

1 **Title**

2 Seroprevalence of *Toxoplasma gondii* and *Neospora caninum* infections in cattle in Mongolia

3

4 **Running title**

5 Distribution of *Toxoplasma* and *Neospora* in cattle in Mongolia

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23 **Authors' contributions**

24 Conceived and designed the experiments: BP, BB, NY, YN.

25 Performed the experiments: BP, RF, YN.

26 Analyzed the data: BP, PM, YN.

27 Contributed reagents/materials/analytical tools: BP, PM, RF, MI, YN.

28 Wrote the manuscript: BP, YN.

29 Critically revised the manuscript: BP, BB, NY, YN.

30

31 **Abbreviations:** HRP, horseradish peroxidase; iELISA, indirect enzyme-linked
32 immunosorbent assay; IgG, immunoglobulin G; NcSAG1, surface antigen 1 of *Neospora*
33 *caninum*; TgGRA7, dense granule protein 7 of *Toxoplasma gondii*.

34

35 **Abstract**

36 *Toxoplasma gondii* and *Neospora caninum* are protozoan parasites that cause huge economic
37 losses in animal industries worldwide. *N. caninum* can cause abortion storms and high culling
38 rates in cattle, whereas *T. gondii* infection is a significant concern in both human and animals
39 because it can induce abortion and clinical symptoms in immunocompromised hosts. The aim
40 of this study was to determine the seroprevalence of *T. gondii* and *N. caninum* in cattle in
41 Mongolia. Specific antibodies to *T. gondii* and *N. caninum* were detected by using an indirect
42 enzyme-linked immunosorbent assay (iELISA) based on recombinant antigens of dense
43 granule protein 7 of *Toxoplasma gondii* and surface antigen 1 of *Neospora caninum*,
44 respectively. A total of 1,438 cattle sera from 20 of 21 provinces of Mongolia and the capital
45 city of Ulaanbaatar were tested. Overall, 18.7% and 26.2% of cattle were positive for specific
46 antibodies to *T. gondii* and *N. caninum*, respectively. Prevalence rates were higher (*T. gondii*
47 infection: $P < 0.0001$, *N. caninum* infection: $P = 0.002$) in the central region of Mongolia (*T.*
48 *gondii* infection: 27.1%, *N. caninum* infection: 30.8 %) compared with western region,
49 suggesting that prevalence rates might be influenced by geographical condition, particularly
50 warmer temperatures around this area in Mongolia. The lowest prevalence rates were observed
51 in the western region of Mongolia (*T. gondii*: 9%, *N. caninum*: 20.8%). In addition, the
52 seroprevalence of *N. caninum* in female animals (27.5%) was significantly higher than that in
53 male animals (20.4%) ($P = 0.018$), suggesting an important risk factor of abortion and stillbirth
54 in cattle. The present results showed that *T. gondii* and *N. caninum* infections might be a risk
55 for public health and economy of the livestock industry in Mongolia. In conclusion, this study
56 demonstrates high seroprevalences of *T. gondii* and *N. caninum* in Mongolia and provides
57 valuable new data for development of control measures against these infections in Mongolia.

58

59 **Keywords:** *Toxoplasma gondii*; *Neospora caninum*; Seroprevalence; Cattle; Mongolia.

60 1. Introduction

61 Toxoplasmosis is a zoonotic disease caused by the protozoan parasite *Toxoplasma*
62 *gondii*, which infects warm-blooded animals including human and livestock as an intermediate
63 host. Domestic cats and other members of the family *Felidae* serve as definitive hosts of *T.*
64 *gondii*. At least 17 species of wild felines have been reported as definitive hosts, i.e., European
65 and African wild cats, Pallas' cat, Bobcat, leopard cat, Amur leopard cat, Iriomote cat, Ocelot,
66 Geoffroy's cat, Pampas cat, jaguarundi, cougar, leopard, jaguar, tiger, lion, and cheetah (Tenter
67 et al., 2000). Oocysts shed by the definitive host cause wide-ranging environmental
68 contamination (Hill et al., 2002). *T. gondii* tissue cysts are commonly seen in meat-producing
69 animals such as pig, sheep, and goats, while they are rare in beef and buffalo meat (Tenter et
70 al., 2000). Therefore, raw or undercooked meat from these animals is potentially hazardous if
71 ingested by humans or other animals.

72 Neosporosis is a serious disease in cattle and dogs caused by the protozoan parasite
73 *Neospora caninum*, which is a *Toxoplasma*-like organism. Major reproductive problems
74 caused by *N. caninum* infection are abortion and stillbirths in cattle (Dubey et al., 2007).
75 Canids, including the Australian dingo (*Canis lupus dingo*) (King et al., 2010), coyote (*Canis*
76 *latrans*) (Gondim et al., 2004), and gray wolf (*Canis lupus*) (Dubey et al., 2011), are definitive
77 hosts and can shed oocysts in their feces. *N. caninum* oocyst-contaminated food or water is
78 considered one route of infection for cattle (De Marez et al., 1999). In addition, vertical
79 transmission of *N. caninum* between dams and calves is another route of the infection
80 (Thurmond et al., 1997; Schares et al., 1998). Although there is no evidence that *N. caninum*
81 infection occurs in humans, anti-*N. caninum* antibodies are detected in humans (Ibrahim et al.,
82 2009; Tranas et al., 1999), suggesting it as a potential zoonotic pathogen.

