

The occurrence of *Orthoreovirus*, *Rotavirus* and chicken anemia virus in chickens of the poultry industry in Minas Gerais, Brazil

[Ocorrência de *Orthoreovirus*, *Rotavirus* e vírus da anemia das galinhas em frangos de corte da indústria avícola de Minas Gerais]

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ABSTRACT

Fifty-four fecal samples taken from broiler chickens from 1 to 45 days of age, and of pullets from 10 to 13 weeks of age, original from eight different poultry regions in the state of Minas Gerais, Brazil, were collected from March 2008 to January 2010 for avian *Orthoreovirus* (ARV) and avian *Rotavirus* (AvRV) analyses. For the assay of ARV, RNA was immediately extracted (Trizol®) and transcribed into cDNA for assaying in a nested-PCR with ARV-specific primers. For AvRV, polyacrylamide gel electrophoresis (PAGE) was performed with RNA extracts obtained by phenol-chloroform extraction. CAV was additionally investigated through a nested-PCR of thymus and spleen. Results found 5.55% positive for ARV and 9.25% for AvRV. Also, CAV and ARV genomes were detected in co-infection, in a highly prostrated and claudicating chicken flock. No ARV or AvRV infections were detected in pullets. Material of a clinically affected flock was inoculated into SPF embryos, resulting in embryonic hemorrhage, whitish foci in the chorio-allantoic membrane and death. Sequencing of ARV amplicons and isolate cDNA grouped local strains with the ARV S1133 strain, historically used in live vaccines, suggesting the continued circulation of this vaccine virus strain in intensive poultry regions. Detection rates for ARV and AvRV, as well as the presence of CAV, were additionally indicative of failing biosecurity strategies for the intensive poultry regions examined.

Keywords: Broiler, layer, chicken anemia virus, CAV, avian reovirus, ARV, avian rotavirus, AvRV, nested-PCR, PAGE.

RESUMO

Avaliou-se a ocorrência de *Orthoreovirus* (ARV) e *Rotavirus* (AvRV) aviários na avicultura industrial de Minas Gerais. Foram colhidas cinquenta e quatro amostras de fezes de frangos de corte entre um e 45 dias e de frangas de postura de 10 a 13 semanas de idade. Para análise de ARV, o RNA foi imediatamente extraído (Trizol), transcrito em cDNA e avaliado em uma PCR com oligonucleotídeos iniciadores específicos para ARV. Para a investigação de AvRV, os extratos de RNA foram obtidos por fenol-clorofórmio e submetidos à eletroforese em gel de poliacrilamida. Todas as amostras foram também avaliadas para o DNA do vírus da anemia das galinhas (CAV) em uma nested-PCR específica. Em frangos de corte, a positividade encontrada para ARV foi de 5,55% e para AvRV de 9,25%. CAV foi detectado em coinfeção em um plantel com refugagem, claudicação e prostração. Nenhuma amostra de poedeiras foi positiva para ARV ou AvRV. Material de plantel com sinais clínicos foi purificado e inoculado em ovos SPF embrionados, sendo obtidas lesões hemorrágicas e focos brancos na membrana cório-alantóide. O sequenciamento dos produtos de PCR e de embrião agrupou os isolados de ARV com a estirpe S1133, historicamente usada como vacina viva. Os resultados sugerem a continuada circulação

Recebido em 9 de março de 2011

Aceito em 6 de setembro de 2012

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da infecção por estirpes assemelhadas a ARV S1133 nas regiões de avicultura industrial. Os índices de detecção de ARV, AvRV e CAV indicam que a intensificação nas regiões produtoras tem resultado em falhas de biossegurança.

Palavras-chave: Frango de corte, poedeira, reovirus, rotavirus, vírus da anemia das galinhas, nested-PCR, PAGE.

