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Standardisation and comparison of serial dilution and single dilution enzyme linked immunosorbent assay (ELISA) using different antigenic preparations of the Babesia (Theileria) equi parasite

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Abstract – Serial dilution and single dilution enzyme linked immunosorbent assays (ELISA) were standardised and their sensitivity and specificity were compared for serodiagnosis of Babesia equi infection. The antibody titres of 24 donkey sera of known identity were determined separately by serial dilution ELISA using three different B. equi antigens namely whole merozoite (WM), cell membrane (CM) and high speed supernatant (HSS). The ratios of the optical density (OD) of known positive and known negative sera at different serum dilutions were calculated and termed as the positive/negative (P/N) ratio. The coefficients of correlation (r) were calculated between the P/N ratios at different dilutions of sera and the log_{10} antibody titres of the same sera were ascertained by serial dilution ELISA. The highest value of ‘r’ was obtained at a serum dilution of 1:200. From log_{10} antibody titre of sera (y) and their P/N ratio at a dilution of 1:200 (x), regression equations (y = a + bx) were calculated separately for the three antigens. Test sera were diluted to 1:200, their OD were read in duplicate wells and were converted to the P/N ratio. Antibody titres were predicted from the P/N ratio using a regression equation separately for the three antigens. Titres obtained by both ELISAs were not significantly different from each other, thus confirming that single dilution ELISA could be successfully used to replace conventional serial dilution ELISA. The sensitivity, specificity and predictive value of single dilution ELISA was validated statistically using 42 B. equi disease-positive sera and 106 B. equi disease-negative sera. The WM antigen was found to be the most sensitive with a higher predictive value for negative test sera as compared to the CM or HSS antigens. Sera positive for other equine infections including Babesia caballi showed no cross-reaction with the three B. equi antigens in ELISA, thus the test was immunologically specific. Antibody titres of 109 unknown field donkey/horse sera obtained by serial and single dilution ELISA using the WM antigen did not show any significant difference. Since the single dilution ELISA was found to be

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more economical, convenient, sensitive, specific than the serial dilution ELISA and has a high predictive value, it is suitable for use in sero-epidemiological studies on B. equi infections in the field.

**Babesia equi / Theileria equi / protozoan / sero-diagnosis / ELISA**

1. INTRODUCTION

Equine babesiosis is an acute, subacute or chronic tick-borne disease of the equidae (horse, pony, donkey, mule and zebra), caused by the intraerythrocytic protozoan Babesia (Theileria) equi and/or Babesia caballi. Babesia equi and B. caballi share the same vectors, but in an enzootic region B. equi is reported to be more common than B. caballi [20]. The disease has attained worldwide importance due to the widespread prevalence of associated tick vectors [4]. Equine babesiosis caused by B. equi is endemic in India, but only isolated clinical cases have been reported [26, 27]. However, outbreaks of the disease have also been reported when clean animals are introduced into endemic zones [12].

Diagnosis of the disease can be made on the basis of clinical signs and demonstration of B. equi / B. caballi organisms in blood smears. However, serological tests are required for diagnosis in latent or carrier animals. A number of techniques have been used to measure B. equi antibodies in carrier animals. The complement fixation test (CFT) is the test recommended by the Office International des Epizooties (OIE) for the detection of the Babesia species antibodies, combined with IFAT in doubtful cases. These two tests have been used by various researchers [14, 22, 30]. However CFT has the inherent disadvantages of giving false positive results and low sensitivity for detecting latent infections [6, 18, 19, 25]. Various forms of ELISA have been standardised and are reported to be more sensitive than CFT [6, 18]. However the cost per sample of ELISA is high and it is tedious and inconvenient when the titre of a sample is to be determined. Recently, single dilution ELISAs have been standardised and used for the sero-diagnosis of the foot and mouth disease [33], bovine babesiosis [1], Pasteurella multocida infection in buffaloes [17] and tropical theileriosis in cattle [17, 24]. In conventional serial dilution ELISA, the end titre of a sample is calculated by serial two fold dilution of the sample until the optical density (OD) becomes negative. However in single dilution ELISA, only one particular dilution of the serum sample is tested (that is why it is named the single dilution ELISA). The OD value of the one dilution tested is used in the regression equation for predicting the end antibody titre. So single dilution ELISA is less cumbersome, more economical in terms of consumption of antigen/conjugate/reagents and suitable for field studies [17, 28] since the titre of a serum sample can be predicted by testing it at only a single dilution. The estimation of exact antibody titres by single dilution ELISA during sero-epidemiological surveys would help in avoiding the movement of naïve horses into endemic regions. We describe here the standardisation of a serial dilution ELISA for detecting antibodies against B. equi using three antigenic preparations. The assay was then adapted to a single dilution ELISA format.

