

Detection of bovine coronaviruses from adult cows with epizootic diarrhea and their antigenic and biological diversities

Brief Report

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Summary. Bovine coronavirus (BCV) was detected by reverse transcriptase-PCR, immune electron microscopy or virus isolation from adult cows at 6 out of 6 outbreaks of epizootic diarrhea in Japan. Six BCVs isolated in feces, intestinal content or tracheal exudate of the cows were analyzed for their antigenic properties by cross virus neutralization (VN) tests. The isolates were divided into two groups, one of which had closely related antigenicity with the reference Mebus and Kakegawa strains of BCV, and another which showed significant differences in VN antibody titers from the reference strains. Two isolates in the latter group, which were from the enteric and respiratory tracts of the same cows, respectively, were distinguished from each other by ELISA using monoclonal antibodies against the Kakegawa strain. The isolates showed various hemagglutination and receptor destroying enzyme titers against chicken or mouse erythrocytes.

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Bovine coronavirus (BCV) belongs to antigenic group 2 of the *Coronaviridae*, and possesses four major structural proteins, the nucleocapsid protein N, the integral membrane glycoprotein (gp) M, the hemagglutinin-esterase gp HE and the spike gp S [4]. It is known that both HE and S gps are involved in hemagglutination (HA) activity. The HE gp is a less potent hemagglutinin than the S gp, as it only agglutinates erythrocytes from mice and rats and not those from adult chickens. The HE gp also exhibits receptor destroying enzyme (RDE) activity [4].

BCV is known as a major causative agent of calf diarrhea [3, 4, 10, 15, 24, 25] and winter dysentery of adult cattle [2, 4, 8, 18, 28, 30]. It still remains difficult to isolate the virus on cell cultures. Thus, there have been few papers on the isolation

and antigenicity of BCV, despite numerous reports on their molecular nature. The purpose of this study was to detect BCV from adult dairy cows at 6 outbreaks of epizootic diarrhea in Japan and to compare the antigenic and biological properties of these BCVs with those of the American and Japanese reference strains of BCV.

Diarrheal feces were obtained from 32 adult dairy cows (2 to 9 years of age) in 6 farms at 6 outbreaks of epizootic diarrhea in Okayama Prefecture, Japan, during 1995 and 1997. Small and large intestinal contents, and tracheal exudate were also obtained from a dead adult cow with bloody diarrhea at one of the outbreaks. In each outbreak, diarrhea spread to most of the adult cows at each farm within 1 week. No respiratory symptoms were observed in any cow. A total of 35 field samples were diluted 1:10 in Eagle's minimal essential medium (EMEM), clarified by low-speed centrifugation ($2,500 \times g$ for 10 min), and used for reverse transcriptase PCR (RT-PCR) (30 samples at 5 outbreaks), immune electron microscopy (IEM) (31 samples at 5 outbreaks), and virus isolation (VI) (20 samples at 6 outbreaks). These feces and intestinal contents were also tested for group A rotavirus by latex agglutination tests (Rotascreen; Microgen Bioproducts Ltd., UK), and for *Salmonella* spp., *Campylobacter* spp., *Coccidium* spp. and *Cryptosporidium* spp. by using standard techniques. Serum samples were also collected from 21 cows at the 3 farms at the acute and convalescent phases of the disease. These paired sera were tested for antibodies to BCV and bovine viral diarrhea virus by virus neutralization (VN) tests.

The Mebus [13] and Kakegawa [1] strains of BCV were used as the reference strains. A hyperimmune sera against these reference strains and 5 isolates of BCV were prepared with guinea pigs according to the method described by Tsunemitsu et al. [30].

The HRT-18 cells derived from human rectal adenocarcinoma [29] were used for VI, indirect immunofluorescence (IF), VN tests. For VI, 0.1 ml of field samples were inoculated onto the HRT-18 cells grown in roller tubes (10×100 mm) that had been washed three times with EMEM. After adsorption for 90 min at 37°C , 0.5 ml of EMEM containing $0.5 \mu\text{g/ml}$ of trypsin was added in each tube. The inoculated cells were incubated in a roller drum for 7 days at 37°C , and observed for cytopathic effects (CPE). After three sequential passages, 6 isolates were cloned 3 times by limiting dilution method before use. Confirmation of the isolates as BCV was done by using indirect IF tests [2] and IEM with antiserum to the Kakegawa or Mebus strain of BCV. VN tests were carried out using 96-well micro plates as described previously [31].

