

ACCEPTED MANUSCRIPT

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34 **Serological diagnosis of bovine neosporosis: a comparative study of commercially**
35 **available ELISA tests**

36

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65 **Abstract**

66 Bovine neosporosis control programs are currently based on herd management
67 and serodiagnosis because effective treatments and vaccines are unavailable. Although a
68 wide variety of serological tools have been developed, enzyme-linked immunosorbent
69 assays (ELISAs) are the most commonly commercialized tests. Partial comparative
70 studies have been performed in the past, and the panel of available ELISAs has notably
71 changed in the last few years. Therefore, diagnostic laboratories are requesting updated
72 information about the performance of these tests.

73 Accordingly, the aim of this study was to compare all of the commercially
74 available ELISAs (n=10) by evaluating their performance and to re-standardize them
75 based on TG-ROC analyses when necessary. For this purpose, a well-characterized
76 serum panel from experimentally and naturally infected bovines and non-infected
77 bovines (n=458) was used. Two different definitions of gold standard were considered:
78 (i) the result of the majority of tests and (ii) pre-test information based on
79 epidemiological, clinical and serological data. Most of the tests displayed high
80 sensitivity (Se) and specificity (Sp) values when both gold standard criteria were
81 considered. Furthermore, all the tests showed near perfect agreement, with the exception
82 of the pair-wise comparisons that included the VMRD and SVANOVIR. The best-
83 adjusted ELISAs were the HIPRA-CIVTEST, IDVET, BIOVET and IDEXX Rum (Se
84 and Sp > 95%). After the TG-ROC analyses, higher Se and Sp values were obtained for
85 the BIO-X, LSI Bov, LSI Rum and IDEXX Bov, though the increases were more
86 significant for the SVANOVIR and VMRD. The Kappa values also increased with the
87 new adjusted cut-offs. This is the first study that offers updated performance evaluations
88 of commercially available ELISAs. Such analyses are essential for diagnostic
89 laboratories and are valuable to the companies that develop and distribute these tests.

90

91 **Key words:** Bovine neosporosis, commercial ELISAs, comparative study, diagnostic
92 performance, agreement, TG-ROC

93

94

95 **1. Introduction**

96

97 Bovine neosporosis is a parasitic disease caused by the cyst-forming coccidian
98 parasite *Neospora caninum* which causes abortion and neonatal mortality in cattle
99 worldwide and, consequently, significant economic losses in the cattle industry
100 (reviewed by Dubey et al., 2007; Reichel et al., 2013).

101 There are numerous studies confirming the importance of bovine neosporosis. A
102 multinational study (Bartels et al., 2006) and another recent study carried out in Spain
103 (Eiras et al., 2011) updated the prevalence rates of *N. caninum* infection in several
104 European countries reporting herd prevalence rates from 16% in Sweden to 80% in
105 Spain.

106 At present, there is no effective treatment or vaccine for *N. caninum* infection,
107 and control measures are based on herd management and diagnosis. The serological
108 diagnosis of neosporosis in adult cattle and precolostral calves is an integral part of
109 control programs because the other commonly adopted control measures include the
110 selective culling of seropositive *Neospora*-associated aborted dams, herd replacement
111 with seronegative cattle and testing during the quarantine period prior to herd entry
112 (Dubey et al., 2007).

113 Serological techniques are primarily employed to detect specific antibodies
114 against *N. caninum* to differentiate infected from non-infected animals. These
115 techniques include a wide variety of enzyme-linked immunosorbent assays (ELISAs)
116 (in-house and commercial tests), indirect fluorescent antibody tests (IFATs) and a *N.*
117 *caninum*-agglutination test (NAT) (reviewed by Ortega-Mora et al., 2007). In addition,
118 western blotting is often recommended to confirm uncertain results in valuable samples
119 (Álvarez-García et al., 2002). Moreover, avidity tests are useful for investigating the
120 route of *N. caninum* transmission in herds because they can differentiate between
121 primary and chronic infections (Bjorkman et al., 2006; Aguado-Martínez et al., 2008).

122 In the last few years, the panel of commercially available serological kits has
123 notably changed; new tests have been developed, several have been modified and other
124 tests are no longer commercialized. Moreover, other in-house developed tests have been
125 recommended for the diagnosis of bovine neosporosis (eg. Osawa et al., 1998; Wouda
126 et al., 1998). Thus, at present, there is a paucity of up-to-date information about the
127 performance of these diagnostic products, which is essential information for diagnostic

128 labs. In fact, the last comparative studies offered a fragmented picture of the diagnostic
129 tools employed in Europe (von Blumröder et al., 2004) and the USA (Wapenaar et al.,
130 2007) and it remains unclear if the current serological tests offer standardized
131 interpretation of results.

132 Therefore, the aim of this study was to evaluate the performance of and re-
133 standardize the commercially available ELISA tests to detect anti-*N. caninum*
134 antibodies. Thus a sera panel that reflects an appropriate spectrum of disease was tested
135 following the recommended procedure for validation of diagnostic tests (Jacobson,
136 1998; Greiner and Gardner, 2000; Gardner et al., 2010).

137

138 **2. Materials and Methods**

139

140 *2.1. Sera and experimental design*

141 A well-defined bovine sera panel was analyzed by ten commercial ELISA tests.
142 This coded panel was composed of 458 bovine serum samples from both experimentally
143 and naturally infected cattle (including aborted and non-aborted dams) as well as non-
144 infected cattle. All the sampled animals were older than 6 months to avoid the presence
145 of colostral antibodies.

146 The animals were categorized into the following groups:

147

148 *2.1.1. Sera from non-infected cattle (Group a; n=125)*

149 Heifers and cows from a dairy herd (Holstein Friesian breed) without a previous
150 history of *N. caninum*-associated abortions tested negative in three consecutive
151 samplings during 6-9 month intervals throughout a period of two years in order to
152 discard antibody fluctuations below the cut-off value that may occur in chronically
153 infected cattle. All the samples were negative using an in house *N. caninum* soluble
154 extract antigen-based ELISA (Álvarez-García et al., 2003; Aguado-Martínez et al.,
155 2008) that discriminates between positive and negative results. Moreover, the samples
156 also tested negative using two recombinant antigen-based ELISAs (rNcGRA7 and
157 rNcSAG4 ELISAs). The NcGRA7 protein is shared by both tachyzoite and bradyzoite
158 stages, whereas NcSAG4 is the first bradyzoite stage-specific protein described. The
159 usefulness of rNcGRA7 and rNcSAG4 ELISAs to detect acute and chronic cattle

160 infections, respectively, that may go undetected by conventional serological tools was
161 previously reported (Aguado-Martínez et al., 2008).

