

Citation: Morioka K, Fukai K, Yoshida K, Kitano R, Yamazoe R, Yamada M, et al. (2015) Development and Evaluation of a Rapid Antigen Detection and Serotyping Lateral Flow Antigen Detection System for Foot-and-Mouth Disease Virus. PLoS ONE 10(8): e0134931. doi:10.1371/journal.pone.0134931

Editor: Xia Jin, University of Rochester, UNITED STATES

Received: March 26, 2015

Accepted: July 16, 2015

Published: August 13, 2015

Copyright: © 2015 Morioka et al. This is an open access article distributed under the terms of the <u>Creative Commons Attribution License</u>, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the paper.

Funding: This study was supported by scientific technique research promotion program for agriculture, forestry, fisheries and food industry, grant No. 25044B (http://www.s.affrc.go.jp/docs/research_fund/2013/sinki_koubo_2013.htm). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

RESEARCH ARTICLE

Development and Evaluation of a Rapid Antigen Detection and Serotyping Lateral Flow Antigen Detection System for Foot-and-Mouth Disease Virus

Kazuki Morioka, Katsuhiko Fukai, Kazuo Yoshida, Rie Kitano, Reiko Yamazoe, Manabu Yamada, Tatsuya Nishi, Toru Kanno*

Exotic Disease Research Station, National Institute of Animal Health, National Agriculture and Food Research Organization, Tokyo, Japan

* kannot@affrc.go.jp

Abstract

We developed a lateral flow strip using monoclonal antibodies (MAbs) which allows for rapid antigen detection and serotyping of foot-and-mouth disease virus (FMDV). This FMDV serotyping strip was able to detect all 7 serotypes and distinguish serotypes O, A, C and Asia1. Its sensitivities ranged from 10³ to 10⁴ of a 50% tissue culture infectious dose of each FMDV stain; this is equal to those of the commercial product Svanodip (Boehringer Ingelheim Svanova, Uppsala, Sweden), which can detect all seven serotypes of FMDV, but does not distinguish them. Our evaluation of the FMDV serotyping strip using a total of 118 clinical samples (vesicular fluids, vesicular epithelial emulsions and oral and/or nasal swabs) showed highly sensitive antigen detection and accuracy in serotyping in accordance with ELISA or RT-PCR. To the best of our knowledge, this is the first report on any FMDV serotyping strip that provides both rapid antigen detection and serotyping of FMDV at the same time on one strip without extra devices. This method will be useful in both FMD-free countries and FMD-infected countries, especially where laboratory diagnosis cannot be carried out.

Introduction

Foot-and-mouth disease (FMD) is one of the most highly contagious viral diseases, and causes devastating economic damage in the countries affected by it. FMD is caused by foot-and-mouth disease virus (FMDV), which belongs to the *Aphthovirus* genus of the family *Picornaviridae*, and consists of 7 immunologically distinct serotypes: O, A, C, Asia1, and South African Territories (SAT) 1, 2 and 3. Within a single serotype, there may be many different topotypes [1]; serotype A in particular displays a large number of topotypes [2]. In Asia, type O, A and Asia1 viruses have prevalently existed and the disease is endemic or sporadic in main countries [3]. Additionally, there have been some cases of FMD whose serotype is unknown in South

East Asia [3]. Although outbreaks of the SAT serotypes have been largely limited to the African continent, it has been reported that the SAT2 serotype invaded Middle Eastern countries in 2000 and 2012 [4].

In an FMD outbreak, antigen serotyping should be conducted immediately in order to implement the appropriate plan for disease control (e.g., preparing vaccine and immunological diagnosis kits for the serotype in question). To serotype FMDV, indirect sandwich enzyme-linked immunosorbent assay (ELISA), which is mentioned in the World Organization for Animal Health (OIE) *Terrestrial Manual 2012* [5], may be used, however it lacks sufficient sensitivity and specificity [6–9]. Although some studies have reported serotyping by reverse transcription-polymerase chain reaction (RT-PCR) or real-time RT-PCR in some regional endemic FMD strains, the mutability of viral RNA makes it difficult to apply either of these as a universal method [10]. Moreover, the recent activation of world trade and human transportation might change the epidemic styles of trans-boundary emerging diseases. The possibility of new outbreaks of the different topotypes in places where they have never previously occurred is increasing. We have developed a monoclonal antibody (MAb)-based direct sandwich ELISA (MSD-ELISA), which demonstrates higher sensitivity than those of current indirect sandwich ELISA methods reported in previous studies [8,9]. ELISA is a useful tool for antigen detection, however this should be carried out in a laboratory with the appropriate apparatus.

