Serological detection of St. Louis encephalitis virus and West Nile virus in equines from Santa Fe, Argentina

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St. Louis encephalitis virus (SLEV) and West Nile virus (WNV) present ecological and antigenic similarities and are responsible for serious human diseases. In addition, WNV is a significant pathogen in terms of equine health. The purpose of our study was to analyse the seroprevalence of SLEV and WNV in equine sera collected in Santa Fe Province, Argentina. The seroprevalence determined using the plaque reduction neutralisation test was 12.2% for SLEV, 16.2% for WNV and 48.6% for a combination of both viruses. These results provide evidence of the co-circulation of SLEV and WNV in equines in Santa Fe.

Key words: SLEV - WNV - horses - PRNT

St. Louis encephalitis virus (SLEV) and West Nile virus (WNV) belong to the Flavivirus genus of the Flaviviridae family; along with the Japanese encephalitis, Cacipacore, Murray Valley encephalitis, Koutango, Usutu and Yaounde viruses, SLEV and WNV constitute the Japanese encephalitis antigenic complex [Index of Viruses. Flaviviridae (2006) (ncbi.nlm.nih.gov/ICTVdb/Ictv/fs_index.htm)]. SLEV and WNV can cause encephalitis in humans and WNV also causes encephalitis in equines. There is a strong antigenic and epidemiological correlation between these viruses.

SLEV is widely distributed in the Americas (from Canada through Argentina) and serological studies have demonstrated a wide circulation of this virus in temperate and subtropical regions (Sabattini et al. 1998). In the central regions of Argentina, SLEV re-emerged as a human pathogen in 2002 (Spinsanti et al. 2003), with the first epidemic of SLEV human encephalitis occurring in 2005 (Spinsanti et al. 2008). During this outbreak, viral strains belonging to genotype III were isolated from *Culex quinquefasciatus* mosquitoes (Diaz et al. 2006).

Several species of wild birds of the Ardeidae, Columbidae, Fringillidae, Furnariidae, Icteridae, Tyranidae and Phytotomidae families are infected with this virus in Argentina (Monath & Heinz 1996, Diaz 2009). The transmission cycles include *Cx. quinquefasciatus* and

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Culex interfor mosquitoes, which act as the primary vectors and Eared doves (*Zenaida auriculata*) and picui ground-doves (*Columbina picui*), which serve as the urban vertebrate hosts of SLEV (Diaz 2009).

WNV originated in the Old World and was detected for the first time on the American continent in USA during 1999 (Petersen & Roehrig 2001). The introduction of this virus was associated with encephalitis outbreaks in humans and massive deaths among birds, particularly Corvidae (Anderson et al. 1999, Nash et al. 2001).

Since 2004, bird and equine infections have been detected from Canada through the northern areas of South America (Komar & Clark 2006, Bosch et al. 2007). Argentina is one of the few countries outside of USA where WNV strains have been isolated from sick equines and the first records of WNV activity in Argentina were reported in 2006 (Morales et al. 2006); however, seroprevalence studies performed on wild birds confirmed WNV activity at the end of 2004 (Diaz et al. 2008). In Argentina, as elsewhere in the world, WNV is maintained through a bird-mosquito transmission cycle. Although the primary mosquito and avian species involved in WNV transmission are still unknown, *C. picui* could be acting as a host (Diaz et al. 2011).

Other flaviviruses that circulate in Argentina include dengue virus, yellow fever virus, Bussuquara virus (BSQV) and Ilheus virus (ILHV) (Sabattini et al. 1998, Aviles et al. 1999).

The antigenic similarities between different flaviviruses lead to wide cross-reactivity, especially for secondary infections (natural sequential infections by the same or a different flavivirus); this cross-reactivity complicates diagnosis in those regions in which two or more viral species co-circulate. In some cases, higher titres of neutralising antibodies (NTAbs) targeting a specific virus indicate which virus was responsible for the first infection ("original antigenic sin") (Kuno 2003).