162

163 *2.1.2. Sera from N. caninum naturally infected cattle (Group b; n=169)*

164

165 *Serum samples from seropositive non-aborted dams:*

166 Serum samples (n=136) were collected from dams from a dairy herd (Holstein
167 Friesian breed) with a history of *N. caninum*-associated abortions. The herd comprised
168 200 cows and had an intra-herd seroprevalence of 85% and a 9.2% annual abortion rate.
169 The *N. caninum*-associated abortions were diagnosed using histopathology and PCR.
170 Moreover, an endogenous transplacental transmission rate of 80% was estimated based
171 on an equal distribution of seropositive animals across the different age groups and a
172 significant association between seropositive dams and their daughters. One hundred and
173 thirty serum samples came from mother-daughter pairs (n=65 pairs), and 6 serum
174 samples came from only mothers or daughters. All the samples tested positive using an
175 in house *N. caninum* soluble extract antigen-based ELISA.

176

177 *Serum samples from seropositive aborted dams and/or dams at risk of abortion:*

178 *Sera from a herd with an endemic pattern of N. caninum-associated abortions*

179 The sera were collected from aborted cows (n=21) belonging to the previously
180 mentioned herd. The samples were collected 2 months prior to or after the abortion
181 when an increase in specific antibody levels is expected (Quintanilla-Gozalo et al.,
182 2000). All the samples tested positive using an in house *N. caninum* soluble extract
183 antigen-based ELISA.

184

185 *Sera from a herd with an epidemic pattern of N. caninum-associated abortions*

186 The sera were collected from animals from a dairy herd (Holstein Friesian
187 breed) of 1080 cows, which did not have a history of reproductive failure and had a
188 *N. caninum* seroprevalence of less than 5% prior to an abortion storm. During this event all
189 abortions were concentrated in a two month period, and the abortion rate during this
190 period was 5.1%. In addition, 81.82% of the abortions occurred in seropositive animals
191 that displayed low avidity anti-*N. caninum* antibodies (Rojo-Montejo et al., 2009). The
192 intra-herd seroprevalence increased to 19.5% after the abortion storm. The tested

193 samples belonged to the seropositive aborted cows or the cows at risk of abortion during
194 the abortion storm (n=12). These animals were located in the same yard according to the
195 gestation period and lactating age.

196

197 2.1.3. Sera from *N. caninum* experimentally infected cattle (Group c; n=150)

198 All experimentally infected animals were seronegative prior to inoculation using
199 an in house *N. caninum* soluble extract antigen-based ELISA and rNcGRA7 and
200 rNcSAG4 ELISAs.

201

202 *Bulls*

203 Three bulls of the Asturiana de los Valles breed were intravenously infected
204 (i.v.) with 10^8 live *N. caninum* tachyzoites of the Nc-1 isolate. Sequential serum
205 samples were collected weekly and fortnightly for approximately 8 months (Serrano-
206 Martínez et al., 2007). Later, these samples were assayed using an avidity ELISA and an
207 ELISA based on the recombinant proteins NcGRA7 and NcSAG4. The results were
208 representative of primary infection until 8–10 weeks p.i. (low IgG avidity values) and
209 chronic infection from 10 weeks p.i. to the end of the experiment (high IgG avidity
210 values). Similar antibody kinetics were observed during primary and chronic infections
211 with the rNcGRA7 and *N. caninum* soluble extract antigen-based ELISAs, but the
212 experimentally infected bovines did not show specific anti-rNcSAG4 antibodies with
213 the rNcSAG4 ELISA. The most feasible explanation for the absence of specific anti-
214 rNcSAG4 antibodies in the study was a low efficiency of tachyzoite–bradyzoite
215 conversion in an experimental bovine model of neosporosis (Aguado-Martínez et al.,
216 2008). For the present study, a total of 55 serum samples were selected (n=24 samples
217 for the first bull, n=11 for the second bull and n=20 for the third bull), and they were
218 collected at 24 different time points throughout the 270 days of the experiment (Fig. 2).
219 These sera were representative of primary and chronic infections.

220

221 *Heifers*

222 Three heifers of less than 24 months old were inoculated i.v. with 10^7 live *N.*
223 *caninum* tachyzoites of the Nc-1 isolate at 70 days of gestation. Sequential serum
224 samples were collected fortnightly prior to and after the inoculation and until the
225 delivery or abortion (Rojo-Montejo et al., under review). For the present study, a total
226 of 18 sera were selected at 7 different time points throughout the 87-day experiment

227 (n=5 sera samples for the first heifer, n=7 sera samples for the second heifer and n=6
228 sera samples for the third heifer). Additionally, three heifers of less than 24 months old
229 were i.v. inoculated with 4×10^8 live *N. caninum* tachyzoites of the Nc-1 isolate at 135
230 days of gestation. Sequential serum samples were collected fortnightly prior to the date
231 of inoculation and after the inoculation up until the delivery or abortion (Rojo-Montejo
232 et al., 2013 submitted). Of these samples, a total of 44 were selected at 16 different time
233 points throughout the experiment (n=16 sera samples for the first heifer, n=13 sera
234 samples for the second heifer and n=15 sera samples for the third heifer). Moreover,
235 three additional non-infected heifers were employed as negative controls (inoculated
236 with PBS) (n=14 sera samples from the first negative control heifer, n=9 sera samples
237 for the second negative control heifer and n=10 sera samples for the third negative
238 control heifer). All these samples were assayed using an in house *N. caninum* soluble
239 extract antigen-based ELISA.

240 In all animals from group “c” all samples remained seropositive (between 2-3
241 weeks post-infection) once they had seroconverted.

242

243 *2.1.4. Sera from animals infected with closely related apicomplexan parasites (Group*
244 *d; n=14)*

245 Nine sera from seropositive cows of Brown Swiss breed naturally infected with
246 *Besnoitia besnoiti* were analyzed. *B. besnoiti* infection was confirmed using an in-house
247 ELISA developed by the SALUVET group (García-Lunar et al., 2013). Five sera from
248 heifers of Holstein Friesian breed with natural *Sarcocystis* spp. infections also were
249 analyzed. *Sarcocystis* infection was detected using the visualization of tissue cysts in the
250 heart via histological examination and *B. besnoiti* infection was discarded by an in
251 house ELISA mentioned above. All samples tested negative using an in house *N.*
252 *caninum* soluble extract antigen-based ELISA. These sera were included in the
253 experiment to study cross-reactivity with other apicomplexan parasites.