2. MATERIALS AND METHODS

2.1. Parasite antigen

A healthy donkey, negative for B. equi parasites by blood smear examination and negative for B. equi antibodies by Dot-ELISA was obtained and maintained under
tick-free conditions. The donkey was also serologically negative for *Trypanosoma evansi* and glanders. The donkey was splenectomised as described by Dennig and Brocklesby [7] and injected with 50 mL of blood collected in an acid citrate solution (3.5% w/v) from a known *B. equi* carrier animal, which was continuously maintained in the laboratory. When the parasitaemia reached about 80% in the donkey, blood was collected in EDTA bottles (1.2 g/L of blood) and was processed for preparation of antigens as described by Brown et al. [5]. Briefly, the plasma and buffy coat were discarded following centrifugation (8000 g × 30 min at 4 ºC) and the erythrocyte pellet was washed three times with normal saline solution (NSS). The erythrocytes were lysed overnight at 4 ºC by adding an equal volume of sterile distilled water. The erythrocyte lysate was centrifuged at 8000 g for 30 min at 4 ºC. The pellet was washed once with NSS and centrifuged again. The sediment was resuspended in NSS, homogenised in a glass homogeniser and centrifuged until the haemoglobin was completely removed from the lysate pellet. The pellet was resuspended in an equal volume of cold NSS containing 1 mM of phenylmethlysulfonyl fluoride (PMSF) and centrifuged at 800 g for 1 min. The supernatant containing *B. equi* merozoites was collected, sonicated 8 times at 100 W for 30 s and was divided into two equal parts. One part was retained as the whole merozoite (WM) antigen. The second part was centrifuged at 145 000 g for 1 h at 4 ºC. The supernatant was collected and designated as the high-speed supernatant (HSS) fraction. The sediment was collected separately, mixed with an equal volume of cold NSS containing 1 mM PMSF, homogenised and designated the cell membrane (CM) fraction. The total protein concentration of these antigens was estimated by the method of Lowry et al. [21] and was expressed in mg/mL. All antigens were stored at 4 ºC.

### 2.2. Reference sera

In a previous study [19], four donkeys were experimentally infected with *B. equi* by intravenous injection of blood with a high parasitaemia (~ 70%). Serum samples were collected daily for the first 7 days post-infection (PI), then biweekly up to 40 days and finally once a week up to 89 days PI. Twenty-four (six samples from each four donkeys) serum samples with antibody titres ranging from low positive (1:200 to 1:800, representing samples collected during 1–10 days PI) to high positive (1:1600 to 1:51200, representing samples collected during 16 to 89 days PI) by Dot-ELISA were selected and used to standardise the serial and single dilution ELISAs. Another batch of 21 serum samples, collected from asymptomatic healthy donkeys (positive for the *B. equi* antibodies by Dot-ELISA) from different areas of the Haryana State, India, were selected and used in the present study to validate the titres obtained by serial dilution and single dilution ELISAs. Three reference negative control sera were collected from such donkeys, which had been repeatedly found negative for the *B. equi* parasite and antibodies by blood smear examination and Dot-ELISA, respectively. A reference *B. caballi* positive serum (1:80 CFT titre) was obtained from the United States Department of Agriculture (USDA), USA. Sera positive for *Trypanosoma evansi*, glanders, strangles, equine infectious anaemia and equine influenza were obtained from a laboratory testing clinically infected equids.