83 The agricultural sector is the backbone of the economy in Mongolia. Livestock products,
84 in particular, are considered an [important](#) source of income. [Mongolian livestock populations](#)

85 (horses, cattle, camels, goats, and sheep) total 66 million head. Among these, the cattle
86 population is estimated to have included 4.3 million individuals in 2017 (National Statistics
87 Office of Mongolia, 2017). Although it is only a subsector of agriculture, livestock produces
88 nearly 30% to the gross domestic product in Mongolia (Shagdar, 2002).

89 In our previous study, the seroprevalence of *T. gondii* among sheep in seven provinces
90 of Mongolia was examined using an indirect enzyme-linked immunosorbent assay (iELISA)
91 based on recombinant *T. gondii* matrix antigen 1 and latex agglutination test (LAT). The overall
92 prevalence rate of *T. gondii* was 24% (42/175) and 16% (29/175) by iELISA and LAT,
93 respectively (Tumurjav et al., 2010). The seroprevalence of *T. gondii* in the wild Pallas cat
94 (*Otocolobus manul*), a small felid species, was 13% (2/15) (Brown et al., 2005). These results
95 imply the presence of *T. gondii* in Mongolia. In contrast, to our knowledge, there is no report
96 of the prevalence of *N. caninum* in Mongolia. Because the current prevalences of *T. gondii* and
97 *N. caninum* in Mongolia have not been well-studied, epidemiological evidence of the
98 prevalences of *T. gondii* and *N. caninum* in livestock is an urgent issue for reducing not only
99 economic loss of livestock production but also public health risks.

100 Previous studies have demonstrated that dense granule protein 7 of *T. gondii* (TgGRA7)
101 is a potential diagnostic marker of immunoglobulin G (IgG) in acute and chronic infections.
102 An iELISA based on recombinant TgGRA7 is available to differentiate *T. gondii* infections
103 from other infections with high specificity and sensitivity, in different animals, including goats,
104 sheep, cattle, donkeys, and pigs (Selseleh et al., 2012; Terwaki et al., 2013; Wang et al., 2014;
105 Ichikawa-seki et al., 2015; Fereig et al., 2016). There is substantial agreement between the
106 TgGRA7-based iELISA and LAT (sensitivity and specificity in cattle 84% and 88%,
107 respectively; in sheep 83% and 83%, respectively; in goats 82% and 88%, respectively) (Fereig
108 et al., 2016). Several studies have confirmed that surface antigen 1 (NcSAG1) of *N. caninum*
109 is a useful diagnostic antigen that can be used to detect specific antibodies against *N. caninum*

110 in cattle during acute and chronic infections (Chanan et al., 2003; Wilkowsky et al., 2011;
111 Hiasa et al., 2012; Takashima et al., 2013; Ichikawa-seki et al., 2016). Thus, the aim of the
112 present study was to conduct a large-scale examination of the seroprevalence of *T. gondii* and
113 *N. caninum* infections and risk factors for such infection in cattle in Mongolia using iELISAs
114 based on the recombinant proteins TgGRA7 and NcSAG1.

115

116 **2. Materials and methods**

117 *2.1. Ethics statement*

118 This study was performed in strict accordance with the recommendations of the Guide
119 for the Care and Use of Laboratory Animals of the Ministry of Education, Culture, Sports,
120 Science and Technology, Japan. The protocol was approved by the Committee on the Ethics
121 of Animal Experiments at Obihiro University of Agriculture and Veterinary Medicine, Obihiro,
122 Japan (permit number 18-15).

123

124 *2.2. Study area and samples*

125 Based on economic and geographical conditions, Mongolia is divided into four major
126 regions comprising 21 provinces: central region (Tov, Omnogobi, Gobisumber, Dornogobi,
127 Selenge, and Dundgobi), western region (Gobi-Altai, Khovd, Bayan-Olgii, Uvs, and Zavkhan),
128 eastern region (Sukhbaatar, Dornod, and Khentii) and khangai region (Khovsgol, Arkhangai,
129 Ovorkhangai, Bayankhongor, Bulgan, and Orknon). A total of 1,438 cattle sera from 20
130 provinces of Mongolia and the capital city of Ulaanbaatar were tested in this study (Table 1,
131 Figs. 1 and 2); no samples were available from Darkhan-Uul province, which is located in the
132 central region of Mongolia. Cattle sera collected from 2014 to 2016 were obtained from the
133 collection of serum samples at the Institute of Veterinary Medicine in Mongolia. Blood samples
134 were collected with venal puncture into glass tubes without anticoagulant from each animal in

135 the field. The blood samples were placed in an icebox and immediately transferred to the
136 laboratory for the preparation of sera. They were kept frozen at -30°C until analysis. Samples
137 were divided according to region, province, age group (1–4, 5–9, and 10–14 years), and sex
138 for risk factor analyses of the infections.