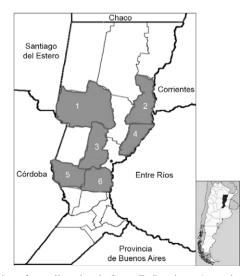
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The purpose of the present study was to analyse the natural co-circulation of SLEV and WNV in equines of Santa Fe Province (Argentina) to provide helpful information to allow better interpretation of the serological data obtained from surveillance programmes.

Study site - Santa Fe, located in the central area of Argentina, is divided into 19 counties and has with a total surface of 133.007 km²; the total population exceeds three million inhabitants. The local weather, particularly with regard to temperature and rainfall, constitutes a transition between tropical and temperate climates. The economy is based on agriculture and livestock activities.

Sample collection - Equine (horse) sera from six different counties in Santa Fe were analysed (Figure). All of the samples were obtained between September-November of 2008. The blood samples were collected from the jugular vein of asymptomatic animals and kept at 4°C until being processed.

Plaque reduction neutralisation test (PRNT) - NTAb detection was performed using a PRNT with Vero cells (African green monkey kidney cells) (Early et al. 1967) and the serum samples that neutralised at least 80% of the inoculated viral plaques were considered positive. SLEV strain 78V-6507 obtained from Cx. quinquefasciatus mosquitoes from Santa Fe and WNV E/7229/06 isolated from sick equines in Buenos Aires were used (Morales et al. 2006). Viral suspensions were prepared using the brains of infected lactating mice and 10% (weight/volume) minimal essential medium with Earle salts supplemented with L-glutamine, non-essential amino acids, 10% foetal bovine serum and 1% antibiotics. The suspensions were centrifuged at 10,000 rpm for 30 min using a refrigerated centrifuge. Based on new serological evidence provided by Ledermann et al.



Location of sampling sites in Santa Fe Province, Argentina. 1: San Cristóbal; 2: San Javier; 3: Las Colonias; 4: Garay; 5: San Martin; 6: Sam Jerónimo.

(2011) and Patiris et al. (2008), the serum samples with antibody titres higher than 20 were considered positive for the tested viruses.

All of the samples were also subjected to PRNTs for ILHV (Be H7445) and BSQV (Be AN 4116).

For the statistical analysis, the chi-square test of independence was applied using InfoStat program v. 2010.

A total of 57 of the 74 analysed serum samples were positive for WNV, SLEV or both viruses. The monotypic seroprevalence values were 12.2% (9/74; titre range 20-1280) for SLEV and 16.2% (12/74; titre range 20-640) for WNV. In addition, 48.6% (36/74) of the equines displayed antibodies for both of the viruses (SLEV + WNV) (Table). No significant differences between the prevalence of the viruses were found with respect to the monotypic infections ($\chi 2 = 1.51$; p = 0.2192). All of the sera were negative for ILHV and BSQV.

The prevalence values were not analysed according to the place of origin of the equines due to the scarcity of representative values for most of the sampled locations.

SLEV is endemic in the north-central regions of Argentina, coexisting with the recently introduced WNV (Díaz et al. 2008). Although WNV had been previously isolated from sick and dead equines in Argentina, these cases were all from Buenos Aires; for this reason, the information presented herein is the first record of the equine seroprevalence in Santa Fe, widening the known range of distribution of WNV in Argentina.

During the 1980s, Monath et al. (1985) reported an NTAb screening test that demonstrated an SLEV prevalence of greater than 50% among equines from different provinces, including Santa Fe; however, there is no recent report of arbovirus circulation in this province.

Our studies found a 12.2% seroprevalence for SLEV, which is lower than the previously reported values (Monath et al. 1985). The seroprevalence of WNV was similar (16.2%) and among animals that were positive only for antibodies against WNV, the titres were lower than those detected for SLEV, indicating patterns of primary response. These seroprevalence values could be underestimates due to the alternative criteria of positivity used in the present study.