INTRODUCTION

Orthoreovirus and *Rotavirus* are genera of *Reoviridae*, which in chickens and other species are usually involved in enteritis, stunted and poorer growth, reduced productive performance and higher mortality rates (McFerran *et al.*, 1983; Andral *et al.*, 1985; Gouch *et al.*, 1988; Decaesstecker *et al.*, 1988; Martins *et al.*, 2009). The denomination reovirus derives from an acronym of respiratory and enteric orphan (REO), due to early isolated viruses not being associated to a clinical condition, the recent suffix *ortho* added to designate the genus for early isolates. Viruses of *Reoviridae* contain double stranded RNA divided, for *Orthoreovirus* into 10 segments or for *Rotavirus* into 11 segments. Virions are characterized by particles of about 100 nm in diameter, with triple-layered icosahedral protein capsid, with 60 external capsid spikes, an external capsid which requires calcium for integrity and virions which contain RNA-dependent RNA polymerase. Avian orthoreoviruses are mostly fusogenic of cell cultures and the S gene sigma (σ) C protein (outer capsid) acts as cell-attachment protein, hemagglutinin and primary serotype determinant (Shiff *et al.*, 2007). Multiple syndromes are described resulting from ARV infections. Older birds are more resistant and remain infected for long periods of time, acting as source for the more susceptible young. Infected breeders transmit ARV vertically, through the egg or horizontally, as for AvRV, via fecal-oral route (Jones *et al.*, 1975). Infected hatchlings and young develop the gastro-intestinal signs which will result in poor growth and lower performance, and survivors may develop arthritis (Van der Heide *et al.*, 1975). A novel *Orthoreovirus* was isolated from the necrotic liver of affected *Anas platyrhynchos* (Pekin ducks) flocks with up to 40% mortality (Chen *et al.*, 2012). For AvRV, the horizontal challenge, due to the high titers available in feces, up to 10^{10} a 10^{12} viral particles per gram (Flewett and Woode, 1978) will result in similarly poor performance in chickens and turkeys mostly

from 6 to 14 days of age (McNulty *et al.*, 1980). The virus appears to be shed mainly from the intestine and for longer periods, suggesting fecal-oral contamination as a principal mode of infection (Rosenberger, 2003). For the diagnosis of ARV, specific nested-polymerase chain reaction (PCR) protocols have been described for amplifying a region of the S1 gene encoding the σ C protein (Liu *et al.*, 1997) region best suited for demonstrating diversity ((Shiff *et al.*, 2007). Nested PCR reactions reduce or eliminate non-specificity and increase sensitivity for ARV detection (Liu *et al.*, 1997).

PAGE (polyacrylamide gel electrophoresis) of AvRV will separate the 11 segments of the viral RNA according to the molecular sizes (McNulty *et al.*, 1980). Previous studies were reported in Minas Gerais (Alfieri *et al.*, 1988; Rios *et al.*, 2010). This study was proposed for the evaluation of the occurrence of ARV and AvRV, as well as CAV, for defined flocks in poultry regions in Minas Gerais.

MATERIALS AND METHODS

The geographical locations of the sampling areas were distributed to evaluate some of the most important poultry areas (Table 1). Fifty-four flocks of broiler or layer chickens were sampled from March 2008 to January 2010. Flocks were selected at random and included normal and diarrheic/clinically ill. Fecal samples were evenly distributed per house (n=20), totaling 1,080 samples. Samples were pooled for each flock for analyses. Broilers were from 1 to 45 days of age and pullets were from 10 to 13-week of age. Each sample was collected into a new autoclaved flip top tube (1.5mL) and frozen at -70 °C until the preparation of pools. A dilution factor was established for feces during standardization, by serially diluting (base 10) the ARV vaccine mixed with SPF chicken feces. A 10^{-2} dilution was chosen for studying all field samples.