139

140 2.3. Serological testing

141 In the present study, recombinant proteins of NcSAG1 and TgGRA7 were used for
142 detection of specific antibodies (Terwaki et al., 2013; Chahan et al., 2003). Purified NcSAG1
143 fused with glutathione S-transferase (GST) was used for detection of *N. caninum*-specific
144 antibodies by iELISA. iELISA based on recombinant TgGRA7 protein was used for detection
145 of anti-*T. gondii* antibodies. Fifty microliters of purified antigen were coated onto ELISA plates
146 (Nunc, Roskilde, Denmark) with a coating buffer (carbonate-bicarbonate buffer, Sigma, St.
147 Louis, MO, USA) at a final concentration of $0.1\ \mu\text{M}$ and incubated at 4°C overnight. After
148 incubation, plates were washed with washing buffer [0.5% Tween 20 in phosphate-buffered
149 saline (PBS)], blocked with $100\ \mu\text{l}$ of blocking solution (3% skim milk in PBS) in each well,
150 and incubated at 37°C for 1 h. Plates were washed with washing buffer once. Cattle sera were
151 diluted at 1:200 with 3% skim milk in PBS, and then $50\ \mu\text{l}$ of positive control, negative control,
152 or test serum sample were added to the wells and incubated at 37°C for 1 h. After washing the
153 plates six times with washing buffer, $50\ \mu\text{l}$ of horseradish peroxidase (HRP)-conjugated anti-
154 bovine IgG (Bethyl Laboratories, Montgomery, TX, USA) diluted at 1:10,000 (1:1 v/v in
155 glycerol) with 3% skim milk in PBS were added to each well and incubated at 37°C for 1 h.
156 After washing the plates six times with washing buffer, $50\ \mu\text{l}$ of TMB substrate reagent (BD
157 Bioscience, San Diego, CA, USA) were quickly added to each well and incubated at room
158 temperature for 10 min in the dark. After incubation, $50\ \mu\text{l}$ of stop solution (1 M H_2SO_4) were
159 added to each well. Absorbance values at 450 nm (A_{450}) of each reaction were determined

160 using an ELISA reader. The readings for recombinant antigens were subtracted from those of
161 GST protein. The cut-off point was determined as the mean A_{450} value for standard negative
162 sera kept in our laboratory ($n = 8$ negative sera) plus 3 standard deviations. The standard
163 positive and negative sera used for the *Neospora* study were confirmed with a commercial
164 indirect fluorescent antibody test (IFAT; VMRD Inc., Pullman, WA, USA). The sera from
165 experimentally infected cattle were used as the standard positive sera (Nishimura et al., 2013).
166 The standard negative sera used for the *Toxoplasma* study were confirmed with a commercial
167 LAT (Toxocheck-MT, Eiken Chemical, Tokyo, Japan). Although we had no sera from
168 experimentally *T. gondii*-infected cattle, the reactivity of the recombinant TgGRA7 to field
169 bovine sera was confirmed in our previous studies (Ichikawa-Seki et al., 2015; Fereig et al.,
170 2016). In those studies, the specificity and sensitivity between a commercial LAT (Toxocheck-
171 MT) and the TgGRA7-based iELISA for cattle sera were calculated (sensitivity 84%,
172 specificity 88%; Fereig et al., 2016). In the case, the specificity and sensitivity between an
173 IFAT and an NcSAG1-based iELISA for cattle sera were also calculated (sensitivity 100%,
174 specificity 100%; Chahan et al., 2003).

175

176 2.4. *Statistical analysis*

177 The data analysis was performed with the GraphPad Prism 6 software (GraphPad software Inc.,
178 La Jolla, CA, USA). A χ^2 test was used to analyze the data. Associations were tested with odds
179 ratios (ORs) and 95% confidence intervals after adjustment. A P value of < 0.05 , determined
180 with the VassarStats online tool (<http://vassarstats.net/>), was considered to indicate statistically
181 significant differences.

182

183 3. Results

184 The overall seroprevalence rate of *T. gondii* in Mongolian cattle was 18.7% (range: 2.6–

185 42.1%) (Table 1). The highest seroprevalence rate of *T. gondii* was 42.1% in Dundgobi, which
186 is located in the central region of Mongolia. In addition, higher seroprevalence rates were
187 observed in Ovorkhangai of khangai region (41.7%) and the capital city of Ulaanbaatar
188 (38.5%). Lower seroprevalence rates were seen in Bayan-Olgii (2.6%) and Khovd (5.7%)
189 provinces in the western region (Table 1, Fig. 1). Higher seroprevalence rates of *T. gondii* were
190 found in the central region (27.1%) and the khangai region (23.3%) compared with that of the
191 western region (9.0%), which had the lowest seroprevalence rate (Table 2). This result suggests
192 that geographical condition may be a potential risk factor of *T. gondii* infection in Mongolia.
193 Although age and sex were also analyzed as potential risk factors of *T. gondii* infection, no
194 significant differences were observed (Table 3).

