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Abstract

Molecular and serological data suggest that noroviruses (NoVs) might be transmitted between humans and domestic carnivores. In this study we screened an age-stratified collection of canine sera (n=516) by using an ELISA assay based on virus-like particles (VLPs) of human NoVs GII.4 and GIV.1 and carnivore NoVs GIV.2 and GVI.2. Antibodies against GII.4 and GIV.1 human NoVs and GIV.2 and GVI.2 NoVs from carnivores were identified in dog sera (13.0%, 67/516) suggesting their exposure to homologous and heterologous NoVs. Analysis of the trends of age-class prevalence showed a gradual increase in the positive rate from 9.0% and 7.0%, in young dogs < 1 year of age to 15.0% in dogs older than 12 years, for GII.4 and GVI.2 NoVs, respectively. A significant difference in the IgG distribution by age classes was observed for GIV.1 NoVs, with the highest rate of antibodies (7.0%) in the age group < 1 year and the lowest (1.0%) in the age-classes 7-9 (P = 0.049). High correlation between the reactivity to GII.4 and GVI.2 NoVs was observed, likely due to conserved epitopes in the capsid structure.

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**To the Editor of
Veterinary Microbiology**

Teramo, 1-25-2017

Dear Editor,

several data suggest that circulation of noroviruses (NoVs) may occur between pets and humans raising concerns of potential cross-species transmission. In this study, antibodies against GII.4 and GIV.1 human NoVs and GIV.2 and GVI.2 NoVs from carnivores were identified in dog sera (13.0%, 67/516) suggesting their exposure to homologous and heterologous NoVs. I hope you will find the manuscript “Seroprevalence for norovirus genogroup II, IV and VI in dogs” by Di Martino et al., written in a format corresponding to a short communication of interest for the journal.

With the best regards,

Barbara Di Martino

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1 **Seroprevalence for norovirus genogroup II, IV and VI in dogs**

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26

27 **Abstract**

28 Molecular and serological data suggest that noroviruses (NoVs) might be transmitted between
29 humans and domestic carnivores. In this study we screened an age-stratified collection of canine
30 sera (n=516) by using an ELISA assay based on virus-like particles (VLPs) of human NoVs GII.4
31 and GIV.1 and carnivore NoVs GIV.2 and GVI.2. Antibodies against GII.4 and GIV.1 human NoVs
32 and GIV.2 and GVI.2 NoVs from carnivores were identified in dog sera (13.0%, 67/516) suggesting
33 their exposure to homologous and heterologous NoVs. Analysis of the trends of age-class
34 prevalence showed a gradual increase in the positive rate from 9.0% and 7.0%, in young dogs < 1
35 year of age to 15.0% in dogs older than 12 years, for GII.4 and GVI.2 NoVs, respectively. A
36 significant difference in the IgG distribution by age classes was observed for GIV.1 NoVs, with the
37 highest rate of antibodies (7.0%) in the age group < 1 year and the lowest (1.0%) in the age-classes
38 7-9 (P = 0.049). High correlation between the reactivity to GII.4 and GVI.2 NoVs was observed,
39 likely due to conserved epitopes in the capsid structure.

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48 **Keywords:** noroviruses (NoVs); genotypes GII.4, GIV.1, GIV.2 and GVI.2; dogs; antibodies.

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53 **Introduction**

54 Noroviruses (NoVs) are major human pathogens associated with acute gastroenteritis (Patel et al.,
55 2008). NoVs belong to the genus *Norovirus* in the family *Caliciviridae*. The genome of NoVs
56 consists of a single-stranded positive-sense RNA molecule of ~7.5kb that is organized into three
57 open reading frames (ORFs). ORF1 encodes a polyprotein that is co-translationally cleaved into
58 seven proteins required for replication, while ORF2 encodes the major capsid protein (VP1) and
59 ORF3, a minor capsid protein (VP2) (Green, 2013). Based on the full-length VP1 capsid protein,
60 the *Norovirus* genus comprises seven genogroups (G), which can be subdivided in at least 40
61 genotypes (Green, 2013; Vinje, 2015). Viruses belonging to GI, GII and GIV can infect humans,
62 with GII.4 strains that are the most prevalent worldwide (Green, 2013). NoVs (GIV.2, GVI.1,
63 GVI.2 and GVII) have been also identified in carnivores (Martella et al., 2007; Martella et al., 2008;
64 Martella et al., 2009; Mesquita et al., 2010; Pinto et al., 2012; Tse et al., 2012; Di Martino et al.,
65 2016). Some NoV strains from carnivores are genetically related to human GIV NoVs, suggesting
66 common pathways in their evolution (Martella et al., 2007).