Primer pairs for RT-PCR were prepared on the basis of the nucleotide sequence of BCV Mebus strain [11]. The sequences of primers were as follows: forward primer, 5'-ACCACCAGTTCTTGATGTGG-3' (45–26 nucleotides upstream from the M initiation site); reverse primer, 5'-GGCCTAACATACCTTCC-3' (391–409 nucleotides downstream from the M initiation site). The predicted PCR product was 455 base pairs. RNA was extracted by using a Sepa-Gene RNA extraction kit (Sanko Jyunyaku, Japan) according to the manufacturer's instruction. RT-PCR was performed using the GeneAmp RNA PCR core kit (Perkin-Elmer, USA). One microliter of RNA was mixed with 100 pmol of the reverse primer in

PCR buffer containing $MgCl_2$, four deoxynucleotides and reverse transcriptase in a total volume of 20 μ l and incubated at 42 °C for 15 min, 99 °C for 5 min, then 5 °C for 5 min. PCR reaction with 100 pmol of the forward primer was carried out in a final volume of 100 μ l for 35 cycles at 95 °C for 15 sec and 60 °C for 30 sec. PCR products were resolved by electrophoresis in a 2% agarose gel and visualized by ethidium bromide staining.

The field samples and isolates were processed and examined by IEM as described previously [18]. These samples were reacted with guinea pig anti-BCV (Mebus or Kakegawa) serum, negatively stained, and examined by using a JEM-1010 electron microscope (Jeol Ltd., Japan).

An ELISA using monoclonal antibodies (Mabs) for antigenic analysis of BCV was conducted as described previously [21]. Four Mabs against the Kakegawa strain of BCV (4H4 and 7F5 to the S gp, and 2C8 and 1D11 to the N protein [21]) were kindly supplied by Dr. M. Sato, Department of Agriculture, Tochigi Prefectural Office, Tochigi, Japan.

HA tests were carried out with 1% and 0.5% suspensions of mouse and chicken erythrocytes, respectively, at 4 °C and 37 °C according to the microplate method described by Sato et al. [22]. The BCV strains were partially purified from infected cell culture supernatants by ultracentrifugation through a 30% sucrose cushion, which were concentrated approximately 50-fold. The plates incubated at 4 °C were moved to 37 °C for 2 h to measure inactivation of receptors reflected by the disaggregation of the BCV-erythrocyte complexes mediated by the RDE activity [27].

PCR products of 455 bp were detected in 16 of 30 field samples (5 of 5 outbreaks of epizootic diarrhea) (Table 1). The small intestinal content produced no PCR product, while the large intestinal content and tracheal exudate obtained from the same cow were positive in the PCR. As shown in Fig. 1, PCR products were detected in 4 of 5 fecal samples of adult cows at the outbreak in August (lanes 1, 3–5). Characteristic coronavirus-like particles were detected in 19 of 31 samples (5 of 5 outbreaks of epizootic diarrhea) by IEM (Table 1). The particles were approximately 70 to 130 nm in diameter and had short and large projections (data not shown). The 4 PCR-positive samples collected in August were also positive in IEM. The agreement of the results between these tests was 93%. The paired sera from 20 of 21 cows (3 of 3 outbreaks of epizootic diarrhea) showed seroconversion (4-fold increase) to BCV, but no seroconversion to bovine viral diarrhea virus was observed. All fecal and intestinal samples were negative for group A rotavirus, *Salmonella* spp., *Campylobacter* spp., *Coccidium* spp. and *Cryptosporidium* spp.

Six BCV strains designated the 95C6, 95C7, 96C1, 96CT2, 96CE3 and 96C4 strains were isolated in 4 diarrheal feces (95C6, 95C7, 96C1 and 96C4), 1 large intestinal content (96CE3), and 1 tracheal exudate (96CT2) from 5 adult cows at 4 outbreaks of epizootic diarrhea (Table 1). CPE characterized by syncytium formation on HRT-18 cells was observed after two to three serial passages. Cross VN tests were performed with antisera against the 5 isolates and 2 reference strains. The antisera against the reference strains showed 8 to 32-fold lower VN

Table 1. Detection of BCV by RT-PCR, immune electron microscopy (IEM) and virus isolation (VI) in diarrheal feces, intestinal contents and a tracheal exudate of adult cows at 6 outbreaks of epizootic diarrhea

| Outbreak | No. sample | RT-PCR | IEM | VN | Seroconversion ^a |
|----------------|-----------------|------------------|-------|------|-----------------------------|
| A ^b | 6 | 4/5 ^c | 4/6 | 0/4 | NT ^d |
| B | 6 | 2/5 | 4/6 | 1/3 | NT |
| C | 13 ^e | 7/13 | 8/12 | 2/7 | + (5/5) ^f |
| D | 1 | 1/1 | 1/1 | 1/1 | NT |
| E | 6 | 2/6 | 2/6 | 0/2 | + (6/7) |
| F | 3 | NT | NT | 2/3 | + (9/9) |
| Total | 35 | 16/30 | 19/31 | 6/20 | (20/21) |