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The RNA extraction (Trizol®) was based on a protocol designed for infectious bursal disease virus (IBDV) (Gomes *et al.*, 2005). The quantified RNA was immediately transcribed (300ng) into ARV cDNA using reverse transcriptase (M-MLV, PROMEGA, USA) according to the manufacturer, with primer 5' ATTGAATTCTCTGTTATCTCAACCTTG 3'. For the first ARV PCR reaction, the external primer oligonucleotides was S1C 5' ATTGAATTCTCTGTTATCTCAACCTTG 3' and S1D 5' AAGGAATTCGTTGAGAACAGAAGTAGG 3', described for amplifying the S1 gene region encoding the σ C protein (Liu *et al.*, 1997), in reactions with 50 μ L volumes. In each assay, the cDNA transcribed from RNA of Poulvac®Maternavac IBD-Reo *Orthoreovirus* strain (Fort Dodge) was used as positive control. In order to prepare the positive control, a sample of reconstituted vaccine (10 μ L) was mixed with the feces (500 mg) of SPF chickens for extraction. As negative controls, reactions with all reagents and without cDNA were used. ARV nested-PCR reactions were performed using the amplicon of the first reaction as a template and the internal oligonucleotides S1E 5'-TCTGAATTCATCGCAGCGAAGAGAGGT CG-3' and S1F 5'-AGTGAATTCAGTAT CGCCGCGTGCGCAG-3' as primers (Liu *et al.*, 1997). Electrophoresis in 1% agarose stained by ethidium bromide and visualized in ultraviolet transillumination was used for the visualization of the 738bp amplified amplicons of the first reaction and 342bp of the nested reaction.

One of the ARV PCR positive pooled fecal samples was suspended in equal parts sterile PBS and clarified at 2.000xg/10min. The supernatant was collected and treated with penicillin G (1,000 units/mL), streptomycin (500 μ g/mL) and amphotericin B (1 μ g/mL) for 30min (25 °C) and inoculated into the 10th day of incubation specific pathogen free (negative for *Orthoreovirus*) chicken eggs (n=10) via the chorioallantoic membrane (CAM). Eggs were candled and death was recorded daily. Dead embryos up to 24 hours post inoculation were discarded and the others examined for lesions and tissues collected for PCR. Harvested CAM were macerated, treated with antibiotics and passed in eggs three times to demonstrate embryonic effects and PCR detection.

DNA extracts were used for nested-PCR reactions specific for chicken anemia virus (CAV), as described previously (Cardona *et al.*, 2000; Barrios *et al.*, 2009).

Amplified PCR products (PTC-100, MJ Research, Inc., USA) were sequenced by chain termination (dideoxynucleotide), as described previously (Sanger *et al.*, 1997), in an automated capillar sequencer (ABI 377, Perkin Elmer, USA) and *Big Dye Terminator Mix* (Applied Biosystems, USA), according to the manufacturer. Each sample was sequenced in both directions, as many times as needed for highest certainty, the quality of sequences evaluated using Bioedit (Hall, 1999) and compared to data at the *National Center for Biotechnology Information* (NCBI – <http://www.ncbi.nlm.nih.gov/>). Four independent PCR reactions (*BigDye* labeled) were performed, twice for the forward primer and twice for the reverse primer, for sequencing each product.

The alignment of cDNA and deduced amino acid sequences were processed using Clustal W version 1.6 (*Molecular Evolutionary Genetics Analysis* – MEGA 3.1/ www.megasoftware.net) version 3.1 for Windows. The alignments of gene coding for ARV protein σ 1 were used for preparing phylogenetic trees (MEGA 3.1 www.megasoftware.net).

For the diagnosis of AvRV, polyacrylamide gel electrophoresis (PAGE) of phenol-chloroform-isoamyl alcohol (25:24:1) extracted fecal RNA was stained with silver nitrate (Herring *et al.*, 1982; Alfieri, 1989; Alfieri, 1992). Stored fecal samples (-20°C) were thawed, separately homogenized with disposable plastic sticks and diluted (1:7) in a *Rotavirus* buffer (TERV) (pH 7.4 -Tris-HCl 50mM; NaCl 10mM; CaCl₂ 13mM; β -mercaptoethanol 1.5mM). Dilutions were mixed (vortex, GENIE BENDER, model K-550G, USA) and clarified at 2,000 x g/15min (FANEM, model 240-N, Brazil). Supernatants were transferred to 1.5mL flip-top microtubes for RNA extraction. Approximately 300 nanograms of total RNA were applied to the 3.75% acrylamide (acrylamide/bisacrylamide) stacking gel. Electrophoresis was run in 7% resolving gel in Tris-glycine (Tris 25mM, glycine 0,2M, pH8,6), with constant voltage (100 V) during 4 hours. Gels were fixed in ethanol-acetic acid, stained with silver nitrate (0,11M), developed in

NaOH-formaldehyde and preserved in 20% ethanol (4°C).

