INTRODUCTION

Borna disease virus (BDV) is an enveloped, negative-stranded RNA virus. It mainly infects the limbic system, hippocampus, and cerebellum. Animals infected with BDV show various symptoms such as aggression, ataxia, hyperexcitability, and stereotypic behaviour reminiscent of psychiatric symptoms in humans. Due to these similarities, BDV has been considered to be a possible zoonotic agent that may contribute to psychiatric disease.1

Although various warm-blooded animals could be infected with BDV, horses have been known to be the main natural hosts.2 BDV infection causes delayed immunopathology and most animals infected with BDV are asymptomatic.3 Inoue et al. has revealed that 18 out of 35 (51.4%) horses infected with BDV could be seropositive for BDV without any clinical symptoms for 4 consecutive years.4

Initially, BDV infection has been known to be endemic mainly in central Europe such as Germany and Austria.1 However, it has now been reported that horses infected with BDV could be found in Asia.5-11 The prevalence of BDV antibodies or RNA in asymptomatic horses in Asia is as follows: 16.7-29.8% in Japan,4,12,13 20% in China,7 and 23.5-41.2% in Iran.6 There have been also evidences that people living near animals could be infected with BDV.12,14

Despite the growing evidences of the BDV infection in Asia, there has been no report on the BDV prevalence in animals and people in Korea. In a previous study, we examined BDV RNA in race horses and jockeys using one-tube reverse transcriptase polymerase chain reaction (RT-PCR) and nested RT-PCR, but failed to detect BDV RNA in horses and jockeys.15 Thus in this study, we aimed to investigate BDV markers by other methods, real time RT-PCR (rRT-PCR). Considering some evidences indicating discrepancies between serologic study and RT-PCR result,16,17 we used both indirect immunofluorescence antibody (IFA) test and rRT-PCR to compare the results of two methods. We also tried to examine the correlation between the prevalence of BDV in the horses and jockeys.
METHODS

Subjects
Thirty-nine race horses (nine were thoroughbred, 30 were cross-bred) were included in the study. Twenty-three horses were female, and 16 male. The mean age of the horses was 3.1 years (SD=3.3, range: 1-15 years). Clinical and neurological examinations of the horses were performed by a veterinarian. All horses were in a healthy condition. The study protocol was approved by the Ethics Committee of Korea University.

Forty-eight jockeys were also included. Their mean age was 31.2 years (SD=5.5, range: 22-40 years), and their mean duration of contact with horses was 11.2 years (SD=5.8, range: 1-22 years). None had any psychiatric or medical illness, as assessed by means of clinical interviews and physical examinations. Three jockeys had a current familial history of psychiatric illness (2 depression, 1 alcoholism). All the subjects in this study were included in our previous study.15 Informed consent was obtained from the jockeys and the Seoul Horse Race Association.

Preparation of peripheral blood mononuclear cells
A sample of fasting blood (20 mL) was drawn from each subject. Preparation of peripheral blood mononuclear cells (PBMCs) were isolated from anticoagulant (lithium heparin)-treated blood by Ficoll-hypaque gradient centrifugation. Total RNA was extracted from PBMCs using RNAzol (Gibco/BRL, Gaithersburg, MD).

Serological detection of BDV infection with indirect immunofluorescent antibody test
Indirect immunofluorescent antibody (IFA) test was conducted according to the method previously described.16 The Madine Darby canine kidney (MDCK) cell line and MDCK cell line persistently infected with BDV (MDCK-BDV)16 cultured for 5 days with Dulbecco’s Modified Eagle Medium (DMEM) (Gibco/BRL, Germany) supplemented with 10% fetal bovine serum (FBS) (Gibco/BRL, Germany), were trypsinized, suspended in phosphate buffered saline (PBS), spotted onto Teflon-coated 10-well slides, and air-dried in room temperature. Cells on the 10-well antigen slide were fixed with anhydrous acetone at -20°C for 7 minutes and dried. Human and horse serum samples diluted in PBS as 1 : 32 were treated to MDCK cell and MDCK-BDV cell on antigen slide and the slide was incubated in humidity chamber at 37°C for 30 minutes. After three washes with PBS, Fluorescein isothiocyanate conjugated anti-human immunoglobulin goat IgG (MP Biomedicals Inc., USA) was added to each well and the slide was incubated in humidity chamber at 37°C for 30 minutes. The slides were mounted with glycine-buffered glycerol under cover slips and examined for characteristic cytoplasmic fluorescent pattern with a fluorescence microscope.