254

255 *2.2. Tests*

256 The samples were analyzed using nine commercial indirect enzyme-linked
257 immunosorbent assays (iELISA) and one commercial competitive enzyme-linked
258 immunosorbent assay (cELISA) (Table 1). The tests were performed, and the cut-off
259 values were applied according to the manufacturer’s instructions.

260

261 2.3. Analysis of data

262 Sensitivity (Se), specificity (Sp) and test agreement (expressed as Kappa-values;
263 κ), including 95% confidence intervals (95% CI), were calculated using WinEpiscope
264 2.0 (<http://www.clive.ed.ac.uk>).

265 Two different definitions of a gold standard were used to calculate the diagnostic
266 characteristics of the tests because a perfect reference test is not available for the
267 diagnosis of bovine neosporosis (Ortega-Mora et al., 2007).

268 The first gold standard was defined by the result of the majority of the tests
269 (‘Majority of tests’). If equal numbers of tests returned positive and negative results, the
270 sample was regarded as doubtful and was discarded.

271 The second gold standard was defined according to the pre-test information
272 (‘Pre-test information’). A sample was considered positive or negative based on
273 epidemiological (previous history of endemic or epidemic *N. caninum*-associated
274 abortions), clinical (aborting or non-aborting cattle) and serological data (seropositive or
275 seronegative using one or two reference tests: *N. caninum* soluble extract antigen-based
276 ELISA and recombinant protein-based ELISAs). Groups “a” and “d” were regarded as
277 negative reference sera as compared to groups “b” and “c”, which were regarded as
278 positive reference sera. Results of the tests were evaluated blinded.

279 TG-ROC analyses were carried out with respect to the gold standard ‘Majority
280 of tests’ (Greiner, 1995), and SPSS 17.0 for Windows (SPSS Inc.) was used. In
281 addition, the Student’s t-test was employed to compare specific antibody levels between
282 serum samples from seropositive aborted dams from a herd with an endemic pattern of
283 *N. caninum*-associated abortions (n=21) and samples from seropositive non-aborted
284 dams (n=136). Both groups were described above (group b: sera from *N. caninum*
285 naturally infected cattle) for each ELISA test. Moreover, the repeated measures
286 ANOVA with Tukey’s multiple comparison test was used for the comparison of
287 antibody levels at different days p.i. in the serum of experimentally infected animals for
288 each ELISA test. A *P*-value of less than 0.05 was considered statistically significant.
289 These statistical analyses were carried out using GraphPad Prism 5 v.5.01 (San Diego,
290 CA, USA) software.

291

292

293 **3. Results**

294

295 *3.1. Sensitivity (Se) and specificity (Sp) of tests according to the 'Majority of tests' and*
296 *'Pre-test information' gold standards*

297 Se and Sp values were calculated for each ELISA based on the cut-offs
298 recommended by each laboratory (Table 2). When “Majority of tests” was regarded as
299 the gold standard only one sample was discarded since equal numbers of tests returned
300 positive and negative results. Irrespective of the chosen gold standard, the Se was high
301 (>95%) for most tests except for the SVANOVIR test (85.9% relative to the gold
302 standard “Majority” and 87.2% relative to the gold standard “Pre-test info”). All the
303 tests showed high Sp values (93–100%) except for the VMRD test (65.1% relative to
304 the gold standard “Majority of tests” and 66.5% relative to the gold standard “Pre-test
305 info”).

306

307 *3.2. TG-ROC analysis*

308 TG-ROC analysis, based on the ‘Majority of tests’, was conducted to confirm
309 the accuracy of the suggested cut-off values. These analyses were conducted for the
310 ELISA tests that showed Se and/or Sp values of less than 95%. Thus, the cut-offs were
311 recalculated for the LSI Bov, LSI Rum, BIO-X, VRMD, IDEXX Bov and SVANOVIR
312 ELISAs (Table 2, Fig. 1).

313 According to the Sp and Se values, the tests that were notably modified after
314 TG-ROC analysis were the VRMD and SVANOVIR ELISAs. In the case of the
315 SVANOVIR ELISA, the new suggested cut-off was percent positivity (PP) ≥ 15 for Se
316 and Sp values of 94.0% and 93.7%, respectively. In the case of the VMRD ELISA, the
317 new suggested cut-off was percent inhibition (% I) ≥ 65 for Se and Sp values of 94.8%
318 and 91.6%, respectively (Table 3).

319 This study also permitted slight readjustments of other tests, including the LSI
320 Bov ELISA (readjusted cut-off ≥ 52 ; Se=98.9% and Sp=97.9%), LSI Rum ELISA
321 (readjusted cut-off ≥ 53 ; Se=99.3% and Sp=97.4%), BIO-X ELISA (readjusted cut-off
322 $\geq 23/13$; Se=98.1% and Sp=96.2%) and IDEXX Bov ELISA (readjusted cut-off ≥ 1 ;
323 Se=98.5% and Sp=97.9%) (Table 3).

324 TG-ROC analysis was not performed for the HIPRA-CIVTEST, IDVET,
325 IDEXX Rum and BIOVET ELISAs because these tests showed Se and Sp values higher
326 than 95%.

327 Interestingly, all the ELISAs that employed cut-off values with a range of
328 doubtful results yielded a low percentage of doubtful results (3.2% for HIPRA-
329 CIVTEST, 2.1% for IDVET, 3.9% for BIO-X and 2.4% for IDEXX Rum).

330

331 3.3. Test agreement (K-statistics)

332 First, K-values were calculated between ELISAs and both gold-standard criteria
333 prior to and after TG-ROC analyses. HIPRA-CIVTEST, IDVET, IDEXX Rum and
334 BIOVET ELISA showed in both cases perfect agreement (K-values higher than 0.95).
335 K-values of LSI Bov, LSI Rum, BIO-X, IDDEX-Bov and SVANOVIR ELISAs were
336 close to or higher than 0.90 and remained similar or slightly increased after TG-ROC re-
337 adjustment of cut-offs. The lowest K-value corresponded to VMRD ELISA, which
338 significantly increased after TG-ROC analysis (Table 4).