This study was approved by the Ethics Committee in Animal Experimentation, certificate 074/2007 of August 8th, 2007.

RESULTS AND DISCUSSION

ARV and AvRV were detected in the broiler industry in Minas Gerais (Table 1). Three out of 54 broiler flocks, at 8, 12 and 31 days-old, the first and third clinically affected, were positive for amplicons compatible in molecular mass to the expected ARV product. Broilers were of the intensified regions of Divinópolis, Pará de Minas and São Sebastião do Oeste. Five out of 54 (9.25%) flocks showed the typical 11-band RNA segments by PAGE, indicative of AvRV, present in the regions of Pará de Minas, São Sebastião do Oeste and Pitangui. Two out of 3 flocks (66%) were positive for ARV and showed enteric signs and one positive flock was clinically normal. All AvRV positive flocks were clinically affected (diarrhea). Considering the overall clinically affected flocks (n=7), the relative indexes for ARV (42%) and AvRV (57%) are indicative of major roles for these viruses. Failed or lack of biosecurity, intensification and proximity of flocks of broiler chickens, may have a role in the detected occurrence of ARV and AvRV, as samples were obtained from chickens of densely broiler populated areas, and susceptible floor raised flocks may facilitate fecal-oral infection (Rosenberger, 2003).

For ARV positive flocks, signs of uneven growth (2/3) and lameness (1/3) were noted. ARV has been associated with stunting of young birds, chickens with less than 15 days of age (Ruff and Rosenberger, 1985), supporting the finding of positive 8-day old broilers. The slightly higher ARV occurrence (13.5%) previously detected by PAGE in broilers with diarrhea in Paraná state (Tamehiro *et al.*, 2003) may be due to biosecurity differences. No inference could be drawn toward the role of ARV or AvRV in the clinical aspects (diarrhea) of flocks. In agreement, two Dutch and one German ARV derived from malabsorption syndrome did not cause weight gain depression in the broilers, although lesions in the small intestine were present from day 1 to day 4 PI and were more severe in broilers than in the white leghorn

chickens (Songserm *et al.*, 2003). However, infection with *Orthoreovirus* strain ARV-CU98 (10 pfu/ml/chick) in broilers resulted in depressed growth, increased mortality and reduced protein concentration in various tissues (Burgos *et al.*, 2006).

A co-infection of ARV and CAV (chicken anemia virus) was detected in the 31-day-old flock, with CAV present in the thymus and spleen. A previous CAV study in the region has shown disseminated infection in broilers (data not shown). CAV mostly causes cellular immune depression in chickens, and may have a role in the clinical effects of prostration and anorexia, in addition to arthritis and diarrhea observed in the flock. Immune compromised flocks will present the most severe effects (McNeilly *et al.*, 1995; Rosenberger, 2003). Field conditions will enable co-infections with a range of primary and opportunistic pathogens (Andral *et al.*, 1985), and the co-infection ARV / CAV may enable sufficient immune depression for an evolutive advantage of ARV variants. However, this is the first study detecting ARV and CAV natural co-infection in Brazil.

ARV was previously reported in chickens with enteric problems in Brazil (Alfieri *et al.*, 1988; Alfieri *et al.*, 1989; Tamehiro *et al.*, 2003; Villareal *et al.*, 2006), and in several countries such as United States, Australia, Sweden, England and Germany (Apple *et al.*, 1991; Lenz *et al.*, 1998). ARV has also been described as affecting growth and causing mortality in broilers by infecting the digestive system, and also the heart and articulations, mainly the hock joint (Rosenberger, 2003).