67 Molecular and serological studies suggest that circulation of NoVs may occur between pets and
68 humans raising concerns of potential cross-species transmission. Human sera may contain specific
69 IgG antibodies against carnivore GIV.2 and GVI.2 NoVs (Mesquita et al., 2013; Di Martino et al.,
70 2014) and in turn, dog sera may contain antibodies against human NoVs of genogroup I and II
71 (Caddy et al., 2015). In a study in Mexico, having dogs in or near home was recognised as a risk
72 factor for acquisition of IgA antibodies specific for NoVs in infants (Peasey et al., 2004). Also,
73 partial capsid sequences of human GII.4 and GII.12 NoVs have been detected in household dogs in
74 contact with human patients affected by NoV gastroenteritis (Summa et al., 2012).

75 In order to draw a more complete picture of NoV epidemiology in dogs, we screened an age-
76 stratified collection of canine sera by using an enzyme-linked immunosorbent assay (ELISA) based
77 on virus-like particles (VLPs) generated from human NoVs of genotype GII.4 and GIV.1 and from
78 carnivore NoVs of genotype GIV.2 and GVI.2.

79 **Materials and methods**

80 *Serum sample collection*

81 A total of 516 serum samples were collected between March 2013 and July 2015 by a convenience
82 sampling of household dogs admitted to veterinary clinics from different Italian regions. The
83 samples were divided on the basis of age groups: <1 year, 1–3 years, 3-year age groups from 4 to
84 12, and >12 years of age.

85

86 *Virus-like Particles (VLPs)*

87 The recombinant baculoviruses carrying the genes for the viral capsid proteins of the
88 Hu/NoV/GII.4/MD14512/1987/US, Hu/NoV/GIV.1/SaintCloud/624/1998/US,
89 Lion/NoV/GIV.2/Pistoia/387/06/ITA and Dog/NoV/GVI.2/FD53/2007/ITA were obtained as
90 previously described (Bok et al., 2009; Di Martino et al., 2010; Di Martino et al., 2015). For large-
91 scale production of VLPs, 100 ml of *Spodoptera frugiperda* (*Sf9*) cells (1×10^6 cell/ml) suspension
92 culture were infected with the recombinant baculovirus at a multiplicity of infection of three plaque
93 forming units/cell. The recombinant capsid proteins were concentrated by ultracentrifugation
94 through a 17% sucrose cushion in TEN-buffer (100 mM NaCl; 50 mM Tris-HCl, pH 7.5; 1 mM
95 EDTA) and purified on a discontinuous 20 to 60% (wt/vol) sucrose gradient. The collected fractions
96 were dialyzed against PBS, and the protein concentration of VLP preparations was determined by
97 measuring the optical density at 280 nm (OD_{280}) and visually by running aliquots on SDS-10%
98 PAGE containing bovine serum albumin (BSA) standards. The presence of VLPs was confirmed by
99 western blotting (WB) and electron microscopy, as previously described (Bok et al., 2009; Di
100 Martino et al., 2010; Di Martino et al., 2014).

101

102 *Enzyme-linked immunoassay (ELISA)*

103 For the development of the antibody-detection ELISA, the supernatant of mock infected cells,
104 GII.4, GIV.1, GIV.2 and GVI.2 VLPs were diluted to a final concentration of 4 μ g/ml in carbonate-

105 bicarbonate buffer (0.05 M, pH 9.6) and 100 μ l of each antigen were added to the well of a 96-well
106 EIA plate (Costar, Italy). The wells were washed five times with 0.1% Tween-PBS (PBS-T) and
107 then blocked with 200 μ l of PBS containing 2% BSA at room temperature for two hours. Each
108 serum sample was tested at the initial dilution of 1:100 and the plates were incubated at 37°C for 1
109 hour. After incubation with horseradish peroxidase-conjugated goat anti-dog immunoglobulin G
110 (IgG) (Sigma-Aldrich, Italy) at dilution of 1:5,000 for 30 min at 37°C, the reaction was developed
111 with the addition of 100 μ l per well of 2,2'-azino-di-(3-ethylbenzthiazoline-6-sulfonate) (ABTS)
112 substrate. The cut-off point of the ELISA test was established as the mean of the OD₄₀₅ readings of
113 50 dog sera negative in WB for each NoV antigen plus 2 standard deviations. For each tested
114 sample a positive/negative ratio (OD₄₀₅ of VLPs/OD₄₀₅ of mock infected cells) \geq 2.0 was used to
115 evaluate the background binding. All the sera with an OD₄₀₅ values \geq 0.5 at the initial dilution of
116 1:100 were considered positive and titrated in twofold dilutions. Mean ELISA antibody titres were
117 calculated and expressed as the reciprocal of the highest serum dilution with a positive absorbance
118 (OD₄₀₅ \geq 0.5) for each NoV antigen.