339 When all ELISAs were compared to each other all the tests showed near perfect
340 agreement (K = 0.8–0.9) with the exception of the pair-wise comparisons, which
341 included the VMRD and SVANOVIR ELISAs (Supplementary Table 1).

342 The K-values were recalculated using the adjusted cut-offs obtained using TG-
343 ROC analysis based on the gold standard ‘Majority of tests’ (Supplementary Table 2).

344 As expected, there was a substantial increment of K-values in the pair-wise
345 comparisons, including the VMRD and SVANOVIR ELISAs, reaching K-values
346 between 0.8 and 0.9 (Supplementary Table 2).

347

348 3.4. Cross-reactions

349 Nine of the 14 sera that were positive against *Sarcocystis* spp. and *Besnoitia*
350 *besnoiti* infections were positive according to four of the 10 evaluated ELISAs. The
351 BIO-X test showed the highest number of cross-reactions (4/14), and all were with sera
352 positive against *B. besnoiti*. The VMRD ELISA showed cross-reactions (3/14) to
353 *Sarcocystis* spp. positive sera, and the LSI Bov and BIOVET ELISAs only yielded one
354 false positive result with *B. besnoiti*-positive serum.

355

356

357

358 *3.5. Antibody titers in aborted vs. non-aborted cattle using Student's t-test analysis*

359 Student's t-test comparisons showed significant differences between aborted and
360 non-aborted dams for the HIPRA-CIVTEST ($P= 0.0055$), Bio-X ($P= 0.0023$) and
361 BIOVET ($P= 0.0012$).

362

363 *3.6. Kinetics of antibody responses using ANOVA analysis*

364 Antibody kinetics were examined in experimentally infected bulls and cows
365 (Fig. 2) throughout the sampling period. The samples included in the statistical analyses
366 comprised 27 sera samples from 3 experimentally infected bulls collected at 9 different
367 sampling points post infection (1, 2, 3, 4, 5, 6, 7, 8 and 9 weeks post-infection) and 33
368 sera samples of 3 experimentally infected cows at 135 days of gestation collected at 11
369 different sampling points post infection (2, 3, 4, 6, 7, 9, 11, 13, 15, 17 and 19 wpi). Both
370 the groups were described above in detail (section 2.1 section, sera and experimental
371 design; group c). In summary, all the kits behaved similarly.

372 All the sera obtained from experimentally infected bulls were negative at 1 wpi
373 irrespective of the kit employed. Seroconversions were mostly detected at 2 wpi, and all
374 the animals remained positive from 3 wpi onwards. Antibody levels peaked at 4wpi or
375 even later depending on the test. Seroconversions (values above the cut-off) were
376 detected in all the animals at 2 wpi with the BIO-X and IDVET tests, whereas the
377 IDEXX Bov and VMRD displayed positive and negative results at 2 wpi.
378 Seroconversions were observed at 3 wpi for the CIVTEST, LSI Bov, LSI Rum, IDEXX
379 Rum, SVANOVIR and BIOVET. Accordingly, the specific antibody levels differed
380 throughout the experiment depending on the test, as evidenced using an ANOVA with
381 Tukey's multiple comparison test analysis. Thus the HIPRA-CIVTEST, IDVET, LSI
382 Bov, IDEXX Bov, IDEXX Rum and BIOVET behaved similarly, and significant
383 differences were observed in the pair-wise comparisons that included 1 and 2wpi.
384 Conversely, significant differences for LSI Rum and BIO-X were detected when all the
385 time points were compared to 1 and 3 wpi, respectively. The VMRD and SVANOVIR
386 showed significant differences between 1 wpi and 4 wpi onwards because large standard
387 deviations were observed at different dpi.

388 When serum antibody levels of experimentally infected cows were analyzed, all
389 the cows were seropositive using the BIO-X, LSI Rum, IDEXX Bov and VMRD,
390 whereas the HIPRA-CIVTEST, IDVET, LSI Bov, IDEXX Rum and BIOVET displayed
391 positive and negative results at 2wpi. Seroconversions were observed at 3 wpi for the

392 SVANOVIR. The antibody level kinetics throughout the experiment were more variable
393 in cows than in bulls. The most relevant results were as follows: significant differences
394 were observed in the 2 wpi pair-wise comparisons between the HIPRA-CIVTEST,
395 IDVET, LSI Bov and BIOVET or the 3 wpi comparisons between the IDVET and BIO-
396 X. For the LSI Rum, VMRD, IDEXX Bov, IDEXX Rum and SVANOVIR, significant
397 differences were only observed for several pair wise comparisons, including 2 wpi.
398 Finally, all samples from negative control heifers were seronegative by all ELISAs
399 employed.

400

401 **4. Discussion**

402

403 This is the first study to compare commercially available ELISA tests for the
404 serological diagnosis of bovine neosporosis. The results in the present work showed
405 generally higher test agreement and performance compared to previous works, which
406 suggests that the serodiagnosis of bovine neosporosis is currently accurate. However,
407 the results also indicate that further refinements are required.

408 Few comparative studies of commercial tests have been performed. Several
409 studies have compared two commercial ELISAs (Wu et al., 2002; Hall et al., 2006) or
410 three tests (two commercial ELISAs and one agglutination test; Waldner et al., 2004).
411 However the most complete studies have compared the serological tests employed in
412 Europe and North America. In particular von Blumröder et al. (2004) studied the
413 performance of six commercial and five in-house serological tests and an in-house IFAT
414 used in Europe, and they reported a high level of agreement among them. Despite this
415 positive result, the authors offered an *a posteriori* adjustment of all the tests to obtain
416 more comparable results irrespective of the test employed. Wapenaar et al. (2007) later
417 studied the performance of serological tests employed in North America (Canada and
418 USA) (three commercial ELISAs, an in house IFAT, an agglutination test and a
419 commercial IFAT that was regarded as the reference test), and they showed several
420 significant discrepancies among the tests.

421 In general, most of the available tests for the diagnosis of bovine *N. caninum*
422 infection have shown strong performances (reviewed by Dubey and Schares 2006).
423 However, reference diagnostic laboratories have noted the need to work with accurately
424 validated commercial ELISAs because discrepancies between tests still exist

425 (previously noted by Aguado-Martínez et al., 2006), and the results obtained from these
426 tests are often employed in control programs. A low number of discrepancies among
427 tests is acceptable because they often correspond to values near the cut-off and are
428 difficult to avoid. However, no information about test performance is available from
429 previous comparative studies for several tests as mentioned below. Furthermore, the
430 performance reported by the manufacturer is either based on the analysis of a short
431 panel of reference sera, is outdated or the target population has changed.