^aDefined as 4-fold or more increase of virus neutralization antibody titers to BCV (Mebus) in convalescent phase sera compared with those in acute phase sera

^bObserved in the summer of 1996

^cNo. positive /No. tested

^dNot tested

^eIncludes samples of large and small intestinal contents and a tracheal exudate from a dead adult cow

^fNo. of seroconverted cows/No. of tested cows

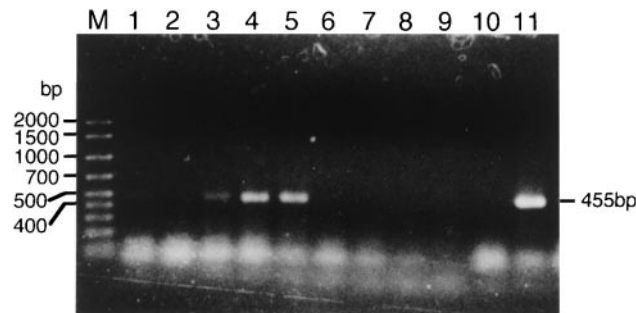


Fig. 1. Agarose gel electrophoresis of the RT-PCR products amplified from fecal specimens obtained from adult cows with epizootic diarrhea in August. *M* Molecular weight marker (Ampli Size DNA size standard; molecular sizes of the marker are indicated on the left in base pair); *1–5* diarrheal feces; *6–9* normal control feces of cows; *10* uninfected cell supernatant; *11* BCV Kakegawa strain infected cell supernatant. A PCR product of approximately 455 bp was detected in *1, 3–5* and *11*

antibody titers to the 96C1, 96CT2, 96CE3 and 96C4 strains than to the homologous strains. The antisera against these 4 isolates also showed 4 to 16-fold lower VN antibody titers to the reference strains than to the homologous isolates (Table 2). According to the results, the isolates were divided into two groups. The first group (95C6 and 95C7) showed closely related antigenicity with the reference strains of BCV. The rest of the isolates showed significant antigenic differences from the reference strains of BCV.

Although all BCVs reacted in ELISA with 2 Mabs recognizing the N protein (1D11 and 2C8), various reactivities of the isolates with 2 Mabs recognizing the S

Table 2. Virus neutralization (VN) tests using antisera against the reference strains (Mebus and Kakegawa) and the isolates of BCV

| BCV strain ^b | VN antibody titers ^a of hyperimmune guinea pig sera to | | | | | | |
|-------------------------|---|-------------|-------------|--------------|-------------|-------------|-------------|
| | Mebus | Kakegawa | 95C6 | 96C1 | 96CT2 | 96CE3 | 96C4 |
| Mebus | 2048 | 2048 | 8192 | <u>1024</u> | <u>512</u> | <u>2048</u> | <u>512</u> |
| Kakegawa | 1024 | 1024 | 4096 | <u>1024</u> | <u>512</u> | <u>1024</u> | <u>512</u> |
| 95C6 | 1024 | 1024 | 4096 | <u>8192</u> | 1024 | <u>2048</u> | 2048 |
| 95C7 | 1024 | 1024 | 4096 | <u>4096</u> | <u>512</u> | <u>2048</u> | <u>1024</u> |
| 96C1 | <u>256</u> | <u>128</u> | 2048 | 16384 | 2048 | 8192 | 4096 |
| 96CT2 | <u>64</u> | <u>32</u> | 4096 | 16384 | 2048 | 8192 | 4096 |
| 96CE3 | <u>128</u> | <u>64</u> | 2048 | 16384 | 4096 | 8192 | 4096 |
| 96C4 | <u>256</u> | <u>64</u> | 2048 | 32768 | 2048 | 8192 | 4096 |

^aExpressed as the reciprocal of the highest dilution of serum inhibiting cytopathic effects. Homologous titers are in bold. Titers which differed by 4-fold or greater with homologous titers are underlined

^bThe 96CT2 and 96CE3 strains were isolated from the tracheal exudate and large intestinal content of a dead adult cow, respectively