195 The overall seroprevalence of *N. caninum* in cattle of Mongolia was 26.2% (range:
196 10.5%–69.2%) (Table 1). The highest seroprevalence rate was observed in the capital city of
197 Ulaanbaatar (69.2%). In addition, Dornogobi (42.9%) and Dundgobi (44.7%) provinces in the
198 central region and Zavkhan province (43.9%) in the western region showed higher
199 seroprevalence rates. The three lowest seroprevalence rates were seen in the western region,
200 Bayan-Olgii (10.5%), Uvs (12.3%) and Gobi-Altai (13.3%) (Table 1, Fig. 2). The
201 seroprevalence rates of *N. caninum* in the central region (30.8%) and the khangai region
202 (30.7%) were higher than that in the western region (20.8%) (Table 4). Risk factor analysis of
203 *N. caninum* seroprevalence showed no significant difference according to age, while a
204 significant difference in *Neospora* seroprevalence was observed according to sex (Table 5). In
205 particular, the seropositive rate of female animals (27.5%) was higher than that of male animals
206 (20.4%) ($P = 0.018$).

207 Mixed infections of *T. gondii* and *N. caninum* were also observed among the cattle in
208 Mongolia. The overall seroprevalence of mixed infections was 6.8% and the infection rates in
209 different provinces ranged from 0% to 38.5% (Table 1). Higher rates of mixed infections were

210 observed in Ulaanbaatar (38.5%), Dundgobi (31.6%), Dornogobi (21.4%), and Ovorkhangai
211 (21.4%) than in the other provinces, in which they ranged from 0% to 12.2% (Table 1).

212

213 **4. Discussion**

214 The livestock sector plays an important economic role in Mongolia because milk and meat
215 products derived from cattle are served as foods in the daily life of Mongolians. The meat most
216 frequently consumed in Mongolia is from sheep (31.5%), goats (27.7%), and cattle (21.3%).
217 An annual report of the milk industry in Mongolia in 2016 estimated the milk production from
218 cows, sheep, and goat to be 522.4, 99.1, and 173.6 million liters, respectively. Therefore, cattle
219 are one of the most important food sources in Mongolia (National Statistics Office of
220 Mongolia, 2017). Although *T. gondii* or *N. caninum* infection affects the livestock industry by
221 decreasing productivity due to abortion and stillbirth in cattle and neurological symptoms in
222 calves, available data of the prevalences of these protozoan parasites in Mongolia are limited.
223 Therefore, in the present study, cattle sera from 20 of 21 provinces of Mongolia were examined
224 for seropositivity to *T. gondii* and *N. caninum*. Overall seroprevalences of *T. gondii* and *N.*
225 *caninum* in cattle of Mongolia were 18.7% and 26.2%, respectively. The mixed infection rate
226 was 6.8%. China and Russia are neighboring countries of Mongolia. In China, the prevalence
227 rate of *T. gondii* in cattle was 5.7% in the southern region when determined with an indirect
228 hemagglutination antibody test (Zhou et al., 2012) and 4.8% in the northwest region when
229 determined with a modified agglutination test (Tan et al., 2015). However, the prevalence of
230 *Toxoplasma* in cattle in Russia has not been reported. When the seroprevalence of *N. caninum*
231 in cattle was determined with an ELISA, the prevalence rates were 13.3% in northeastern China
232 (Wang et al., 2010), 18.9% in southern China (Xia et al., 2011), and 9.97% in Russia (Hemphill
233 and Gottstein, 2000). Thus, the seroprevalences of *T. gondii* and *N. caninum* in cattle of

234 Mongolia were higher than those reported in China and Russia, indicating that *T. gondii* and
235 *N. caninum* infections may be a hazard to the livestock industry in Mongolia.

236 In our study, both *N. caninum* and *T. gondii* prevalence rates were significantly higher
237 in the central region of Mongolia, suggesting that one risk factor of such infections might be
238 climate condition, particularly warmer temperatures around this area. In contrast, the overall
239 lowest *T. gondii* prevalence rate was observed in the western region of Mongolia. According
240 to climate and geographical conditions, the average annual temperature of the central region is
241 higher because most of the largest desert (Gobi) is located around this area. Conversely, the
242 western region of Mongolia is the coldest and has the highest elevation (Altai Mountain area).
243 Although *T. gondii* oocysts (and likely *N. caninum* oocysts) are resistant to environmental
244 conditions such as high temperature, drying, and freezing (Dubey et al, 1970; Dubey et al,
245 1998), these infections may increase more readily in warmer and humid conditions than in
246 colder and drier climates (Tenter et al., 2000). Thus, climate and geographical conditions in
247 Mongolia may affect both *N. caninum* and *T. gondii* seroprevalence rates.