119

120 *Statistical analysis*

121 The data were analysed using Prism Graphpad Software. Fisher's exact test was used to determine
122 the differences in seroprevalence among the age groups. Pearson's chi-squared test was applied to
123 assess the correlation among NoV genotype serum titers detected in the tested dogs. A P value of
124 <0.05 was considered statistically significant.

125

126 *Evaluation of serological cross-reactivity between GII.4 and GVI.2 NoVs*

127 In order to assess the antigenic relationships between GII.4 and GVI.2 VLPs, we performed
128 blocking assay experiments. Briefly, 6 dog sera positive for both the antigens with titres $>1:800$
129 were pre-incubated with optimized concentrations of GII.4 and GVI.2 VLPs (2, 4, 8, 16 μ g/ml) for
130 1 h at 37 °C. After incubation, the sera were tested in ELISA for the presence of GII.4 and GVI.2

131 antibodies, starting at dilution of 1:100. The possible inter-genogroup serological cross reactivity
132 between GII.4 and GVI.2 VLPs was also investigated in WB analysis, by assessing the reactivity of
133 five human sera positive for GII.4 with titres of 1:200 (Di Martino et al., 2014) against GVI.2
134 VLPs.

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157 **Results and discussion**

158 Out of 516 dog sera, 13.0% (67/516) reacted with at least one NoV antigen. In detail, 52 sera
159 reacted with the GII.4 VLPs with a prevalence rate of 10.1%, at dilutions ranging from 1:100 to
160 1:3,200. A total of 23 (4.5%) samples reacted with GIV antigens, with titres ranging from 1:100 to
161 1:800. Of these, 17 (3.3%) resulted positive for both GIV.1 and GIV.2, while 3 (0.58%) sera reacted
162 only with GIV.1 at dilutions from 1:200 to 1:800 and an additional 3 (0.58%) samples reacted only
163 with GIV.2, at final dilution of 1:100. Forty-six (8.9%) sera reacted with GVI.2 VLPs, at dilutions
164 ranging from 1:100 to 1:3,200 (Table 1).

165 Analysis of the trends of age-class prevalence by Fisher's exact test showed a gradual increase in
166 the positive rate from 9.0% and 7.0% in young dogs <1 year of age to 15.0% in dogs older than 12
167 years for GII.4 and GVI.2 NoVs, respectively. A significant difference in the IgG distribution by
168 age-classes was observed for GIV.1, with the highest rate of antibodies (7.0%) in the group <1 year
169 and the lowest (1.0%) in the age-class 7-9 ($P = 0.049$). A similar age-related pattern was also found
170 for GIV.2 (Fig. 1).

171 The results obtained are consistent with the hypothesis that dogs may be exposed to both human
172 and carnivore NoVs throughout their life. The magnitude of GII.4 (10.1%) circulation in household
173 dogs was higher in our serological survey than in a previous investigation conducted in UK, where
174 4.9% and 8.8% of kennel and household dogs, respectively, possessed antibodies for NoVs using a
175 pool of GII.3, GII.4, GII.6 and GII.12 VLPs (Caddy et al., 2015). By converse, the overall
176 prevalence of antibodies against carnivore GIV.2 (3.9%) was lower than that reported (56.6%) in
177 UK (Caddy et al., 2013), but similar to the rate (4.8%) previously reported in an Italian dog
178 population (Di Martino et al., 2010). Furthermore, in our study IgG antibodies reacting only against
179 human GIV.1 VLPs were found in dog sera, suggesting exposure to human GIV NoVs. In general,
180 the seroprevalence for carnivore GVI.2 NoVs revealed in our investigation was lower than the
181 prevalences reported in previous studies (Caddy et al., 2013; Mesquita et al., 2014). In UK (Caddy

182 et al., 2013) the seropositivity for GVI.2 VLPs was 15.9%, while in a multi-centric study conducted
183 in 14 European countries the overall rate was 36.0% (Mesquita et al., 2014).

