

LETTER TO THE EDITOR

INFLUENZA VIRUS SURVEILLANCE BY THE INSTITUTO ADOLFO LUTZ, INFLUENZA SEASON 2014: ANTIVIRAL RESISTANCE

São Paulo, August 4, 2014

Dear Sir,

Neuraminidase (NA) inhibitors (NAIs) are the only antivirals that are effective for prophylaxis and the treatment of seasonal influenza A and B infections. There are currently two NAIs approved in most countries: oseltamivir (Tamiflu; F. Hoffmann - La Roche) and zanamivir (Relenza; GlaxoSmithKline plc.). The development of drug resistance is a major drawback for any antiviral therapy, and these specific anti-influenza drugs are not excluded from this rule. Thus, the proper use of NAIs and worldwide monitoring for the presence and spread of drug resistant influenza viruses are of the utmost importance. The existence of a global surveillance network for influenza, underpinning vaccine strain selection, is a valuable asset when seeking to track the emergence of antiviral resistance.

The Instituto Adolfo Lutz, São Paulo, SP, Brazil, plays a role in national and global influenza surveillance. The objective of the present study was to monitor antiviral resistance to assist public health authorities with decisions regarding prophylaxis and treatment strategies.

Using the real time polymerase chain reaction assay (rRT-PCR), influenza viruses of type A, subtype H1N1pdm09 and H3N2, as well as type B viruses, were identified and antiviral resistance testing was conducted using pyrosequencing² and Sanger dideoxy sequence analysis⁵.

Prior to the emergence of the pandemic virus in 2009, the presence of the oseltamivir resistance-conferring marker, H275Y, was identified in seasonal influenza A (H1N1). In 2014, influenza virus surveillance identified the same marker, H275Y, in an influenza A (H1N1) pdm09 strain isolated from a 20 year-old pregnant woman living in Mato Grosso/Cuiabá, the Midwest region of Brazil. The virus was collected in March 2014. In addition, two permissive secondary NA mutations; V241I and N369K were detected in the virus isolated in the Midwest region of Brazil¹. These mutations are known to negate the impact of the NA H275Y oseltamivir resistance mutation on viral replicative fitness. This patient was treated with oseltamivir, rocephin, azithromycin and made a full recovery from the respiratory disease.

The choice of assay for assessing the susceptibility of the influenza virus to NAIs depends on factors pertaining to appropriateness of the setting, cost, sustainability, speed in obtaining valid results, reliability in terms of predictive values, and accessibility. The high sensitivity of genotypic assays allows for testing of clinical specimens, thus eliminating the need for virus propagation in cell culture. In addition, rapid genotypic testing facilitates more appropriate patient management and can significantly advance and assist in large-scale epidemiological

studies of drug-resistant variants⁴.

Katia Corrêa de Oliveira SANTOS (1)
Daniela Bernardes Borges da SILVA (1)
Margarete Aparecida BENEÇA(1)
Renato de Sousa PAULINO(1)
Elián Reis E SILVA Jr(1)
Dejanira dos Santos PEREIRA(2)
Aparecida Duarte Hg MUSSI(2)
Valéria Cristina da SILVA(1,3)
Larissa V. GUBAREVA(4)
Terezinha Maria de PAIVA(1)

(1) Laboratory of Respiratory Viruses NIC/WHO,
Instituto Adolfo Lutz, São Paulo, SP, Brazil.

(2) Central Laboratory of Public Health of the
Mato Grosso State, Cuiabá, Brazil

(3) Epidemiologic Surveillance Center of the
Mato Grosso State, Cuiabá, Brazil.

(4) Molecular Epidemiology Team Virus Surveillance
and Diagnosis Branch Influenza Division,
Centers for Disease Control and Prevention, Atlanta, GA, USA

Correspondence to: Terezinha M. de Paiva, Instituto Adolfo Lutz,
Núcleo de Doenças Respiratórias

Av. Dr. Arnaldo 355, 01246-902 São Paulo, SP, Brasil

Phone: 55 11 30682913, Fax: 55 11 30853505

E-mail: tterezinha@uol.com.br

REFERENCES

1. Butler J, Hooper KA, Petrie S, Lee R, Maurer-Stroh S, Reh L, *et al.* Estimating the fitness advantage conferred by permissive neuraminidase mutations in recent oseltamivir - resistant A (H1N1) pdm09 influenza viruses. *PLOS Pathog.* 2014;10(4):e1004065.
2. Deyde VM, Gubareva LV. Influenza genome analysis using pyrosequencing method: current applications for a moving target. *Expert Rev Mol Diagn.* 2009;9:493-509.
3. Marx C, Gregianini TS, Lehmann FK, Lunge VR, Carli SD, Dambrós BP, *et al.* Oseltamivir-resistant influenza A(H1N1)pdm09 virus in southern Brazil. *Mem Inst Oswaldo Cruz.* 2013;108:392-4.
4. Okomo-Adhiambo M, Sheu TG, Gubareva LV. Assay for monitoring susceptibility of influenza viruses to neuraminidase inhibitors. *Influenza Other Respir Viruses.* 2013;7(Suppl 1):44-9.
5. Sheu TG, Deyde VM, Okomo-Adhiambo M, Garten RJ, Xu X, Bright RA, *et al.* Surveillance for neuraminidase inhibitor resistance among human influenza A and B viruses circulating worldwide from 2004 to 2008. *Antimicrob Agents Chemother.* 2008;52:3284-92.
6. Souza TM, Resende PC, Fintelman-Rodrigues N, Gregianini TS, Ikuta N, Fernandes SB, *et al.* Detection of oseltamivir-resistant pandemic influenza A(H1N1)pdm2009 in Brazil: can community transmission be ruled out? *PLOS One.* 2013;8(11):e80081.