### 2.3. Serological tests

#### 2.3.1. Serial dilution ELISA

The optimum dilution of the three antigens (WM, CM and HSS) and the conjugate [rabbit anti-horse gamma globulin (IgG) horse radish peroxidase, HRP, Sigma Aldrich Co., St. Louis, USA] were determined using a block titration method.
The optimum dilution considered was the highest dilution of antigen/conjugate that gave a maximum contrast in terms of optical density (OD) between known positive and known negative serum dilutions. The technique described by Bose and Peymann [2] for *B. caballi* was adopted. Dilutions of 1:5, 1:10, 1:20 and 1:40 of the WM, CM and HSS antigens were prepared in 0.1 M carbonate bicarbonate buffer, pH 9.6 and 100 µL of each dilution was used to coat the wells of a 96 well microtitre ELISA plate (Greiner, Germany). The plates were incubated at 37 ºC for 1 h and were kept at 4 ºC overnight. These were then washed three times with PBS using an ELISA plate washer (Organon Teknika washer, Veedijk, 58, 2300, Turnhout, Belgium). The unoccupied sites in each well were blocked by adding 200 µL of 3% bovine serum albumin (BSA) dissolved in PBS, pH 7.2 (PBS–3% BSA) for 1 h at 37 ºC. The plates were again washed three times with PBS and 100 µL of two fold serum dilutions, from 1:50 to 1:51200 in PBS containing 1% BSA (PBS–1% BSA) were added to wells 1 to 11 in each row. Eight samples were loaded in each plate. The last (12th) column’s wells of ELISA plate were used for controls, which included (a) a known positive sera; (b) a known negative sera; (c) serum + conjugate without antigen; (d) antigen + conjugate without serum; (e) antigen + serum + conjugate without substrate; (g) PBS alone and (h) conjugate + substrate. The plates were incubated for 1 h at 37 ºC, followed by washing with PBS–0.05%Tween-20 (PBS-T). Dilutions of the conjugate (1:500, 1:1000, 1:2000 and 1:4000) were prepared in PBS–1% BSA. The desired dilution of the conjugate (100 µL/well) was added to the appropriate wells and the plates were incubated for 1 h at 37 ºC, followed by three more washes with PBS-T. Ortho-phenylene diamine dihydrochloride (OPD, Sigma) solution was prepared as per the manufacturer’s instructions and 100 µL dispensed into the wells except in well ‘e’ of control column 12. The plates were kept in the dark at room temperature for 20–30 min. On the development of a yellow colour and when the OD of the positive control well (h12) was 1.0–1.2, 50 µL of 2 M H₂SO₄ was added to each well to stop the enzymatic reaction. The plates were read at 492 nm in an ELISA plate reader (Organon Teknika Reader 530, Veedijk, 58, 2300, Turnhout, Belgium). The mean OD of the negative control wells (b12 to d12) was calculated. Any sample showing an OD above the mean + (3 × standard deviation) of three negative wells was considered positive. Standard deviation (SD) of the OD of three negative wells (b12 to d12) was calculated as per a standard procedure [29]. The highest dilution of the serum sample showing a positive reaction was recorded as the titre.

### 2.3.2. Single dilution ELISA

Regression analysis, as used by Briggs and Skeeles [3] and Manuja et al. [24] to predict the end point of antibody titres to *Pasteurella multocida* in chicken sera and *Theileria annulata* infection in bovine sera, respectively, was adopted.

The log₁₀ end point antibody titres of 24 known positive (y₁) and 3 known negative control sera were determined by diluting the sera from 1:50 to 1:51200 and using the three antigens viz; WM, CM and HSS separately by serial dilution ELISA. The mean OD values of the 24 known positive sera at 1:50, 1:100, 1:200, 1:400, 1:800 dilutions were divided by the mean OD of the three known negative serum samples at the respective dilution and were termed as positive/negative (P/N) ratios separately for each antigen. The coefficient of correlation (r) between the log₁₀ end titre and the P/N ratio (at serum dilutions of 1:50 to 1:800) was calculated separately for the three antigens as per the method described by Snedecor and Cochran [29]. The dilution of serum (1:50 to 1:800) showing the maximum value of ‘r’ was chosen for the derivation of the regression equation (\( y = a + bx \)), where \( y = \text{log}_{10} \) antibody titre of the
test serum, \( x = \) the \( P/N \) ratio of the test
serum at the chosen dilution, \( a = \) the con-
tant and \( b = \) the regression coefficient) for
predicting \( \log_{10} \) end titre by single dilution
ELISA. The values of ‘\( a \)’ and ‘\( b \)’ were cal-
culated separately for the three antigens
(WM, CM and HSS), to have a unique
regression equation for each antigen.