The embryo isolated strain caused 70% mortality of embryos at 24, 48 and 72h post infection at the third passage and was successfully detected by PCR. Chorioallantoic membranes were swollen and with whitish foci and embryos presented with superficial, hepatic and renal hemorrhages, and hepatic enlargement. Negative control eggs did not show lesions and were negative for ARV by PCR.

The sequencing of ARV PCR products has enabled comparisons with the data available (GenBank, NCBI, USA). ARV sequences were grouped (77%) with S1133-derived strains (Figure 1). The attenuated S1133 vaccine strains

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were used for a few decades for vaccinating breeding flocks in Brazil and the virus may have been maintained through cyclic infections among flocks in densely populated areas, and might have reversed in pathogenesis causing the

localized arthritis lesions. However, a large degree of variation in the N-terminal region of ARV σ C when USA and Australian isolates were compared and correlation among USA isolates was consistent with serotyping (Liu *et al.*, 1997).

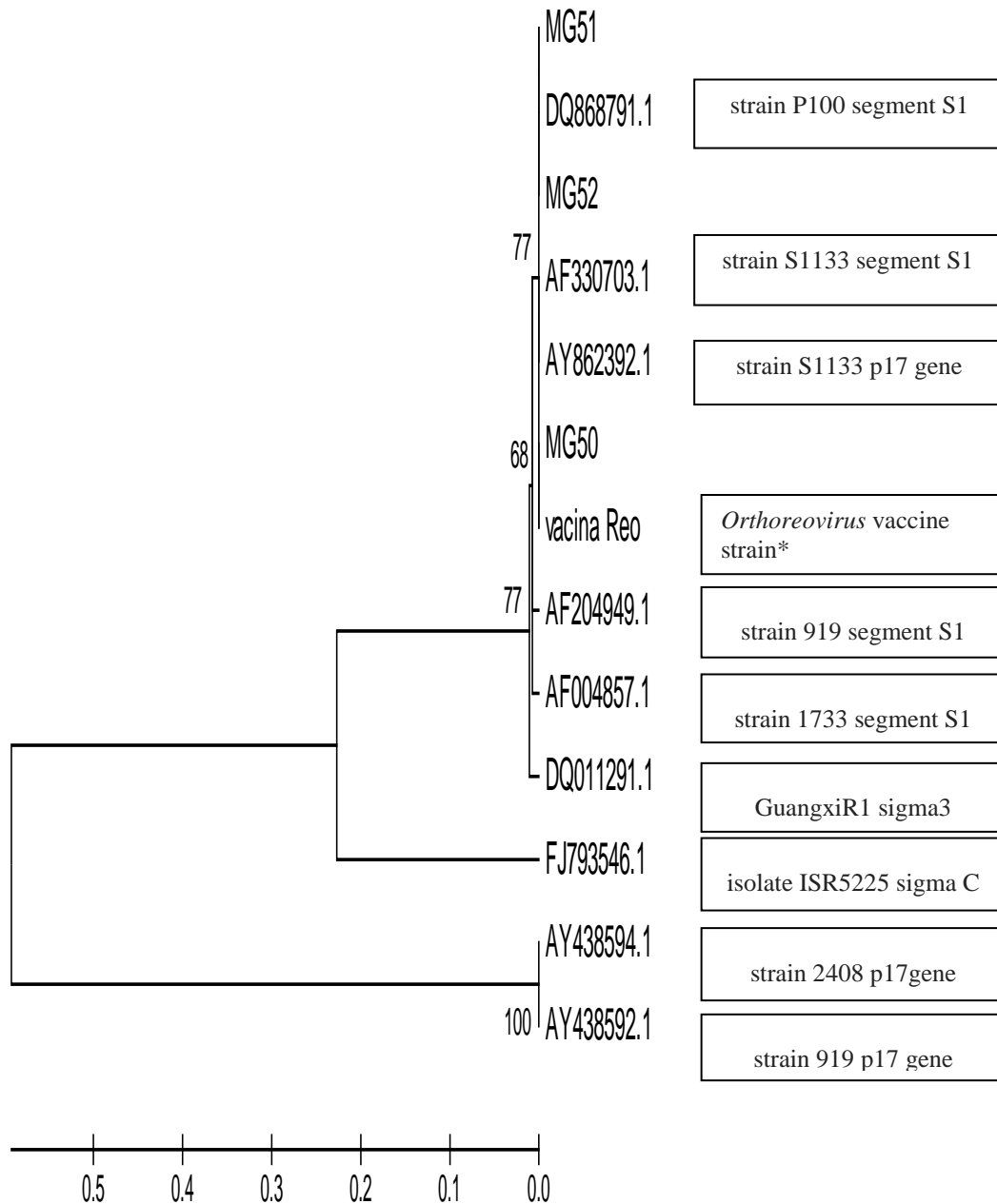


Figure 1. Phylogenetic tree of *Orthoreovirus* based on sequences of cell attachment protein σ 1 encoding gene. Isolates MG50, MG51 and MG52 were grouped with the S1133 strains of Europe, United States and strains employed as attenuated vaccines in Brazil. The scale bar indicates nucleotide substitutions per site. GenBank accession numbers are employed for each reference sequence.

* PoulvacMaternavac IBD-Reo vaccine (Fort Dodge).

AvRV was only detected in broilers, in agreement with previous studies (McNulty *et al.*, 1983, Alfieri *et al.*, 1989). Rotaviruses have been shown to be the most important etiologies associated to gastro-enteric pathology, including results in Brazil (Alfieri *et al.*, 1988 and 1989; Alfieri *et al.*, 1992; Tamehiro *et al.*, 2003; Villareal *et al.*, 2006). The occurrence in two broiler flocks of the same region could be associated to the concentration of flocks and lack of biosecurity. The preclinical incubation infection period might be the reason for its presence in a normal flock, although replicating the virus would last for less than 5 days (McNulty, 2003). Positive samples were found in broilers from 16 to 31 days of age, results which are in agreement with the literature, which reports the occurrence after the second week of age, possibly associated to the decline of passive immunity and increased ambient titers for challenge (McNulty *et al.* 1983).