184 In our analysis, the majority of the NoV-seropositive dogs showed reactivity for multiple genogroup
185 antigens. Pearson's rank analysis revealed a high correlation between the levels of antibodies to
186 GIV.1 and GIV.2 antigens (Fig. 2a), while a weak correlation was found when comparing the
187 antibody titers to GIV.1 and to GIV.2 antigens with the antibody titers to GII.4 and to GVI.2 VLPs,
188 respectively (Fig. 2b, 2c, 2d, 2e). Of interest, we found a high correlation between the level of
189 antibodies to GII.4 and GVI.2 VLPs (Fig. 2f), in particular in the dog sera that strongly reacted
190 against both antigens (from >1:800 to 1:3,200). Previous evidence indicates that GII.4 VLPs are
191 antigenically unrelated to GIV NoVs, whilst antigenic cross-reactivity has been observed between
192 GIV.1 and GIV.2 genotypes (Di Martino et al., 2014).

193 The antigenic relationships between GII.4 and GVI.2 VLPs was investigated by blocking assay
194 experiments. In our analysis, we found for all the six sera examined that at concentrations higher
195 than 8 µg of each antigen per ml, binding of GVI.2 antibodies was blocked (~50% reduction of
196 OD₄₀₅ values) by GII.4 VLPs (Fig. 3a), and in turn binding of GII.4 antibodies was blocked by
197 GVI.2 VLPs, although with a ~25% reduction of the OD₄₀₅ values (Fig. 3b). Also, by testing in WB
198 analysis five human sera positive for GII.4 with titres of 1:200, we found reactivity against GVI.2 at
199 the initial dilution of 1:100 (Fig. S1). Overall, these findings suggest a strong serologic cross-
200 reactivity between human GII.4 and carnivores GVI.2 NoVs, likely due to the existence of
201 conserved epitopes in the capsid VP1 protein (Parra et al., 2012).

202 In conclusion, screening of canine sera with multiple NoV antigens demonstrated that dogs are
203 exposed to both human (heterologous) and animal (homologous) NoV strains. Different age-related
204 patterns were observed between the antibody prevalence to GII/GVI and to GIV antigens.
205 Understanding the ecology and dynamics of transmission of NoVs in carnivores will be helpful to
206 assess more precisely if and to which extent pets may pose a risk of infection by homologous and
207 heterologous NoV strains for humans.

208 **Acknowledgements**

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210 Ministry of University and Research.

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312 **Figure captions**

313

314 **Fig. 1** - IgG antibodies to GII.4, GIV.1, GIV.2 and GVI.2 VLPs in dog sera of different age groups.

315

316 **Fig. 2** - Pearson's rank analysis of the levels of antibodies among GII.4, GIV.1, GIV.2 and GVI.2
317 VLPs. Each scatterplot shows the correlation between genotype serum titers detected in the tested
318 dogs (a: GIV.1/GIV.2; b: GII.4/GIV.1; c: GII.4/GIV.2; d: GVI.2/GIV.1; e: GVI.2/GIV.2; f:
319 GII.4/GVI.2). All Pearson ranking values were statistically significant with a $p < 0.0001$.

320

321 **Fig. 3** – Blocking assay experiment. A canine serum positive for GII.4 and GVI.2 antigens with
322 titres $\geq 1:800$ was pre-incubated with GII.4 (**a**) and GVI.2 (**b**) VLPs at concentrations of 2, 4, 8 and
323 16 $\mu\text{g/ml}$ and tested in ELISA for the presence of antibodies against both VLPs.

324

325 **Fig. S1** - WB analysis of GII.4 and GVI.2 VP1 with a human serum sample positive in ELISA for
326 GII.4 NoVs. Line 1: Precision Plus protein Standards (Bio-Rad, Italy); line 2: GII.4 VP1; line 3:
327 GVI.2 VP1. Serum dilution: 1:100.

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338 **Conflict of interest statement**

339 All Authors declare that there are no financial or other relationships that might lead to a conflict of
340 interest. All authors have seen and approved the manuscript and have contributed significantly to
341 the work.