The test sera were diluted to the chosen
dilution (which showed a maximum ‘\( r \)’
value as described above) and were added
into duplicate wells of ELISA plates
coated separately with three different anti-
gens. The assay was run as described under
serial dilution ELISA and the OD was
recorded at 492 nm. The mean OD for each
sample (recorded in three ELISA plates
each coated with the WM or CM or HSS
antigen) was converted to the \( P/N \) ratio (\( x \))
and the \( \log_{10} \) end antibody titres (\( y \)) were
predicted by using the value of ‘\( x \)’ in the
regression equation separately derived for
the three antigens.

2.4. Validity of single dilution ELISA

The antibody titres of 21 known posi-
tive field sera from \( B. equi \) antibody carrier
donkeys were determined by serial dilution
ELISA and were predicted by single dilu-
tion ELISA for the three antigens sepa-
rate. The observed \( \log_{10} \) antibody titres
and predicted \( \log_{10} \) antibody titres were
compared for significance (\( P < 0.05 \)) using
factorial complete randomised design by
ANOVA test and their coefficient of corre-
lation was also calculated [29].

2.5. Sensitivity, specificity, predictive
value and cross-reactivity

The statistical sensitivity, specificity
and predictive value of single dilution
ELISA with the three antigens were vali-
dated using the method described by Tyler
and Cullor [31]. For this purpose, samples
from the previously collected known
\( B. equi \) disease-positive and disease-nega-
tive animals were used [19]. The protocol
for this experiment is described in
paragraph 2.2 of this article. Serum sam-

ples were collected after the patency of
\( B. equi \) parasitaemia (3 days PI onwards)
in the four infected donkeys. Forty-two
sera were selected (with at least 10 samples
from each of the four donkeys) as disease-
positive samples. One-hundred and six
sera were also collected simultaneously
as \( B. equi \) disease-negative samples from
50 healthy donkeys that were reared in tick
free housing facilities. These animals were
found to be negative for the \( B. equi \) para-
site by repeated blood smear examination
and their sera were negative for antibodies
by Dot-ELISA and the complement fixa-
tion test.

The immunological cross-reactivity of
the \( B. equi \) antigens (WM, CM, HSS) with
sera positive for \( B. caballi, Trypanosoma
evansi, \) glanders, strangles, equine infec-
tious anaemia and equine influenza were
tested by performing serial dilution ELISA
as standardised above using two fold
serum dilutions from 1:50 to 1:51200. The
immunological sensitivity was determined
by serially diluting high titred \( B. equi \)
positive serum and a negative serum (sera
used as described in paragraph 2.2 refer-
cence sera) in serial dilution ELISA and by
comparing the OD of the positive serum at
a particular dilution with the OD of the
negative serum.

2.6. Unknown field sera

The \( \log_{10} \) end antibody titres of 109
field horse/donkey serum samples were
determined by serial and single dilution
ELISA in order to compare the sensitivity
of these two ELISAs on unknown samples.
Samples were obtained from different
parts of the Haryana State, India.

3. RESULTS

3.1. Optimum antigen and conjugate
dilutions

A dilution of 1:20, 1:20 and 1:5, respec-
tively of the WM, CM and HSS antigens
proved optimum for the ELISA. The protein contents were 0.370, 0.350 and 0.500 mg/mL at the respective dilutions. The anti-horse HRP conjugated secondary antibody showed optimum results at a dilution of 1:1000. The optimum dilution of WM, CM and HSS antigens and conjugate, as standardised for the serial dilution ELISA, were also used for the single dilution ELISA.

3.2. Serial dilution ELISA

At a dilution of 1:200 (log₁₀ value = 2.3) and above, the absorbance of the negative sera plateaued whereas, the positive sera continued to show a high absorbance value (Fig. 1). The observed reciprocal antibody end titre of 24 known positive serum samples was converted to the log₁₀ value. Amongst the 24 samples tested, 15 showed a log₁₀ antibody titre of 4.1 (1:12800) and above with the WM antigen whereas, 13 and 8 samples, respectively, showed a log₁₀ antibody titre of 4.1 and above with the CM and HSS antigens (Fig. 2).