432 The panel of commercially available ELISAs has notably changed in the last few
433 years. Indeed, some of the previously validated tests are either no longer available or
434 have been modified. The ELISA kits that are currently on the market and have been
435 used in different diagnostic and/or epidemiological studies are: the HIPRA-CIVTEST
436 (Álvarez-García et al., 2003; von Blumröder et al., 2004), IDEXX Bov (Pare et al.,
437 1995; Bien et al., 2012), SVANOVIR (Bjorkman et al., 1997; von Blumröder et al.,
438 2004; Malmsten et al., 2011), BIOVET (Pare et al., 1995; Waldner et al., 2004) and
439 VMRD (Kyaw et al., 2004; Wapenaar et al., 2007). For example, only the in-house
440 ELISA on which the SVANOVIR was based, not the SVANOVIR itself, has previously
441 been compared in a multi-centered study (von Blumröder et al., 2004). Other ELISAs
442 are relatively new, and their performances need to be corroborated. Such tests include
443 the IDVET (Spilovska et al., 2009; De Craeye et al., 2011), BIO-X (Ghalmi et al.,
444 2009), IDEXX Rum (Pare et al., 1995), LSI Bov (Bartels et al., 2005) and LSI Rum. On
445 the contrary, previously employed commercial ELISAs, such as the Pourquier (Institut
446 Pourquier, Montellellier, France), Cypress (Cypress Diagnostics CV), p38 (AFOSA),
447 Chekit Bommeli/ Intervet and Mastazyme (MastazymeTM MAST Diagnostics) (von
448 Blumröder et al., 2004) are no longer marketed.

449 Most of the tests used in this study showed high levels of agreement and high Se
450 and Sp values irrespective of the gold standard considered, indicating good or excellent
451 performance. This result was expected based on the similarity of the tests (antigen,
452 principles and technical aspects) and according to previous reports (von Blumröder et
453 al., 2004). The best-adjusted ELISAs were HIPRA-CIVTEST, IDVET, BIOVET and
454 IDEXX Rum that showed excellent Se and Sp values (>95%). A slight increase in
455 agreement and more balanced Se and Sp values were obtained with the BIO-X, LSI
456 Bov, LSI Rum and IDEXX Bov when the adjusted cut-offs from the TG-ROC analysis
457 were applied. However, the most significant adjustment was performed for the
458 SVANOVIR and VMRD after TG-ROC analysis, indicating that the tests were not

459 inferior, but they were improved notably after the validation study. Indeed, in a previous
460 study, the in-house ELISA version of the SVANOVIR that employed ISCOM
461 tachyzoite extract as an antigen showed 98% Se and Sp values after the “majority of
462 tests”-based adjustment (von Blumröder et al., 2004). Moreover previous VMRD results
463 showed 89% Se and 99% Sp values (Wapenaar et al., 2007). The contradictory results
464 obtained with this test may be explained by the different gold standard employed (based
465 on the IFAT test results) and the composition of the panel of sera. In general, the
466 differences observed in the pair wise comparisons of all the tests are likely due to the
467 validation processes conducted by the manufactures. This is evidenced by an increase in
468 the Se and Sp values upon re-adjustment based on TG-ROC analysis. Although most of
469 the ELISAs studied here were indirect assays based on whole or soluble tachyzoite
470 extract, there were three tests based on single antigens or a mixture of purified antigens.
471 These tests showed variable results, with either high Se and Sp (BIO-X), high Se and
472 low Sp (VMRD) or high Sp and low Se values (SVANOVIR). Moreover, cut-off values
473 for the different tests to differentiate between positive and negative results were
474 similarly adjusted for both gold standards employed in the present work. These results
475 may be explained by the restrictive criteria to classify animals as infected or non-
476 infected. There is no consensus on the most appropriate gold standard for this type of
477 study. However, it seems reasonable that results may be less biased with “the majority
478 of the tests” than by relying on a single gold standard based on previous comparative
479 studies that included assays with varying principles (von Blumröder et al., 2004).

480 A similar validation study is highly recommended for commercially available
481 IFAT assays (IFAT VMDR Inc.), taking into account a certain degree of subjectivity
482 inherent to such tests that rely on visualizing results under a fluorescence microscope
483 (reviewed by Bjorkman and Uggla, 1999).

484 Other in-house ELISAs have been suggested for the serological diagnosis of
485 neosporosis. Validation of these tests with a well-referenced sera panel would be
486 similarly recommended prior to considering them as routine diagnostic tools. This issue
487 is of particular relevance for recombinant protein-based ELISAs that may not provide
488 Se and Sp values as high as tachyzoite extract-based ELISAs (Aguado-Martínez et al.,
489 2008) or ELISAs that have been standardized only with sera from experimental
490 infections (Hiasa et al., 2012; Yin et al., 2012). These ELISAs can provide additional
491 information about the route or time of infection, but at present they cannot replace the
492 commercial ELISAs validated herein for conventional serodiagnosis (Aguado-Martínez

493 et al., 2008). The same is true for the agglutination tests (NAT), which are very useful
494 techniques for seroprevalence studies in several species that do not have available
495 secondary antibodies (reviewed by Bjorkman and Uggla, 1999). However, they are
496 presently not recommended for the diagnosis of bovine neosporosis by reference
497 diagnostic laboratories. In this sense, recent studies have shown a preference for the
498 competitive ELISA VMRD for the detection of anti-*N. caninum* antibodies in the sera
499 of species other than cattle, such as dogs (Sharma et al., 2008), cats (Millán et al.,
500 2009), wild ruminants and sheep (Panadero et al., 2010), wild boars (Bartova et al.,
501 2006), horses (Bartova et al., 2010a), hares (Bartova et al., 2010b) and pigs (Bartova
502 and Sedlak, 2011).

503 We included three multi-species commercial tests in our study (IDEXX Rum for
504 bovines, caprines and ovines, and LSI Rum and IDVET for ruminants). Further
505 validations should be conducted with the appropriate reference sera to confirm their
506 performance for species other than bovines. This recommendation should also apply to
507 the VMRD test for any use other than cattle sera. In the absence of positive and negative
508 reference sera, it is highly recommended to check the sera employed in the validation
509 study using another *a posteriori* and confirmatory diagnostic tool, such as the western
510 blot (Malmsten et al., 2011).