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Fig. 1

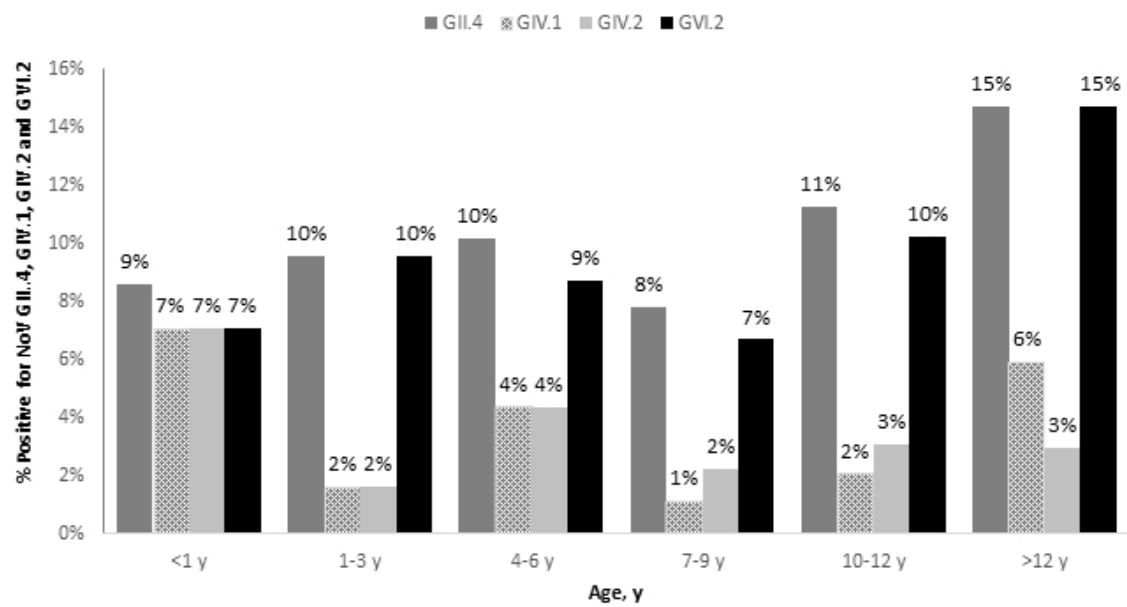


Fig. 2

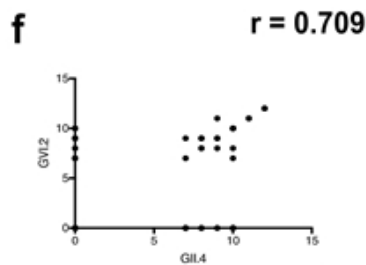
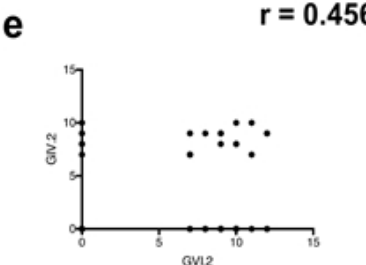
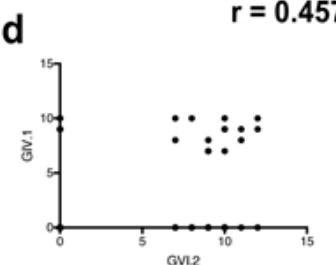
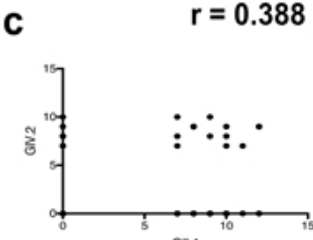
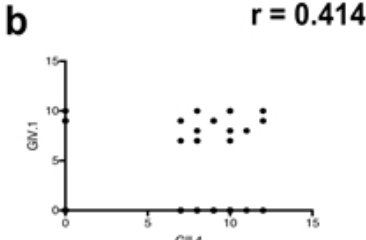
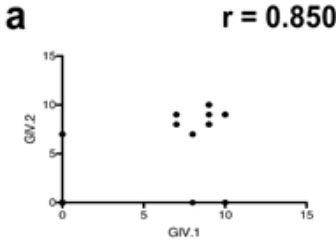