3.2.1. Coefficient of correlation

The coefficients of correlation (r) between the log₁₀ antibody titre of the 24 known positive sera with their respective P/N ratios at 1:50, 1:100, 1:200, 1:400 and 1:800 are shown in Table I. The maximum value of ‘r’ was 0.938 and 0.943 observed at a serum dilution of 1:200 for the WM and HSS antigens, respectively. However with the CM antigen, the highest value of ‘r’ (0.977) was observed at a serum dilution of 1:50 followed by 0.974 at 1:200 dilution. But at the same time, known negative control sera also showed high OD values at 1:50 and 1:100 serum dilution (Fig. 1). Hence the 1:200 serum
Single dilution ELISA using *Babesia equi* antigens

Dilution was considered most favourable for the CM antigen as well. So a 1:200 serum dilution was chosen for calculating the regression equation for each of the three antigens and also for testing of an unknown sample at only one dilution by single dilution ELISA for all three antigens.

### 3.2.2. Regression equations

The regression equations calculated for the three antigens were as follows:

1. WM antigen: \( y = 2.05 + 0.51x \) (SE for ‘a’ = ±0.147, SE for ‘b’ = ±0.041).

2. CM antigen: \( y = 1.63 + 0.57x \) (SE for ‘a’ = ±0.073, SE for ‘b’ = ±0.020).

3. HSS antigen: \( y = 1.55 + 0.71x \) (SE for ‘a’ = ±0.164, SE for ‘b’ = ±0.055).

### 3.2.3. Validity of the regression equations

Reciprocal log_{10} antibody titres of the 21 sera observed by serial dilution ELISA did not differ significantly \((P > 0.05)\) from reciprocal log_{10} antibody titre by serial dilution ELISA.

![Figure 2](https://example.com/figure2.png)

Figure 2. Comparison of reciprocal log_{10} antibody titres obtained by serial dilution ELISA with the respective P/N ratio at 1:200 serum dilutions.

### Table I

Coefficient of correlation between the positive/negative (P/N) ratio and reciprocal log_{10} antibody observed titres by serial dilution ELISA at different two-fold dilutions of known positive control sera \((n = 24)\).

<table>
<thead>
<tr>
<th>Antigen</th>
<th>Coefficient of correlation at respective serum dilution</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1:50</td>
</tr>
<tr>
<td>Whole merozoite (WM)</td>
<td>0.931</td>
</tr>
<tr>
<td>Cell membrane (CM)</td>
<td>0.977</td>
</tr>
<tr>
<td>High speed supernatant (HSS)</td>
<td>0.936</td>
</tr>
</tbody>
</table>
the corresponding predicted antibody titres obtained by single dilution ELISA for the WM, CM and HSS antigens, respectively. Furthermore, the antibody titres obtained by serial dilution and single dilution ELISA using WM, CM and HSS antigens showed very high coefficients of correlation (‘r’ = 0.95, 0.95 and 0.92, respectively for the three antigens as shown in Fig. 3).

3.3. Sensitivity, specificity, predictive value and cross-reaction

As shown in Table II, the statistical specificity of the single dilution ELISA for detecting antibodies against *B. equi* was 100% with all three antigens, whereas, the sensitivity ranged from 80–88%. The predictive value for positive sera was 100% and ranged from 93–95% for negative sera using the three antigens. The WM antigen was the most sensitive of the antigens and had the highest predictive value.

Known positive sera against *B. caballi*, *Trypanosoma evansi*, glanders, strangles, equine infectious anaemia and equine influenza did not show any cross-reactivity (< 1:50 dilution) with the three *B. equi* antigens used in this study indicating the immunological specificity of the test. The OD of known *B. caballi* positive serum at 1:50, 1:100 and 1:200 dilution was lower than the mean OD of known negative sera (Fig. 1). ELISA was found to be very sensitive since positive OD values of a high titred *B. equi* serum were recorded up to a dilution of 1:51200 (Fig. 1).

3.4.Unknown field sera

A frequency distribution of antibody titres of 109 field sera tested by serial and single dilution ELISA using the WM antigen (Tab. III) shows that the results obtained using the two tests are comparable. The only difference was that two sera that were negative with serial dilution...
Single dilution ELISA using *Babesia equi* antigens

### Table II. Sensitivity, specificity and predictive value efficacy of single dilution ELISA with WM, CM and HSS antigens.