Genetic detection of viral genomic RNA with real-time reverse transcription-polymerase chain reaction
Real-time quantitative RT-PCR for nucleoprotein (p40) and phosphoprotein (p24) genes of BDV was performed to detect RNA genome of BDV in PBMCs. The primers and probes described in previously study were used for rRT-PCR.20 For the construction of the quantitative standard, p40 and p24 genes were amplified by RT-PCR and cloned to pGEM-T Easy vector system (Promega, USA). Cloned p40 and p24 genes were transcribed to RNA using the MEGAscript T7 transcription kit (Ambion Inc., USA) in accordance with the manufacturer’s instructions. The amounts of p40 and p24 RNA transcribed using by in vitro transcription system were measured by optical density. p40 and p24 RNA were used for the reverse transcription reaction template from 1×1,010 copies/μL to 1×103 copies/μL diluted by 10-fold dilution method. After the reverse transcription reaction, the Taqman Universal PCR Master Mix (applied Biosystems, USA) was used for the real-time PCR reaction as recommended by manufacturer’s instruction. The real-time PCR reaction mixtures were incubated for 10 minutes at 95°C; 40 cycles of amplification were performed using ABI PRISM 7000 Sequence Detection System (Applied Biosystems, USA). Each cycle consisted of a denaturation step (15 seconds at 95°C) and an annealing-elongation step (1 minute at 60°C). Threshold cycle (Ct) values, i.e., the number of cycles for fluorescence to reach to clearly detectable levels, over 40 were regarded as negative.

Statistical analysis
To analyze demographic data, two-tailed t-test was used for continuous covariates. For discrete covariates, chi-square test was used. The null hypothesis was rejected at p<0.05. The statistical package used for the analysis was Statistical Package for the Social Sciences 11.01.

RESULTS

The p24 and p40 RNA were not detected by rRT-PCR in the horses and jockeys. Indirect IFA was conducted for the serological detection of BDV antibodies. No BDV antibody was detected by IFA in the horses and jockeys.

DISCUSSION

We failed to demonstrate BDV RNA and antibody from PBMCs in race horses and jockeys. This result is contrary to several previous studies demonstrating BDV infection in healthy horses in Asia. In Japan, Nakamura et al.13 found that 17 out
of 57 (29.8%) healthy horses were RNA positive and Takahashi et al.\textsuperscript{12} reported that 7 out of 54 (12.9%) healthy thoroughbred race horses and 4 out of 57 (7.0%) blood donors living near those horses were RNA positive, while in Iran, Bahmani et al.\textsuperscript{6} reported an RNA positive rate of 23.6% in healthy race horses (17 out of 72).

Two main reasons might account for the difference between this study and previous Asian reports of BDV occurrence. The first possibility is the difference in diagnostic tools. All of the above mentioned studies detected BDV RNA by nested RT-PCR, which is highly sensitive to contamination, indeed Dürrwald et al.\textsuperscript{21} suggested that most of the BDV positive reported using nested RT-PCR might actually be the result of inadvertent sample contamination. In this study we used rRT-PCR to detect BDV RNA, which has no carry-over contamination risk.\textsuperscript{22} The second possibility is that, in contrast to other areas in Asia, Korea may not be a BDV-endemic area. This agrees with our previous studies which have failed to detect BDV antibodies and RNA in psychiatric patients in Korea\textsuperscript{23} and Japan.\textsuperscript{24} We also used lithium heparin-treated blood for preparation of total RNA from PBMCs. However, ethylenediaminetetraacetic acid is better than heparin as anticoagulant because heparin inhibits the reaction of RT-PCR.

Limitations to this study include its small sample size and restricted region. To accurately determine the BDV prevalence in Korea, large-scaled study should be performed in several different regions.

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