Fig. 3

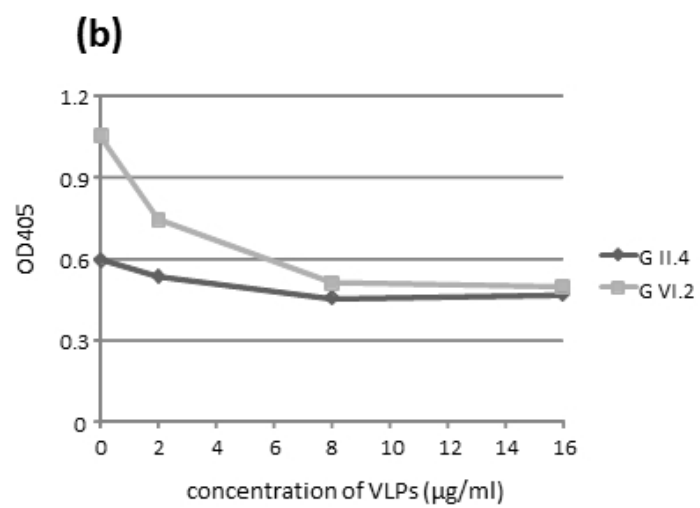
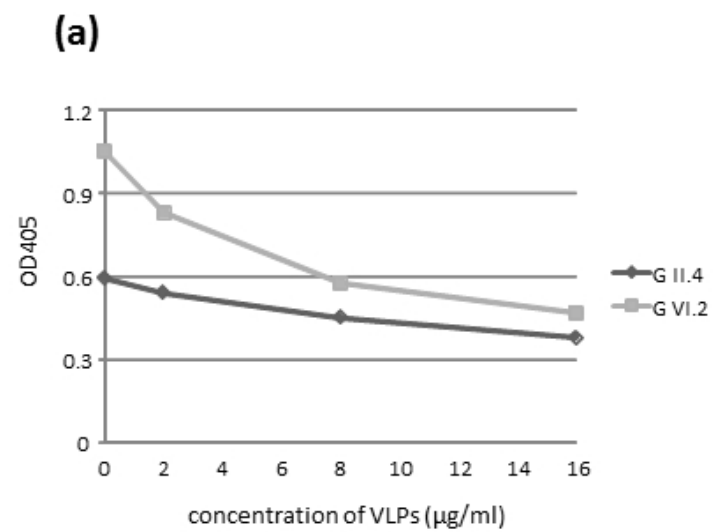


Fig. S1

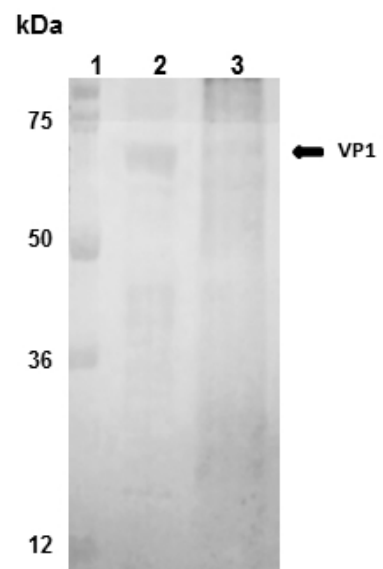


Table 1

IgG antibody titres to GII.4, GIV.1, GIV.2 and GVI.2 VLPs in dog sera. Mean ELISA antibody titres were calculated and expressed as the reciprocal of the highest serum dilution with a positive absorbance ($OD_{405} \geq 0.5$) for GII.4, GIV.1, GIV.2 and GVI.2 VLPs.

| NoV† VLPs* | Serum dilutions | | | | | | Total (%) |
|---------------|-----------------|--------------|---------------|---------------|--------------|--------------|----------------|
| | 100 (%) | 200 (%) | 400 (%) | 800 (%) | 1,600 (%) | 3,200 (%) | |
| GII.4 | 7/516 (1.4%) | 9/516 (1.7%) | 7/516 (1.4%) | 22/516 (4.3%) | 3/516 (0.6%) | 4/516 (0.8%) | 52/516 (10.1%) |
| GIV.1 | 4/516 (0.8%) | 3/516 (0.6%) | 7/516 (1.4%) | 6/516 (1.2%) | 0/516 (0%) | 0/516 (0%) | 20/516 (3.9%) |
| GIV.2 | 5/516 (1.0%) | 5/516 (1.0%) | 7/516 (1.4%) | 3/516 (0.6%) | 0/516 (0%) | 0/516 (0%) | 20/516 (3.9%) |
| GVI.2 | 5/516 (1.0%) | 5/516 (1.0%) | 12/516 (2.3%) | 16/516 (3.1%) | 4/516 (0.8%) | 4/516 (0.8%) | 46/516 (8.9%) |

†NoV, norovirus

*VLPs, virus-like particles

Highlights

- Noroviruses (NoVs) are a major cause of epidemic gastroenteritis in humans.
- We detected IgG antibodies against human and carnivore NoVs in household dog sera.
- This finding indicates that dogs are exposed to homologous and heterologous NoVs.