	
	Sex					
	Male	1.00 (ref)	1.00 (ref)	1.00 (ref)	1.00 (ref)	
Summer	Female	1.18 (1.04, 1.33)	1.54 (1.36, 1.73)	1.18 (1.04, 1.33)	1.54 (1.36, 1.73)	
	Cattle per 10 acres	1.31 (1.19, 1.44)	1.02 (0.90, 1.14)	1.30 (1.18, 1.42)	0.99 (0.87, 1.11)	
	Goats per 10 acres	3.31 (0.25, 14.32)	21.22 (1.85, 88.68)	2.86 (0.22, 12.04)	19.42 (1.67, 79.69)	
	Sheep per 10 acres	2.36 (1.14, 4.20)	2.14 (0.94, 3.95)	2.36 (1.13, 4.20)	2.16 (0.95, 4.01)	
	Swine per 10 acres	1.00 (0.97, 1.03)	1.00 (0.97, 1.03)	1.00 (0.97, 1.03)	1.00 (0.96, 1.03)	
	Ruminant contact					
	No	1.00 (ref)	1.00 (ref)	1.00 (ref)	1.00 (ref)	
	Yes	-	-	5.10 (4.46, 5.81)	7.47 (6.58, 8.45)	
	Age					
	0-4	1.00 (ref)	1.00 (ref)	1.00 (ref)	1.00 (ref)	
	5-9	0.50 (0.29, 0.81)	0.23 (0.13, 0.37)	0.55 (0.31, 0.88)	0.27 (0.15, 0.43)	
	10-49	0.50 (0.36, 0.69)	0.53 (0.41, 0.68)	0.50 (0.35, 0.68)	0.52 (0.41, 0.67)	
	50+	0.46 (0.32, 0.65)	0.28 (0.20, 0.37)	0.40 (0.28, 0.56)	0.22 (0.16, 0.30)	
	Sex					
	Male	1.00 (ref)	1.00 (ref)	1.00 (ref)	1.00 (ref)	
Winter	Female	1.67 (1.34, 2.07)	1.64 (1.38, 1.95)	1.68 (1.34, 2.07)	1.65 (1.38, 1.96)	
	Cattle per 10 acres	0.97 (0.78, 1.16)	1.04 (0.87, 1.22)	0.91 (0.74, 1.10)	1.00 (0.83, 1.17)	
	Goats per 10 acres	19.42 (0.05, 136.53)	4.18 (0.03, 26.50)	18.10 (0.04, 128.20)	3.72 (0.02, 24.22)	
	Sheep per 10 acres	4.49 (1.45, 9.41)	2.65 (0.86, 5.64)	4.57 (1.49, 9.62)	2.66 (0.83, 5.75)	
	Swine per 10 acres	0.98 (0.94, 1.03)	0.98 (0.94, 1.03)	0.98 (0.94, 1.02)	0.98 (0.94, 1.03)	
	Ruminant contact					
	No	1.00 (ref)	1.00 (ref)	1.00 (ref)	1.00 (ref)	
	Yes	-	-	4.94 (3.77, 6.31)	9.02 (7.42, 10.82)	

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