<table>
<thead>
<tr>
<th>Parameters</th>
<th>WM antigen</th>
<th>CM antigen</th>
<th>HSS antigen</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sensitivity(^a) (%)</td>
<td>88.0 (37/42)</td>
<td>86.0 (36/42)</td>
<td>81.0 (34/42)</td>
</tr>
<tr>
<td>Specificity(^b) (%)</td>
<td>100.0 (106/106)</td>
<td>100.0 (106/106)</td>
<td>100.0 (106/106)</td>
</tr>
<tr>
<td>Predictive value/positive test(^c) (%)</td>
<td>100.0</td>
<td>100.0</td>
<td>100.0</td>
</tr>
<tr>
<td>Predictive value/negative test(^d) (%)</td>
<td>96.0</td>
<td>95.0</td>
<td>93.0</td>
</tr>
</tbody>
</table>

\(^a\)Probability of accurately identifying true-positive (diseased) animals out of 42 *B. equi* disease-positive samples.

\(^b\)Probability of accurately identifying true-negative (normal) animals out of 106 *B. equi* disease-negative samples.

\(^c\)Probability that ELISA-positive animals are *B. equi* disease-positive.

\(^d\)Probability that ELISA-negative animals are *B. equi* disease-negative.

Figures in parenthesis indicate the number of positive sera / total number of samples tested.

### Table III. Frequency distribution of *Babesia equi* antibody titres as detected by serial dilution and single dilution ELISA on unknown equine sera collected from the field.

<table>
<thead>
<tr>
<th>Reciprocal antibody titre</th>
<th>No. of samples observed by</th>
<th>Interpretation</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Serial dilution ELISA</td>
<td>Single dilution ELISA</td>
</tr>
<tr>
<td>Up to 200</td>
<td>26</td>
<td>24</td>
</tr>
<tr>
<td>201–800</td>
<td>11</td>
<td>13</td>
</tr>
<tr>
<td>801–3200</td>
<td>38</td>
<td>38</td>
</tr>
<tr>
<td>3201–12800</td>
<td>25</td>
<td>25</td>
</tr>
<tr>
<td>&gt;12800</td>
<td>9</td>
<td>9</td>
</tr>
<tr>
<td>Total</td>
<td>109</td>
<td>109</td>
</tr>
</tbody>
</table>

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ELISA were detected as weak positives with the single dilution ELISA.

### 4. DISCUSSION

The commonly used CF test has to be combined with the IFA test for sero-diagnosis of *B. equi* antibodies in doubtful cases and this test has been demonstrated to lack the required specificity and sensitivity [11, 15, 19]. The conventional ELISA for sero-diagnosis of *B. equi* is cumbersome, time consuming and requires large quantities of reagents when the end titre of a sample is to be determined. However, the single dilution ELISA has been found to be cheaper, faster and reliable than conventional serial dilution ELISA as reported for other pathogens [17, 24]. Thus, the single dilution ELISA was developed to predict the *B. equi* antibody titre of a sample. Estimation of end antibody titre by single dilution ELISA could also be useful in monitoring the titres of passively transferred antibodies in young animals [8, 13]. The end antibody titres could also be important in deciding the effects of a drug against *B. equi* infection since antibody
titres are reported to decline when B. equi carrier equids are treated [16, 32].

An advantage of the single dilution ELISA is that the exact antibody titre can be predicted since the colour change (OD) is directly proportional to the amount of antibody present. The P/N ratio was used instead of the OD values at a particular dilution for standardisation of single dilution ELISA since a great variation in OD values for a particular sample is observed depending upon the assay conditions, time of incubation etc. However, the P/N ratio for a particular serum sample at the same dilution remains relatively constant under varying test conditions, as reported previously for standardisation of similar tests for other pathogens [3, 24]. Regression equations as derived for use in single dilution ELISA, express the correlation between the two variables (log10 end titre and P/N ratio) and are necessary in predicting the value of the dependent variable (log10 end titre) when the value of the independent variable (P/N ratio) is known.