511 Another important issue is the study of cross-reactions with closely related
512 apicomplexan parasites with relevance to cattle, such as *Sarcocystis* spp. and *Besnoitia*
513 *besnoiti*. It is well known that 100% of cattle are infected with *Sarcocystis* spp. (Dubey
514 et al., 1989). However, bovine besnoitiosis is a re-emerging disease in Europe (EFSA,
515 2010). Therefore, these organisms frequently co-exist with *N. caninum* infections. Thus,
516 cross-reactions should be avoided for an accurate diagnosis. In the present study, most
517 of the tests did not yield false positive results, and only three tests showed 7-28% false
518 positive results. Cross-reactions may be responsible for a high percentage of false
519 positive results when using a panel of bovine sera from a different origin (Nasir et al.,
520 2012). Thus, it would be desirable to discard cross-reactions with these three tests by
521 employing a wide panel of appropriate sera.

522 Commercial ELISAs are employed not only for conventional diagnoses by
523 reference diagnostic laboratories but also by research laboratories with varying
524 purposes. Interestingly, only three tests (HIPRA-CIVTEST, Bio-X and BIOVET)
525 showed significant differences in specific antibody levels between aborted and non-
526 aborted cows. It has been reported that aborting cows develop higher specific antibody

527 levels than non-aborting cows (Pereira-Bueno et al., 2000). Therefore, the results of this
528 study show the ability of these ELISA kits to be used in observational and experimental
529 studies

530 As expected, all the tests showed similar antibody level kinetics for the
531 experimental infections. Because experimental infections are performed under
532 controlled conditions (isolate, dose, route of inoculation, management measures),
533 seroconversions occur in a similar manner. However, more discrepancies were observed
534 when the sera from cows were analyzed. In this case, the antibody level kinetics varied
535 depending on the kit employed. This finding may be explained by the fact that the sera
536 came from pregnant cows, and variations in antibody levels during pregnancy could
537 influence the performance of the kits. Indeed, individual variations in antibody levels
538 were more evident in cows compared to bulls. Although sera from experimental
539 infections are convenient for initial validations of serological tests, field sera are always
540 recommended for further refinements; results observed with sera from experimental
541 infections do not necessarily correlate to results extended to a target population.

542 In summary, we have evaluated the performance of all the commercially
543 available ELISA tests through the analysis of a well-defined panel of sera and
544 subsequent refinement based on TG-ROC analyses. Serological assays are essential
545 tools for control programs, they are preferred by diagnostic laboratories and they are
546 also used in many epidemiological studies. The information obtained here increases the
547 robustness of the tests, which is essential for providing confidence in assay performance
548 to the reference diagnostic laboratories. Moreover, the results obtained here may help
549 assay developers to follow the guidelines for validation and certification of diagnostic
550 assays elaborated by the OIE (Wright et al., 2007).

551 According to Jacobson (1998), the validation of diagnostic assays, rather than
552 relying on a small number of experiments that are based on limited reference samples, is
553 a process involving constant surveillance and readjustment of performance
554 characteristics for each target population, an observation that has been corroborated by
555 the present work. Moreover, accredited diagnostic laboratories are highly encouraged to
556 conduct multi-centered studies with validated assays (eg. Dargatz et al., 2004). This is
557 the best way to assess reproducibility and to provide consistent results among
558 laboratories according to the principles and methods of diagnostic test validation. This
559 is of great importance because reproducibility may significantly influence results.

560

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564

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717 **Tables**

718

719 **Table 1:** ELISA tests used in the comparative study.

720

721	Trademark (ID Test)	Antigen	Type	Cut-off value	References
722	CIVTEST Bovis Neospora (HIPRA-CIVTEST)	Sonicate lysate of tachyzoites	iELISA	$>10/6^a$ RIPC=(ODs -ODnc/ODpc- ODnc)x100	Álvarez-García et al., 2003
723	IDVET ID Screen (IDVET)	Sonicate lysate of tachyzoites	iELISA	$\geq 50/41^a$ S/P=ODs -ODnc/ODpc- ODnc	-
724	LSIVet Bovine (LSI Bov)	Sonicate lysate of tachyzoites	iELISA	≥ 30 RIPC=(ODs-ODnc/ODpc-ODnc)x100	-
725	LSIVet Ruminant (LSI Rum)	Sonicate lysate of tachyzoites	iELISA	≥ 30 RIPC=(ODs-ODnc/ODpc-ODnc)x100	-
726	Bio-X Diagnostics (BIO-X)	NcSRS2 purified protein	iELISA	$>15/10^a$ Val=(Delta ODs)x100/(Delta ODp)	Ghalmi et al., 2009
727	VMRD Inc. (VMRD)	Surface protein antigen (GP65) captured using a monoclonal antibody	cELISA	≥ 30 %I=100-[(ODsx 100)/(ODmnc)]	Baszler et al., 1996 Baszler et al., 2001
728	IDEXX Neospora X2 (IDEXX Bov)	Sonicate lysate of tachyzoites	iELISA	≥ 0.50 S/P=ODs-ODnc/ODpc-ODnc	Paré et al., 1995
729	IDEXX Chekit Neospora (IDEXX Rum)	Detergent lysate of tachyzoites	iELISA	$\geq 40/30^a$ RIPC=(ODs -ODnc/ODpc- ODnc)x100	Paré et al., 1995
730	Nc iscom ELISA. Svanovir (SVANOVIR)	Tachyzoite proteins incorporated into iscoms	iELISA	≥ 20 PP=[(mODs or nc)/mODpc]x100	Björkman et al., 1997
731	Biovet- <i>Neospora caninum</i> (BIOVET)	Sonicate lysate of tachyzoites	iELISA	≥ 0.60 R=(mODs-mODwsc)/(mODpc mODwsc)	Paré et al., 1995 Waldner et al., 2004

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a, doubtful cut-off; i, indirect; c, competitive; OD, optical density; IRPC, relative index per cent; S/P, sample/positive; Val, validation;

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%I, percent inhibition; PP, percent positivity; R, ratio; s, sample; pc, positive control; nc, negative control; m, mean; wsc, wash solution

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control.

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737 **Table 2:** Se and Sp values relative to gold standard criteria on the basis of the cut-offs suggested by manufacturers.