248 *Toxoplasma gondii* is not the main causative agent of abortion in cattle, and the
249 contribution of milk and meat from infected cattle to the prevalence of *Toxoplasma* in humans
250 is unknown (Dubey, 1986). However, the consumption of undercooked or raw beef and cow's
251 milk are considered risk factors for human infection. In Mongolia, although the prevalence of
252 *T. gondii* in cattle may be an infectious source to human, human *T. gondii* infection is not well
253 studied. The TORCH test is a blood screen for *T. gondii*, rubella virus, cytomegalovirus, herpes
254 simplex, human immunodeficiency virus, and other organisms, which lead to severe fetal
255 anomalies and fetal loss (Kaur et al 1999). The TORCH test is recommended for pregnant
256 women with high-risk pregnancies in Mongolia. Although TORCH infections were tested
257 among 100 pregnant women in Mongolia, *T. gondii* infection was not detected by ELISA
258 (Otgontsetseg et al., 2013). However, large-scale screening of *T. gondii* is required to

259 understand the impact of toxoplasmosis in Mongolia. Meat and meat-derived products
260 containing *T. gondii* tissue cysts are potential sources of infection in human (Tenter et al.,
261 2009). However, consumption of undercooked and raw meat is not a typical practice among
262 Mongolians. Therefore, intake or consumption of contaminated water, soil, and raw vegetables
263 might be a potential risk factor of *T. gondii* infection in human.

264 Although another source of *T. gondii* infection is cats and members of the family *Felidae*,
265 cats are not common domestic animals in Mongolia. Generally, domestic cats are not
266 widespread in any region of Mongolia because herders do not usually keep cats as pets.
267 However, small numbers of domestic cats are kept in households to control rodents. Several
268 species of wild cats inhabit Mongolia such as the wild cat (*Felis silvestris*), Eurasian lynx (*Lynx*
269 *lynx*), Pallas's cat (*O. manul*), and snow leopard (*Uncia uncia*). Among them, wild cats are
270 distributed in desert areas (Gobi) and Mongolian Altai Mountain area. However, hybridization
271 between wild and domestic cats has been observed, which influences the genetic purity of wild
272 cats, and competition between prey domestic cats and wild cats can lead to disease
273 transmission. The Eurasian lynx (*L. lynx*) is widely distributed in Mongolia (Clark et al., 2006).
274 The snow leopard is distributed in the mountainous area of western Mongolia, but they are rare
275 in all parts of Mongolia. The snow leopard is also thought to prey on livestock (Mallon 1984;
276 Clark et al., 2006; Shehzad et al., 2012). Thus, the transmission of *T. gondii* infection via
277 oocysts shed by wild felines may also facilitate infection in human (especially nomads) and
278 livestock.

279 In the present study, the seroprevalence of *N. caninum* was significantly higher in female
280 cattle (27.5%) compared with male cattle (20.4%), suggesting sex is an important risk factor
281 because of expected reproductive problems such as abortion and stillbirth in cattle. *N. caninum*-
282 specific antibody titers increase during pregnancy and peak within a month after calving in *N.*
283 *caninum*-infected dams (Ybañez et al., 2013). Therefore, the detection rate of *N. caninum*-

284 specific antibodies in female cattle might be higher than that in male animals. However, this
285 result may be attributable to the unequal sample sizes tested (female: 1,174, male: 264).

286 Furthermore, the *N. caninum* infection rate in cattle was shown to increase in the
287 presence of dogs on farms as a neighborhood animal (Dubey et al., 2007). Mongolian herders
288 keep dogs in their households as herding dogs. Although most herders and farmers maintain
289 several dogs, they generally do not check the health of their dogs or medicate them. Therefore,
290 the high prevalence of *N. caninum* infection in Mongolian cattle may be attributable to their
291 contact with the feces of *N. caninum*-infected domestic dogs. Consequently, further studies are
292 urgently required to determine the infection rate of *N. caninum* in dogs in this region to better
293 understand the infection source. In addition, wild canids should be considered as the definitive
294 host of *N. caninum*. Wild canids such as gray wolf (*Canis lupus*), Asiatic wild dog (*Cuon*
295 *alpinus*), raccoon dog (*Nyctereutes procyonoides*), and corsac fox (*Vulpes corsac*) are
296 distributed in Mongolia. Among them, the gray wolf is widely distributed and considered prey
297 livestock in Mongolia (Clark et al., 2006). However, to our knowledge, there are no reports
298 about the prevalence of *N. caninum* infection in these dogs in Mongolia. Further investigations
299 are needed to assess the relationship between dogs and cattle for *N. caninum* infection in
300 Mongolia.

301

302 **5. Conclusions**

303 We conducted the first study of the seroprevalence of *T. gondii* and *N. caninum* infections
304 among cattle in Mongolia. Because the seroprevalences of the infections in cattle of Mongolia
305 were higher than those reported in China and Russia, *T. gondii* and *N. caninum* infections may
306 be a hazard to the livestock industry in Mongolia. Additionally, the seroprevalence rates were
307 affected by climate and geographical conditions in Mongolia. However, to understand the
308 distribution patterns of such infections, identify major risk factor(s), and establish more potent

309 control measures, additional surveys for toxoplasmosis and neosporosis in Mongolia among
310 different livestock are required.

