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Antigenic and molecular characterization of rabies virus in Argentina

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Abstract

The nucleoprotein genes of 54 human, domestic and wild animals rabies isolates obtained in Argentina between 1995 and 2002 were characterized using monoclonal antibodies and partial gene sequence analysis. The antigenic and genetic diversities of rabies virus in samples from bat and bat-related cases were studied, leading to the identification of five distinct genetic variants. Rabies viruses isolated from vampire bat related cases were very similar to each other, showing 98.9% overall similarity. Specific antigenic variants (AgV) were detected associated with different insectivorous bats species, in samples from *Tadarida brasiliensis* and *Eumops patagonicus* bats. In contrast, isolates from *Myotis* sp. and *Histiotus* sp. bats could not be matched to any antigenic type. Additionally, bat rabies cases were also detected in southern provinces previously considered rabies-free. Finally, two independent antigenic and genetic variants co-circulating in northern Argentina were found in isolates obtained from dogs and dog-related cases, suggesting two independent cycles of virus transmission.

This is the first national coordinated study of antigenic as well as molecular epidemiology of rabies in Argentina. The information presented here will improve our knowledge about rabies epidemiology and therefore, will assist preventing fatal human cases. © 2004 Elsevier B.V. All rights reserved.

Keywords: Rabies; Antigenic variant; Molecular epidemiology

1. Introduction

Rabies is a fatal infection of the central nervous system generally acquired through virus-contaminated saliva transmitted by the bite of a rabid animal. The virus circulates through two different epidemiological cycles: urban rabies, where the domestic dog is the main reservoir and transmitter, and sylvatic rabies, where several wildlife species act as reservoirs and/or transmitters. The prototype of rabies virus (RV), lyssavirus, contains a non-segmented 12 Kb negative-sense RNA genome that encodes for five genes: N, P, M, G and L (Tordo et al., 1986). The highly conserved and abundant nucleoprotein (N), a key structural component of the viral ribonucleoprotein core essential to viral propagation (Yang et al., 1998), constitutes the main target for rabies diagnosis (Dean, 1996).

Analysis of rabies isolates from human and animal cases using monoclonal antibodies (Mabs) directed to the nucleoprotein, provides useful information about the geographical and temporal distribution of viral variants associated with outbreaks of rabies. A previous study conducted in 17 several

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Table 1
Distribution of rabies isolates in Argentina

Isolates	Species	Location	Date	Antigenic variant	Genetic varian
Bamsbt121	Myotis sp bat	3 de febrero, Buenos Aires	2001	ND	A5
Batbbt122	Tadarida brasiliensis bat	La Matanza, Buenos Aires	2001	4	A2
Batbbt123	Tadarida brasiliensis bat	3 de febrero, Buenos Aires	2001	4	A2
Batbbt125	Tadarida brasiliensis bat	3 de febrero, Buenos Aires	2001	4	A4
Chabv129	Cattle	Gral San Martín, Chaco	2002	3	A1
Chabv66	Cattle	San Martin, Chaco	2000	3	A1
Chabv72	Cattle	Puerto Veloz, Chaco	2000	3	A1
Chabv74	Cattle	Colonia Mixta, Chaco	2000	3	A1
Chabv76	Cattle	La Leonesa, Chaco	2000	3	A1
Chabv78	Cattle	Colonia Mixta, Chaco	2000	3	A1
Chabv86	Cattle	La Leonesa, Chaco	2001	3	A1
Chabv94	Cattle	Las Palmas, Chaco	2001	3	A1
Chadg120	Dog	Pampa Almiron, Chaco	2001	2	Bb1
Chafx08	Fox	Laguna Limpia, Chaco	1999	NA	A1
Chafx119	Fox	Pampa Almiron, Chaco	2001	2	Bb1
Chahr90	Horse	Colonia Mixta, Chaco	2001	3	A1
Chutbbt124	Tadarida brasiliensis bat	Puerto Madryn, Chubut	2001	4	A2
Ctebv01	Cattle	Beron de Estrada, Corrientes	1999	3	A1
Ctebv55	Cattle	Colonia Romero, Corrientes	2000	3	A1
	Cattle	Concepción Corrientes	2000	3	A1
Ctebv77 Ctebv79	Cattle	San Miguel, Corrientes	2000	3	A1 A1
	Human	÷			
Ctehm82		San Luis del Palmar, Corrientes	2001	3	A1
Forbv07	Cattle	Laishi, Formosa	1999	3	A1
Forbv107	Cattle	Potrero de los caballos, Formosa	2001	3	A1
Fordg53	Dog	Pozo del Tigre, Formosa	2000	2	B1b
Fordg71	Dog	Pirane, Formosa	2000	2	B1b
Forhr96	Horse	Presidente Irigoyen, Formosa	2001	3	A1
Jujdg40	Dog	La Quiaca, Jujuy	1997	2	B1a
Salct41	Cat	Oran, Salta	1998	1	B2
Salct47	Cat	Oran, Salta	1999	1	B2
Salct49	Cat	Oran, Salta	1999	1	B2
Salct50	Cat	Oran, Salta	1999	1	B2
Saldg126	Dog	Oran, Salta	2002	1	B2
Saldg146	Dog	Oran, Salta	2002	1	B2
Saldg15	Dog	Hipolito Irigoyen, Salta	1999	1	B2
Saldg17	Dog	Oran, Salta	1999	1	B2
Saldg19	Dog	Ingeniero Tabacal, Salta	1999	1	B2
Saldg20	Dog	Hipolito Irigoyen, Salta	1999	1	B2
Saldg21	Dog	Hipolito Irigoyen, Salta	2000	1	B2
Saldg42	Dog	Oran, Salta	1999	1	B2
Saldg43	Dog	Oran, Salta	1999	1	B2
Saldg44	Dog	Oran, Salta	1999	1	B2
Saldg45	Dog	Oran, Salta	1999	1	B2
Saldg46	Dog	Rio Blanco, Salta	1999	1	B2
Saldg48	Dog	Hipolito Irigoyen, Salta	1999	1	B2
Saldg51	Dog	Hipolito Irigoyen, Salta	1999	1	B2
Sfeepbt118	Eumops patagonicus bat	Rosario, Santa Fe	2001	4	A2
Sfemnbt116	Myotis nigricans bat	Rosario, Santa Fe	2001	- ND	A5
Stchmbt80	Histiotus montanus bat	Rio Turbio, Santa Cruz	2001	ND	A3
Tucdg35		Concepcion, Tucuman	1995	2	B1a
Tucdg35 Tucdg36	Dog	Aguilares, Tucuman	1993	2	Bla
-	Dog				
Tucdg37	Dog	Rio Seco, Tucuman	1995	2	B1a P1o
Tucdg38	Dog	Aguilares, Tucuman	1996	2	B1a
Tucdg39	Dog	Chigliasta, Tucuman	1996	2	B1a

NA: Not available; ND: not determined.

Latin American and Caribbean countries, identified eight distinct antigenic variants (AgV) of rabies virus, concluding that different animal reservoirs might have been responsible for maintenance and transmission of these variants (Diaz et al., 1994). In addition, molecular characterization of rabies virus isolates enables the identification of epidemiologic links that cannot be otherwise established by antigenic typing (de Mattos et al., 1996, 1999, 2000).

Dog vaccination programs successfully reduced the number of cases of rabies in the northern and center provinces of Download English Version:

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