Forty-eight percent of the analysed samples exhibited serological profiles compatible with a heterotypic serological response. According to the Centers for Disease Control and Prevention's guidelines, a titre difference of > four-fold is used to identify the etiologic agent; when a four-fold difference is not observed, the specific etiological agent cannot be defined (Table). However, this criterion is insufficient to diagnose the etiological agent in some scenarios of arboviral infection; indeed, it is a frequent error to assume that the infecting virus is the one for which higher antibody titres are obtained in two or more dilutions (Kuno 2003). In chickens and equines, Patiris et al. (2008) and Ledderman et al. (2011), respectively, observed that an initial SLEV infection followed by a WNV infection greatly amplified the antibody levels against SLEV, precluding the detection of the subsequent WNV infection using the four-fold titre difference criterion. These authors also demonstrated that crossreaction among heterologous antibodies is not frequent

in primary infections. Based on these facts, we consider that the infectious agent can indisputably be identified only in monotypic infections. Moreover, we observed that a heterotypic immunological response can be the result of the presence of sequential infections by the same or different viruses (SLEV or WNV) (Table), most often in those animals that presented titres > 160 for both vi-

TABLE

Heterotypical serological response by plaque reduction neutralisation test (PRNT) against St. Louis encephalitis virus (SLEV) and West Nile virus (WNV) in horse samples collected in Santa Fe Province, Argentina

	PRNT (80%)		Interpretation criteria	
Code	SLEV	WNV	CDC's	Alternative
270512	160	640	WNV	SLEV + WNV
27116	20	80	WNV	SLEV + WNV
271531	160	20	SLEV	SLEV + WNV
270871	160	640	WNV	SLEV + WNV
271546	160	20	SLEV	SLEV + WNV
1786	80	320	WNV	SLEV + WNV
1788	80	320	WNV	SLEV + WNV
1789	80	80	Inconclusive	SLEV + WNV
1790	1280	160	SLEV	SLEV + WNV
1794	20	40	Inconclusive	SLEV + WNV
1798	640	40	SLEV	SLEV + WNV
1799	640	40	Inconclusive	SLEV + WNV
1800	160	80	Inconclusive	SLEV + WNV
1801	40	1280	WNV	SLEV + WNV
1802	20	1280	WNV	SLEV + WNV
1812	20	1280	WNV	SLEV + WNV
1813	320	160	Inconclusive	SLEV + WNV
1815	80	1280	WNV	SLEV + WNV
1816	20	640	WNV	SLEV + WNV
1817	20	20	Inconclusive	Inconclusive
1818	80	20	SLEV	SLEV + WNV
1819	80	640	WNV	SLEV + WNV
1826	20	1280	WNV	SLEV + WNV
1827	80	160	Inconclusive	SLEV + WNV
1828	20	20	Inconclusive	Inconclusive
1831	20	80	WNV	SLEV + WNV
41	20	20	Inconclusive	Inconclusive
43	160	160	Inconclusive	SLEV + WNV
40	20	20	Inconclusive	Inconclusive
19	20	20	Inconclusive	Inconclusive
27	20	80	WNV	SLEV + WNV
23	80	20	SLEV	SLEV + WNV
10	20	20	Inconclusive	Inconclusive
8	640	80	SLEV	SLEV + WNV
35	640	1280	Inconclusive	SLEV + WNV
12	640	320	Inconclusive	SLEV + WNV

CDC: Centers for Disease Control and Prevention.

ruses or in animal infected with another antigenically related flavivirus (i.e., sample numbers 1789, 1817, 1828 and 41) (Table). Ledermann et al. (2011) reported high titres of antibodies (160-1280) with considerable cross-reactions and a highly heterogeneous response in more than 70% of the equines previously inoculated with SLEV and then challenged with SLEV or WNV.

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These results demonstrate the difficulties encountered in the interpretation of the data and showed that the equines had been exposed to both viruses, with high levels of high cross-reactivity. Thus, the analysis of a single sample may provide misleading or inaccurate results regarding the etiologic agent, underscoring the importance of the timing of sample collection and of multiple-sample testing. To better estimate the level of recent circulation of these viruses in the future, surveillance should be based on IgM assays.

In conclusion, the circulation of more than one flavivirus in Santa Fe indicates that the surveillance of these infectious agents in Argentina should be intensified.

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