311

312 **Conflict of interests**

313 All authors declare that they have no financial or personal conflicts.

314

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442 **Table titles**

443 **Table 1.** Seroprevalence of *T. gondii* and *N. caninum* in cattle in different provinces of
444 Mongolia.

445 **Table 2.** Seroprevalence of *T. gondii* in different regions of Mongolia.

446 **Table 3.** Analysis of risk factors associated with *T. gondii* infection in Mongolian cattle.

447 **Table 4.** Seroprevalence of *N. caninum* in different regions of Mongolia.

448 **Table 5.** Analysis of risk factors associated with *N. caninum* in Mongolian cattle.

449

450 **Figure legends**

451 **Fig. 1.** Geographical distribution of *T. gondii* infection in Mongolian cattle used in this study.

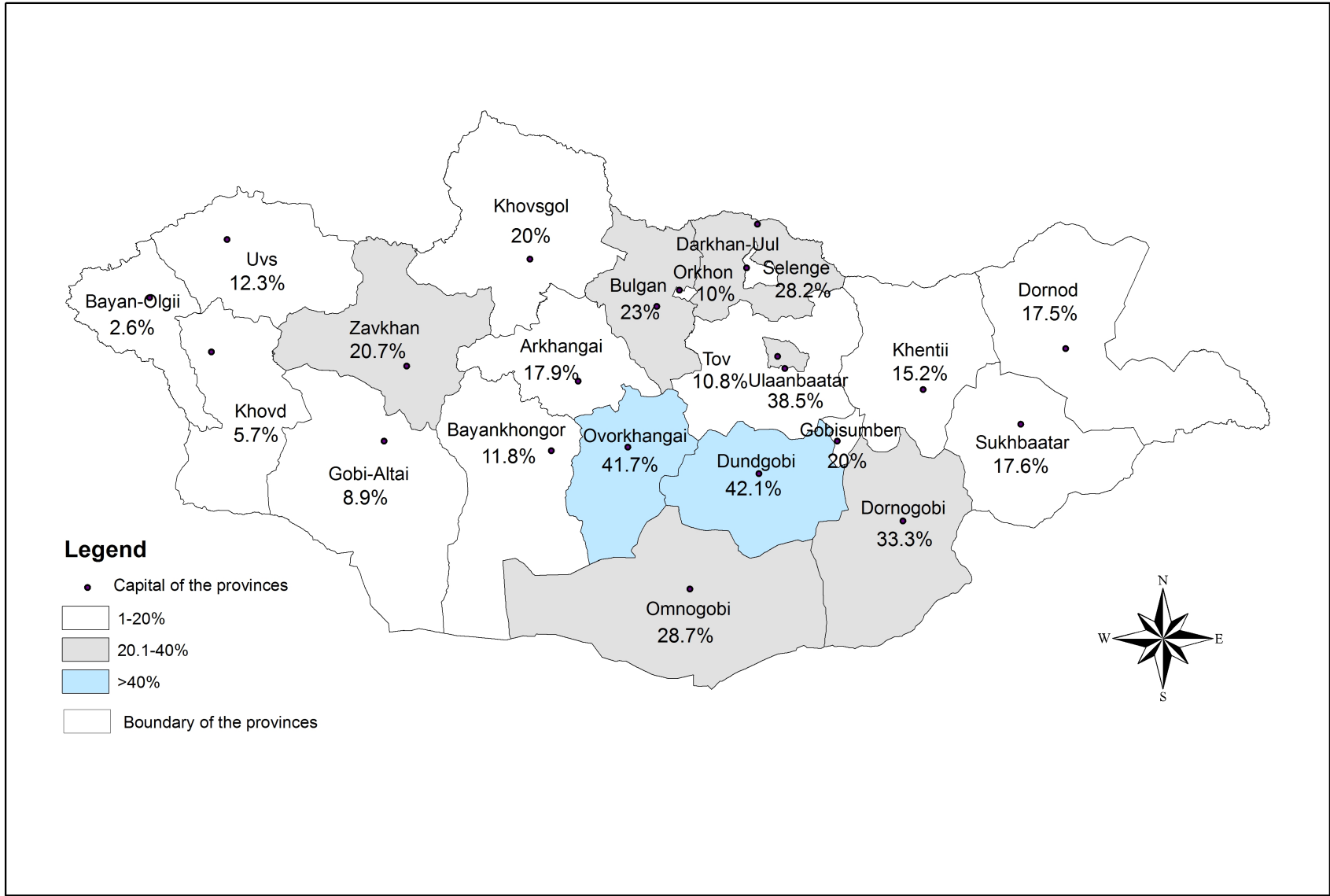
452 Infection rates are indicated as follows: white 1%–20%, gray 20.1%–40%, and light blue
453 >40%.