Table 3. Reactivity of the BCV strains to monoclonal antibodies (Mabs) against the Kakegawa strain by enzyme-linked immunosorbent assay (ELISA)

| BCV strain ^a | ELISA titers of Mabs to | | | |
|-------------------------|-------------------------|------------------|-------------------|------------------|
| | 4H4 ^b | 7F5 ^b | 1D11 ^c | 2C8 ^c |
| Mebus | 6400 | 25600 | 400 | 800 |
| Kakegawa | 25600 | 12800 | 1600 | 3200 |
| 95C6 | <25 | 6400 | 1600 | 6400 |
| 95C7 | <25 | 6400 | 800 | 6400 |
| 96C1 | 400 | 12800 | 200 | 1600 |
| 96CT2 | <25 | <25 | 1600 | 6400 |
| 96CE3 | 12800 | 6400 | 1600 | 6400 |
| 96C4 | <25 | <25 | 1600 | 3200 |

^aThe 96CT2 and 96CE3 strains were isolated from the tracheal exudate and large intestinal content of a dead adult cow, respectively

^bMabs to the spike glycoprotein of the Kakegawa strain

^cMabs to the nucleocapsid protein of the Kakegawa strain

gp (4H4 and 7F5) were observed. The 96C1 and 96CE3 strains reacted with both anti-S gp Mabs, while the 95C6 and 95C7 strains lacked reactivity to the Mab 4H4, and the 96CT2 and 96C4 strains lacked reactivity to both anti-S gp Mabs. These results clearly show that the isolates were divided into three groups (Table 3). Interestingly, the 95C6 and 95C7 strains were distinguished from the reference strains by the reactivities of the Mab 4H4, whereas these two isolates revealed similar antigenicity to the reference strains by VN tests using polyclonal antisera. Similarly, although the 96CE3 strain was differentiated from the reference strains by the VN tests, the reactivities of this isolate to the Mabs were similar to those of the reference strains. Surprisingly, the 96CE3 and 96CT2 strains isolated in the large intestinal content and tracheal exudate from the same cow, respectively, had different reactivity patterns against anti-S gp Mabs.

The HA and RDE titers of semipurified BCV strains are summarized in Table 4. All strains agglutinated mouse erythrocytes at 4 °C with similar HA titers, and no distinct differences except for the 95C6 and 95C7 strains were observed in HA titers against mouse erythrocytes at 4 °C and 37 °C. All strains also agglutinated chicken erythrocytes at 4 °C, but the HA titers varied among the strains. At 37 °C, most of the strains did not agglutinate chicken erythrocytes. RDE activity with chicken erythrocytes was detected in all strains of BCV. RDE activity with mouse erythrocytes was detected only in the 95C6 and 95C7 strains. The differences in HA titers at 4 °C and 37 °C showed good agreement with the RDE titers. This suggests that comparison of HA titers at 4 °C and 37 °C might provide an alternative method for evaluating RDE activity.

Table 4. Hemagglutination (HA) and receptor destroying enzyme (RDE) activities of semipurified BCV strains

| BCV strain ^b | Infectivity titer ^c | HA titer | | | | RDE titer ^a | |
|-------------------------|--------------------------------|----------|---------|-------|---------|------------------------|---------|
| | | 4 °C | | 37 °C | | mouse | chicken |
| | | mouse | chicken | mouse | chicken | | |
| Mebus | 8.00 | 16384 | >32768 | 16384 | 8192 | <2 | >4 |
| Takegawa | 8.00 | 4096 | 8192 | 8192 | 4096 | <2 | 2 |
| 95C6 | 8.00 | 16384 | 8 | 16 | <2 | 2048 | >4 |
| 95C7 | 7.75 | 65536 | 8 | 32 | <2 | 2048 | >4 |
| 96C1 | 7.75 | 512 | 8 | 2048 | <2 | <2 | >4 |
| 96CT2 | 8.25 | 4096 | 8192 | 8192 | <2 | <2 | >4096 |
| 96CE3 | 8.25 | 16384 | 128 | 32768 | <2 | <2 | >64 |
| 96C4 | 8.25 | 8192 | 8192 | 8192 | <2 | 2 | >4096 |

^aExpressed as the reciprocal of the highest dilution of virus causing complete disappearance of HA patterns at 4 °C after 2 h incubation at 37 °C

^bThe 96CT2 and 96CE3 strains were isolated from the tracheal exudate and large intestinal content of a dead adult cow, respectively

