

عنوان مقاله:

ژنوتاُیپینگ و تقسیم بندی ویروس های برونشیت عفونی پرندگان بر اساس ژن کد کننده پروتئین S2و تقسیم بندی ویروس های برونشیت عفونی پرندگان بر اساس ژن کد کننده پروتئین S2

محل انتشار:

فصلنامه طب دامي ايران, دوره 10, شماره 4 (سال: 1395)

تعداد صفحات اصل مقاله: 8

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خلاصه مقاله:

Background: Avian infectious bronchitis (IB), with avianinfectious bronchitis virus (IBV) as the causing agent, is a ubiquitous endemicdisease of the chicken with devastating effects on its industry. A viralmembrane surface protein called S notonly induces neutralizing antibodies but also plays an important role in virusbinding and entry to host cells. Technically, S1 protein gene sequencing alsohelps greatly in IBV genotyping. OBJECTIVES: The aim of this study wasto characterize Iranian IBV based on S2 gene. METHODS: After RT-PCRamplification, the S2 gene of nine Iranian IBV isolates were sequenced and thencompared with reference strains. RESULTS: The isolates were classifiedinto genotype I as Massachusetts like IB Vs, genotype VII which clustered intotwo branches, VIIa (IS-1494 like IB viruses), and VIIb, and was related to QX-like viruses and Genotype VIII as 793/B like IBVs. CONCLUSIONS: As faras we know, this is the first S2-based classification study on Iranian IBVisolates providing a firm experimental basis to correlate with genotypiccharacterization

كلمات كليدى:

Characterization, infectious bronchitis virus, genotyping, phylogenetic study, spike

لینک ثابت مقاله در پایگاه سیویلیکا:

https://civilica.com/doc/886828