454

455 **Fig. 2.** Geographical distribution of *N. caninum* infection in Mongolian cattle used in this study.

456 Infection rates are indicated as follows: white 1%–20%, gray 20.1%–40%, and light blue
457 >40%.

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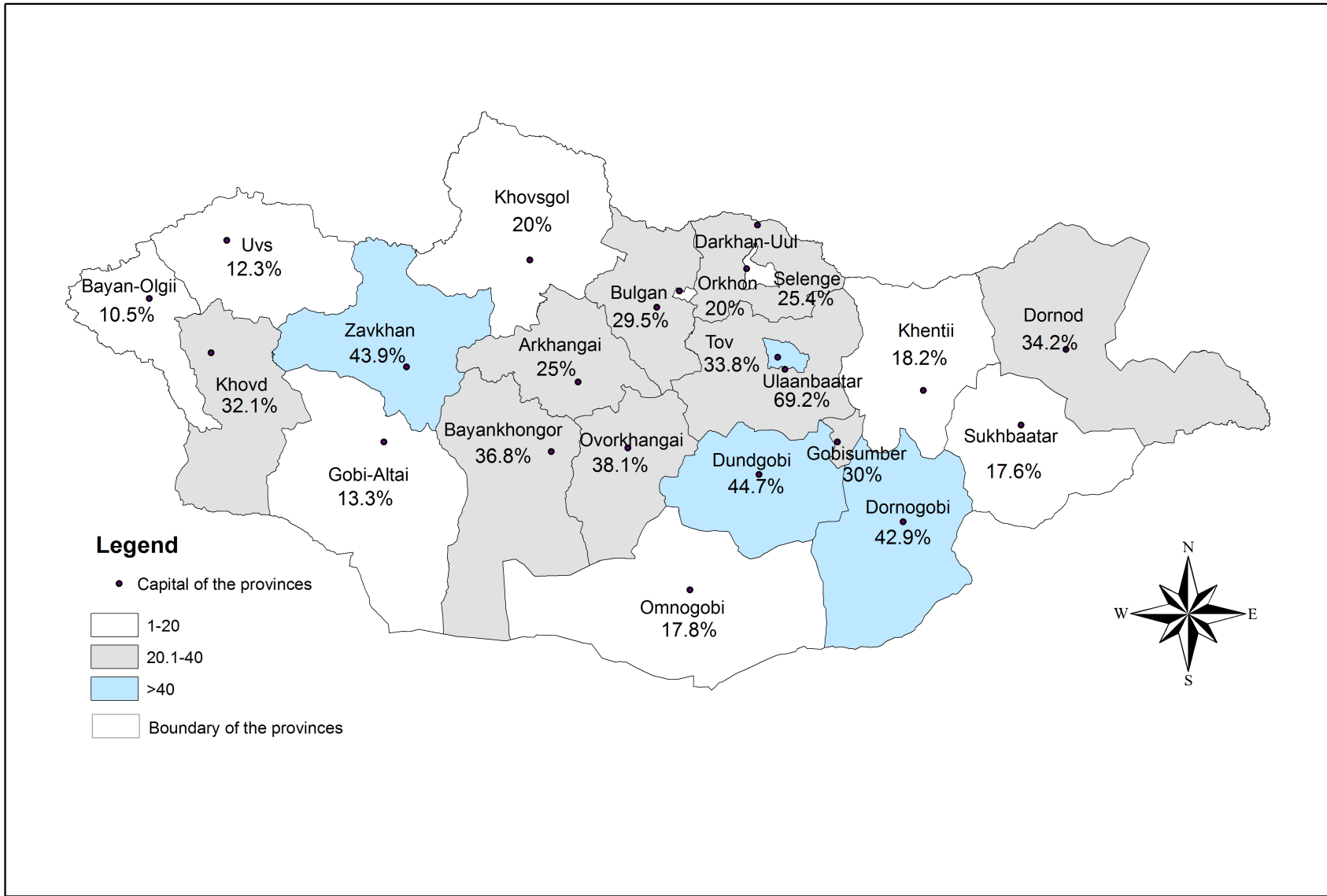


Table 1Seroprevalence of *T. gondii* and *N. caninum* in cattle in different provinces of Mongolia.