Previous AvRV studies in Brazil have found an 8.5% occurrence (Tamehiro *et al.*, (2003), results which are similar to those here described (9.25%). The presence of AvRV in mostly clinically affected flocks might suggest an association of its detection with infection, clinical signs and retarded growth (Barnes, 1977; Villareal, 2006). A higher prevalence of 13.81% of avian rotavirus infection was found in chickens in Bangladesh (Karim *et al.*, 2007). However, rotavirus-like electrophoretic mobility, distinct of groups A, D and F, was detected (0 to 2.43%) in broilers in Bangladesh (Ahmed and Ahmed, 2006). Our investigation did not allow such comparison.

Other etiologies might have a significant role for the enteric impairment. It was interesting to know that, for one diarrheic flock, no ARV or AvRV were detected, suggesting other causes of diarrhea, the most commonly being errors in ration formulation, coccidiosis, mycotoxins, other toxins and possibly other viral etiologies, such as other enteric virus (Alfieri *et al.*, 1988; McNulty, 2003; Reynolds and Schultz, 2003; Rosenberger, 2003; Tamehiro *et al.*, 2003).

CONCLUSIONS

After 23 years of the initial description in Minas Gerais, this is an updating study, regarding the ARV and AvRV epidemiology in chickens in the

state. ARV and AvRV have been detected in broilers of the poultry industry in three different intensive poultry *mesoregions*. However, no ARV or AvRV detection was achieved for pullets. Other etiologies for diarrhea might be present, as for one clinically affected flock, no ARV or AvRV were detected. The co-infection of ARV and CAV was detected in a flock severely affected by stunting and arthritis, suggesting synergism. Results reinforce the need for strict biosecurity of poultry in the state in order to reduce losses derived from these and other infections.

ACKNOWLEDGEMENTS

This work is part of the National Institute of Science and Technology (Instituto Nacional de Ciência e Tecnologia-INCT), Genetic and Health Information of the Brazilian Livestock (Informação Genético-Sanitária da Pecuária Brasileira). Authors are indebted to CNPq, FAPEMIG and FEP-MVZ for financial support.

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