We tried in vitro cultivation of the B. equi parasite by the macroaerophilous stationary phase culture (MASP) technique [15, 35] for preparation of antigens but sufficient quantities of the parasite could not be propagated, hence the antigens were prepared from high parasitaemic blood collected from a splenectomised infected experimental donkey. The different antigens used, WM, CM and HSS represented the whole parasite, the cellular fraction and the soluble fraction, respectively [5]. The idea of using three antigen preparations in ELISA was to identify the best antigenic fraction which could be recommended for future use in ELISA. In ELISA, all three antigens detected antibodies against B. equi in test sera and the results were comparable: those samples showing low titres with the WM antigen also had low titres with the CM and HSS antigens. However, the WM antigen was found to be the most sensitive in the detection of antibody titres. The sensitivity and predictive values for the negative tests were also greater when using the WM antigen as compared to the two other antigens. Furthermore, the WM antigen required less processing during preparation and was, therefore, the antigen of choice. This antigen was further used to compare the sensitivity of the serial dilution and single dilution ELISA with the unknown field serum samples.

No cross-reactions were observed with sera positive for B. caballi, Trypanosoma evansi, glanders, strangles, equine infectious anaemia or equine influenza with the three B. equi antigens in ELISA, indicating good immunological specificity of the standardised tests in the present study. Frerich et al. [9, 10] observed cross-reactivity of the B. equi antigen with B. caballi antiserum in CFT, while Weiland [32] reported cross-reactivity of two parasites in ELISA and not in CFT. Zwart and Broklesby [34] reported that the technique adopted for the preparation of the Babesia antigen was crucial and affected the specificity/cross-reactivity in the test. Furthermore Frerich et al. [9] observed a loss of specificity of the B. equi antigen if it was obtained after a number of serial needle passages among animals. Therefore, serial needle passages of the parasite were avoided in the present study and parasitaemia was boosted in the animal by splenectomy leading to immuno-suppression. Improvements in the techniques used for antigen preparation might have helped in reducing non-specific reactions.

Antigenic differences are known to exist between B. equi isolates from various regions [13]. Cross-reactions were observed when B. equi positive antigen/serum of different origin (USDA and Brazilian) were used in IFAT. However, B. caballi positive serum from USDA, USA gave no cross-reaction with the B. equi antigen prepared from the parasite of Indian origin in the present study. We were not able to obtain B. caballi positive serum from equids of Indian origin since
B. caballi has not been reported to be prevalent in India by using OIE [23] recommended serological and biological tests. It seems likely that the Indian isolate of B. equi might be antigenically different from the USDA strain. These observations at least support the fact that single dilution ELISA can be used with more confidence in countries such as India where B. caballi is not prevalent or large numbers of B. caballi positive animals are unlikely to be present.

Serum samples of known identity were deliberately selected from a previous study for the standardisation of serial and single dilution ELISA to increase the confidence of the regression equation so that unknown field sera with different antibody titres could be evaluated perfectly in this system. This was in conformity with earlier studies of Briggs and Skeeles [3]; Manuja et al. [24] and Khatri et al. [17], who adopted a similar strategy while standardising single dilution ELISA for detecting antibodies to other pathogens.

The analogy in antibody titres obtained by serial and single dilution ELISA revealed that the results obtained using the two tests were statistically similar and thus increased the confidence of the regression equation. This was also confirmed graphically (Fig. 3) by the close distribution of log10 antibody titres observed by the serial dilution ELISA and predicted by single dilution ELISA. Non-significant differences between the antibody titres of unknown field sera by serial and single dilution ELISA further validated the suitability of the single dilution ELISA for field surveys. These observations indicate that the test is immunologically and statistically specific and sensitive.

In the present study, the terms statistical sensitivity and statistical specificity indicate the probability of ELISA to accurately identify B. equi infected (true-positive) and non-infected (true-negative) animals, respectively. The predictive value of a positive test result indicates the probability that a test-positive (i.e. ELISA positive) animal really has the disease condition (B. equi infection). The immunological sensitivity of ELISA indicates the ability of the test to detect very low concentrations of antibodies in a positive sample (up to 1:51200 dilution in the present study, Fig. 1).

On the basis of this study, it can be concluded that the single dilution ELISA is as sensitive and specific as serial dilution ELISA in determining the end B. equi antibody titre of a serum sample. It is also more economical and less time consuming, since only one serum dilution is tested. Thus, single dilution ELISA using the WM antigen can be used in the field to detect antibodies to B. equi in test sera and is suitable for mass field epidemiological studies. To derive the regression equation, it is recommended that a statistically significant number of serum samples ranging from low to high titres should be used to validate each newly prepared batch of antigen.

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REFERENCES


Single dilution ELISA using *Babesia equi* antigens