Regions of Mongolia	Provinces	Seroprevalence of <i>T. gondii</i> % (No. positive / No. sampled)	95% CI of <i>T. gondii</i> seroprevalence	Seroprevalence of <i>N. caninum</i> % (No. positive / No. sampled)	95% CI of <i>N. caninum</i> seroprevalence	Mixed seroprevalence % (No. positive / No. sampled)	95% CI of mixed seroprevalence
Central region	Tov	10.8 (7/65)	4.8-21.5	33.8 (22/65)	22.8-46.7	0.0 (0/65)	0.0-6.9
	Omnogobi	28.7 (29/101)	20.3-38.7	17.8 (18/101)	11.1-26.9	5.9 (6/101)	2.4-12.9
	Gobisumber	20 (4/20)	0.6-44.2	30 (6/20)	12.8-54.3	5.0 (1/20)	0.2-26.9
	Dornogobi	33.3 (14/42)	20.0-49.6	42.9 (18/42)	28.0-58.9	21.4 (9/42)	10.8-37.2
	Selenge	28.2 (20/71)	18.4-40.2	25.4 (18/71)	16.1-37.3	11.3 (8/71)	5.3-21.5
	Dundgobi	42.1 (16/38)	26.7-59.0	44.7 (17/38)	29-61.5	31.6 (12/38)	18.0-48.7
Western region	Gobi-Altai	8.9 (4/45)	2.8-22.1	13.3 (6/45)	5.5-27.4	0.0 (0/45)	0.0-9.0
	Khovd	5.7 (3/53)	1.4-16.6	32.1 (17/53)	20.3-46.4	0.0 (0/53)	0.0-8.0
	Bayan-Olgii	2.6 (4/153)	0.8-6.9	10.5 (16/153)	6.2-16.7	1.3 (2/153)	0.2-5.1
	Uvs	12.3 (8/65)	5.8-23.3	12.3 (8/65)	5.8-23.3	3.1 (2/65)	0.5-11.6
	Zavkhan	20.7 (17/82)	12.8-31.3	43.9 (36/82)	33.1-55.2	12.2 (10/82)	6.3-21.7
Eastern region	Sukhbaatar	17.6 (18/102)	11.0-26.7	17.6 (18/102)	11.0-26.7	3.9 (4/102)	1.2-10.3
	Dornod	17.5 (20/114)	11.2-26.0	34.2 (39/114)	25.7-43.7	4.4 (5/114)	1.6-10.4
	Khentii	15.2 (20/132)	9.7-22.6	18.2 (24/132)	12.2-26.0	4.5 (6/132)	1.8-10.0
Khangai region	Bulgan	23 (14/61)	13.5-35.8	29.5 (18/61)	18.8-42.7	4.9 (3/61)	1.2-14.6
	Khovsgol	20 (11/55)	10.8-33.3	20 (11/55)	10.8-33.3	9.1 (5/55)	3.3-20.7
	Orknon	10 (1/10)	0.5-45.8	20 (2/10)	3.5-55.7	0.0 (0/10)	0-34.4
	Ovorkhangai	41.7 (35/84)	31.1-52.9	38.1 (32/84)	27.9-49.3	21.4 (18/84)	13.5-32
	Bayankhongor	11.8 (9/76)	5.8-21.7	36.8 (28/76)	26.2-48.7	0.0 (0/76)	0.0-6.0
	Arkhangai	17.9 (10/56)	9.3-30.8	25 (14/56)	14.8-38.6	3.6 (2/56)	0.6-13.3
Capital city	Ulaanbaatar	38.5 (5/13)	15.1-64.4	69.2 (9/13)	38.8-89.6	38.5 (5/13)	15.1-67.7
	Total	18.7 (269/1,438)	16.7-20.8	26.2 (377/1,438)	23.9-28.5	6.8 (98/1438)	5.6-8.2

CI: confidence interval.

Table 2Seroprevalence of *T. gondii* in different regions of Mongolia

Regions of Mongolia	No. tested	No. positive	No. negative	Seroprevalence (%)	OR (95%CI)	<i>P</i> -value
Central region	350	95	255	27.1	0.26 (0.17-0.40)	<0.0001
Western region	398	36	362	9	-	
Eastern region	348	58	290	16.6	0.49 (0.31-0.77)	0.002
Khangai region	342	80	262	23.3	0.32 (0.1-0.49)	<0.0001

OR: odds ratio, CI: confidence interval, -: reference group.

Table 3Analysis of risk factors associated with *T. gondii* infection in Mongolian cattle

Risk factors		No. sampled	No. positive	No. negative	Seroprevalence (%)	OR(95%CI)	P-value
Age groups (Years)	1--4	710	135	575	19	-	
	5--9	430	75	355	17.4	1.11 (0.81-1.51)	0.507
	10--14	44	10	34	22.7	0.79 (0.38-1.65)	0.543
	Unknown	254	49	205	19.2	0.98 (0.68-1.41)	0.92
Sex	Female	1,174	213	961	18.1	-	
	Male	264	56	208	21.2	0.82 (0.59-1.14)	0.248

OR: odd ratio, CI: confidence interval, - : reference group.

Table 4Seroprevalence of *N. caninum* in different regions of Mongolia

Regions of Mongolia	No. tested	No. positive	No. negative	seroprevalence (%)	OR (95%CI)	<i>P</i> -value
Central region	350	108	242	30.8	0.59 (0.42-0.82)	0.002
Western region	398	83	315	20.8	-	
Eastern region	348	81	267	23.2	0.86 (0.61-1.22)	0.427
Khangai region	342	105	237	30.7	0.59 (0.42-0.83)	0.002

OR: odds ratio, CI: confidence interval, - : reference group.