^clog₁₀TCID₅₀/ml

Bovine coronavirus is known as a primary pathogen in neonatal calf diarrhea. The virus is also associated with winter dysentery of adult cattle. In Japan, BCV has been reported as the etiological agent in adult cattle with epizootic diarrhea [28] that occurred throughout the entire country. In the present study, we tried to detect BCV in feces, intestinal contents or tracheal exudate from 33 adult cows at 6 outbreaks of epizootic diarrhea by the RT-PCR, IEM and VI tests. The rate of positive reactions was very high in the RT-PCR (16 of 30 samples at 5 of 5 outbreaks) and IEM (19 of 31 samples at 5 of 5 outbreaks), and the results of these 2 tests showed good agreement. No association with other pathogens was observed in these samples. These results suggest that BCV might be associated with epizootic diarrhea of adult cattle at these outbreaks. Also, the RT-PCR developed in this study may be useful for detecting BCV. Interestingly, BCV was detected by RT-PCR and IEM in 4 of 5 fecal samples from adult cows at the outbreak of epizootic diarrhea in August. Several papers have reported that cold stress, e.g. chilling due to low environmental temperatures including cold drinking water or wide fluctuations in temperatures, is a important risk factor to initiate winter dysentery [17, 32]. The present study indicated that cold stress might not always be needed to initiate adult cow diarrhea associated with BCV.

The previous reports described that BCV also has a tissue tropism to the upper respiratory tract in calves and young cattle [7, 12, 16, 20, 26, 31] and causes respiratory diseases [4, 26]. However, BCV has not been detected in the upper respiratory tract in adult cattle although the possibility of respiratory infection of BCV in adult cattle was pointed out [26]. In this study, BCV (96CT2) was isolated from a tracheal exudate as well as a large intestinal content (96CE3) of a dead adult cow with bloody diarrhea although this cattle did not show clinical signs of respiratory disease. This result suggests that BCV may have a tissue tropism to the upper respiratory tract in adult cows. Further studies are needed to clarify whether BCV can cause respiratory disease in adult cattle because some cows showing winter dysentery develop nasolacrimal discharge and cough [17].

Both the S and HE gps cause HA to bind surface receptors containing Neu5, 9Ac2 [23]. The HE gp is a less potent hemagglutinin than the S gp, therefore it only agglutinates mouse and rat erythrocytes but not chicken erythrocytes which possess fewer surface receptors [23]. Some BCV-infected cell culture supernatants did not agglutinate chicken erythrocytes at 4 °C (data not shown), as similar results were reported previously [27]. Thus, the BCV strains were semipurified from infected cell culture supernatants, which were concentrated approximately 50-fold. All semipurified strains agglutinated chicken erythrocytes at 4 °C, but variations of HA titers were observed among strains. This result suggests that the differences of receptor binding properties of the S gp exist among BCV strains. High RDE activity against mouse erythrocytes was observed in some strains, which was not observed in the previous reports [27, 30]. These high RDE titers with mouse erythrocytes could be explained by the fact that the RDE activity of these strains is higher than other strains because detection of this activity with mouse erythrocytes is harder than with chicken erythrocytes for these abundant surface receptors.

Although BCV has been classified as a single serotype [4, 16], some reports described antigenic variations among BCV strains [2, 5, 6, 9, 14, 30]. Tsunemitsu and Saif [30] reported that some BCV isolates were antigenically distinguished from the Mebus strain in one-way direction by cross VN tests using polyclonal antisera. In the present study, the reference strains Mebus and Kakegawa were closely related, and the isolates were divided into 2 antigenic groups by cross VN tests: the first group (2 strains) was closely related to the antigenicity of the reference strains, and the second group (4 strains) showed significant differences of the antigenicity of the reference strains. These results support the conclusion that BCVs are divided into at least 2 serological subtypes, one is Mebus-like and the others is not [30]. These antigenic variations should be considered for the development of a vaccine against BCV.

The viruses were also divided into several antigenic groups according to the reactivity of the Mabs. The antigenic diversities of BCV with the reactivity of the Mabs and polyclonal antisera exhibited no correlation with each other. The reason for this discrepancy is unknown, but it might be explained that the epitopes recognized by the Mabs were not immunodominant sites, which were not reflected by polyclonal antisera. Interestingly, the 96CE3 and 96CT2 isolated from the enteric and respiratory tracts of the same cow, respectively, belonged to different antigenic groups according to the reactivity of the Mabs. Recently, Storz et al. [26] reported that BCV isolated from respiratory tracts had different antigenic and biological properties from those of BCV isolated from enteric tracts, whereas some investigations could not find any difference between enteric and respiratory BCVs [16, 31]. Our result might indicate that the cow was infected with BCVs having different antigenicity simultaneously. Alternatively, mutation of the virus might occur during infection. Further studies are in progress to compare the nucleotide sequences of the S and HE genes of the isolates to determine the domains which cause antigenic diversity.

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