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Test ID	Majority of tests		Pre-test information	
	Se 95%(CI)	Sp 95%(CI)	Se 95%(CI)	Sp 95%(CI)
HIPRA-CIVTEST	96.1 (93.9-98.4)	100 (100-100)	95.7 (93.2-98.2)	100 (100-100)
IDVET	99.6 (98.9-100)	98.9 (97.4-100)	98.9 (97.6-100)	98.3 (96.3-100)
LSI Bov	99.3 (98.2-100)	94.1 (90.7-97.5)	98.9 (97.6-100)	98.9 (97.3-100)
LSI Rum	99.6 (99.0-100)	93.0 (89.3-96.7)	98.9 (97.6-100)	97.2 (94.7-99.6)
BIO-X	98.9 (97.6-100)	94.9 (91.7-98.2)	98.5 (97.0-100)	94.0 (90.4-97.6)
VMRD	98.9 (97.6-100)	65.1 (58.2-71.9)	98.5 (97.0-100)	66.5 (59.5-73.5)
IDEXX Bov	100 (100-100)	93.0 (89.3-96.7)	99.3 (98.2-100)	96.6 (93.9-99.3)
IDEXX Rum	95.8 (93.3-98.2)	100 (100-100)	96.1 (93.8-98.5)	100 (100-100)
SVANOVIR	85.9 (81.8-90.1)	99.5 (98.5-100)	87.2 (83.2-91.2)	100 (100-100)
BIOVET	98.9 (97.6-100)	98.9 (97.4-100)	98.5 (97.0-100)	98.8 (97.3-100)

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763 **Table 3:** Se and Sp relative to gold standard criteria on the basis of the re-calculated cut-offs after TG-ROC analysis.
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765	Test ID	Cut-off employed	Majority of tests		Pre-test information	
766			Se 95%(CI)	Sp 95%(CI)	Se 95%(CI)	Sp 95%(CI)
767	HIPRA-CIVTEST	>10/6 ^a	96.1 (93.7-98.5)	99.5 (98.4-100)	95.7 (93.6-98.3)	100 (100-100)
768	IDVET	≥50/41 ^a	99.6 (98.9-100)	97.3 (94.9-99.6)	98.9 (97.7-100)	98.3 (96.3-100)
769	LSI Bov	>52 *	98.9 (97.6-100)	97.9 (95.9-99.9)	98.2 (96.7-99.8)	99.4 (98.3-100)
770	LSI Rum	>53 *	99.3 (98.2-100)	97.4 (95.1-99.6)	98.6 (97.2-100)	98.9 (97.3-100)
771	BIO-X	>23/13 ^a *	98.1 (96.4-99.7)	96.2 (93.4-99.0)	97.8 (96.0-99.5)	97.0 (94.5-99.6)
772	VMRD	>65 *	94.8 (92.1-97.4)	91.6 (87.6-95.5)	94.3 (91.6-97.0)	94.3 (90.9-97.7)
773	IDEXX Bov	>1 *	98.5 (97.1-100)	97.9 (95.9-99.9)	97.9 (96.2-99.6)	98.3 (96.4-100)
774	IDEXX Rum	≥40/30 ^a	96.5 (94.3-98.7)	99.5 (98.4-100)	96.1 (94.1-98.6)	100 (100-100)
775	SVANOVIR	>15 *	94.0 (91.2-96.9)	93.7 (90.2-97.1)	94.0 (91.2-96.7)	94.3 (90.9-97.7)
776	BIOVET	≥ 0.60	99.2 (98.2-100)	98.4 (96.6-100)	98.5 (97.1-100)	98.8 (97.3-100)

778 ^a cut-off with a range of doubtful results

779 *re-calculated cut-offs after TG-ROC analysis

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784 **Table 4:** Test agreement between ELISAs and gold-standard criteria.

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Gold standard criteria		K-values (95%CI)*					
		LSI Bov	LSI Rum	BIO-X	VMRD	IDEXX Bov	SVANOVIR
Majority of tests	Prior to TG-ROC	0.94 (0.91-0.97)	0.94 (0.90-0.97)	0.94 (0.91-0.97)	0.67 (0.60-0.74)	0.94 (0.91-0.97)	0.83 (0.78-0.88)
	After TG-ROC	0.97 (0.95-0.99)	0.97 (0.95-0.99)	0.95 (0.92-0.98)	0.88 (0.83-0.92)	0.97 (0.94-0.99)	0.87 (0.82-0.92)
Pre-test information	Prior to TG-ROC	0.98 (0.96-1)	0.96 (0.94-0.99)	0.93 (0.89-0.97)	0.69 (0.61-0.76)	0.96 (0.94-0.99)	0.85 (0.79-0.89)
	After TG-ROC	0.97 (0.95-0.99)	0.97 (0.95-0.99)	0.94 (0.91-0.97)	0.88 (0.83-0.92)	0.95 (0.93-0.98)	0.87 (0.83-0.92)

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* Almost perfect agreement is marked in bold (K-values > 0.95)

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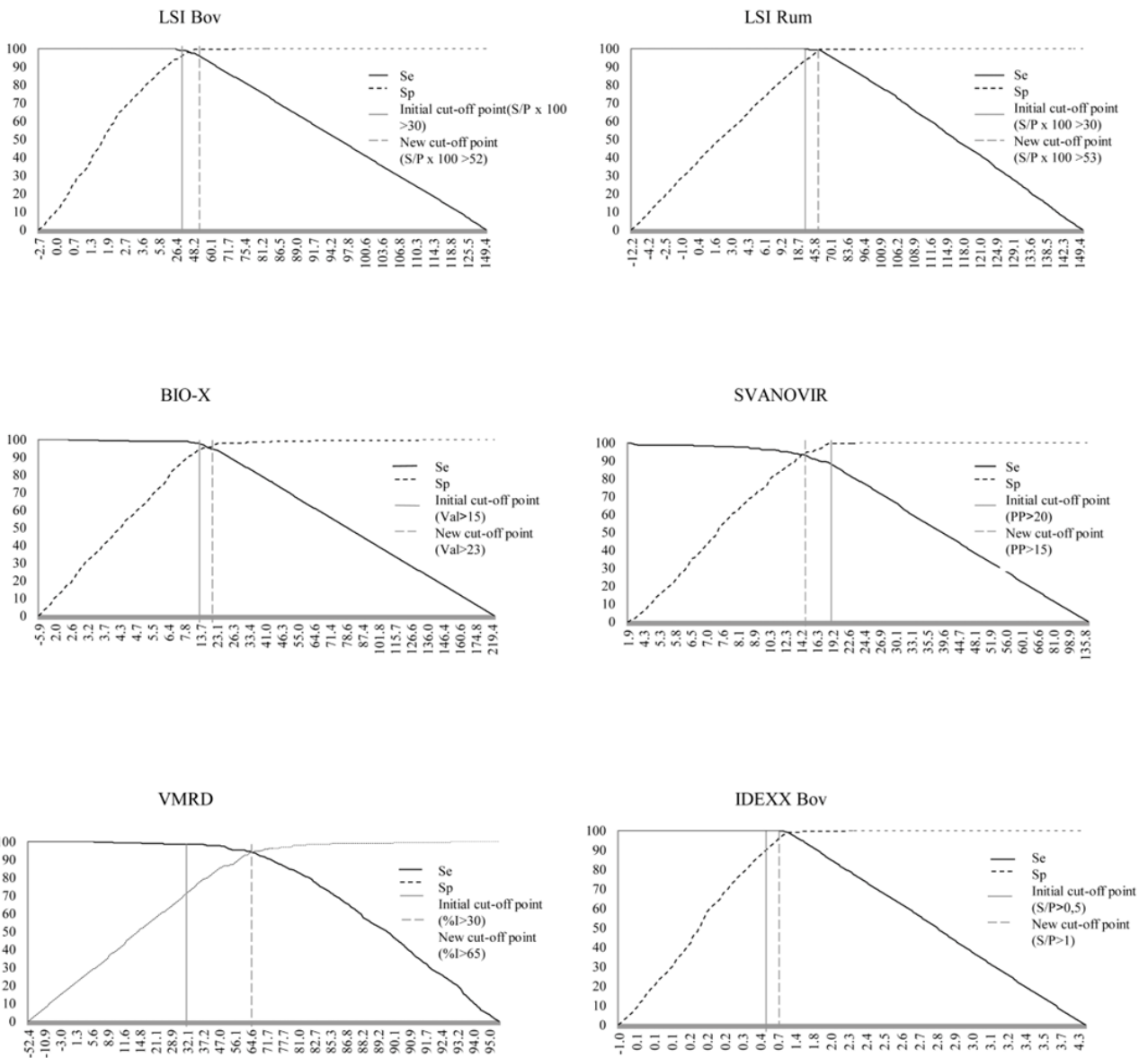
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794 **Figures**

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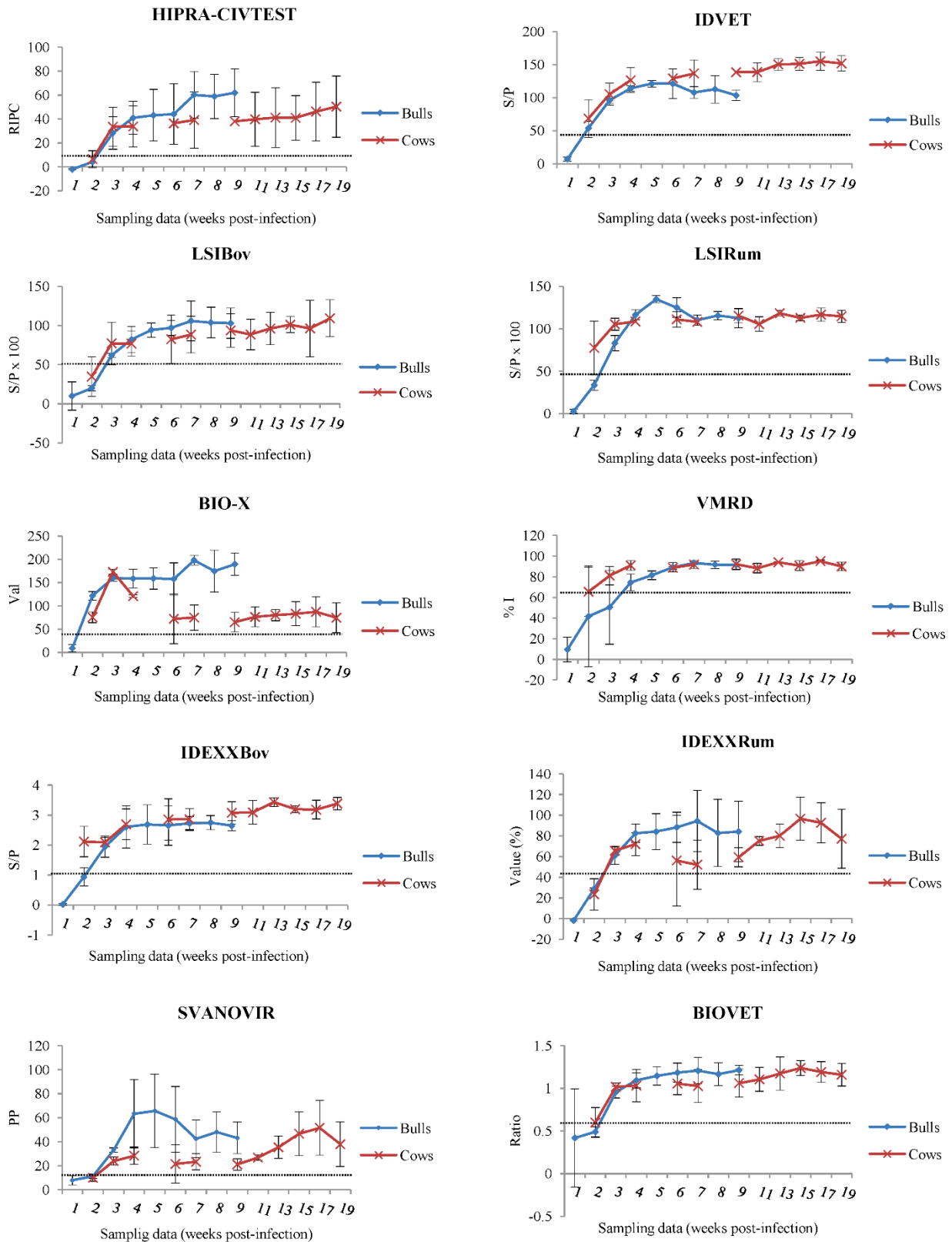
799 **Figure 1:** TG-ROC analysis of 6 commercial ELISA tests based on the gold standard ‘Majority of
800 tests’.

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806 **Figure 2:** Specific anti-*N. caninum* antibodies developed over time by experimentally infected bulls
 807 and cows at 135 days of gestation. Recalculated cut-offs were employed for the LSI Bov, LSI Rum,
 808 BIO-X, VRMD, IDEXX Bov and SVANOVIR ELISAs.