In this report, we developed a lateral flow assay using MAbs (FMDV serotyping strip), which allows for rapid FMD antigen detection for all 7 serotypes and FMD serotyping for types O, A, C and Asia1, specifically in the field, and especially in countries where laboratory diagnosis cannot be carried out, in vast countries in which it would take a long time to transport clinical samples to a laboratory, and/or in countries lacking transportation facilities.

Materials and Methods

Cells and viruses

The virus strains FMDV O/JPN/2000 (ME-SA topotype, Pan-Asia lineage) [11–13], O/JPN/ 2010 (SEA topotype, Mya-98 lineage) [14], O1 Manisa (TUR 8/69) (ME-SA topotype, Pan-Asia lineage), O1 BFS 1860 (Euro-SA topotype), O/TAW/97 (Cathay topotype) [15, 16], O/ TUR/5/2009 (ME-SA topotype, Pan-Asia2 lineage), A15 TAI 1/60 (Asia topotype), A22 IRQ 24/64 (Asia topotype), A/IRN/1/2011 (Asia topotype, Iran-05 ^{FAR-11} lineage), A/TAI/10/2011 (Asia topotype, Sea-97 lineage), C PHI 7/84 (Euro-SA topotype), Asia1 Shamir (ISR 3/89) (Asia topotype), Asia1/TUR/49/2011 (Asia topotype, Shindh-08 lineage) [17, 18], SAT1/KEN/ 117/2009 (topotype I (NWZ)), SAT2/SAU/6/2000 (topotype VII), SAT3/ZIM/3/83 (topotype I (SEZ)) and swine vesicular disease virus (SVDV) J1/73 [19, 20] were grown on monolayers of IBRS-2 [21] and/or BHK-21 [22] cells and used for this study.

Monoclonal antibodies

In the present study, MAbs producing hybridomas 13F1 and 2A1 were established against C PHI 7/84 and A/IRN/1/2011, respectively, by the general method as previously described [12]. MAb 1H5 and MAb 70C4 originating from O/JPN/2000, MAb16C6 originating from A15 TAI 1/60, and MAb 12C7 originating from Asia1 Shamir (ISR 3/89) were used in this study [8, 9].

Conjugation of MAb 1H5 with colloidal gold

A mixture of 0.1 ml of purified MAb 1H5, which reacts with all 7 serotypes of FMDV (200 μ g/ml) and 0.8 ml of colloidal gold (pH 8.0) (Winered Chemical Corporation, Tokyo, Japan) was stabilized for 8 min at room temperature. Next, 0.1 ml of 10% bovine serum albumin (BSA) and

0.01 ml of 5% polyethylene glycol (PEG) were added, and the mixture was centrifuged at 10,000 rpm at 20°C for 30 min. After removal of the supernatant, the colloidal gold-labeled MAb 1H5 was resuspended with 1.5 ml of phosphate buffered saline (PBS) containing 0.5% BSA and 0.05% PEG, centrifuged again under the same conditions mentioned above, and resuspended with 20 mM Tris-HCl (pH 7.5) containing 10% trehalose, 10% BSA and 5% PEG. After the density was adjusted to 2.0 of OD₅₂₀, 50 µl of the colloidal gold labeled MAb 1H5 (G-MAb) was freeze-dried in a 2-ml tube.

Production of the FMDV serotyping strip

Anti-FMDV MAbs against each serotype, MAb 70C4 (type O), 16C6 and 2A1 (type A), 13F1 (type C) and 12C7 (type Asia1), multi-serotype reactive MAb 1H5 and AffiniPure goat antimouse IgG (H+L) (Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA) were suspended with 5 mM phosphate buffer (pH 7.5) in 1000–1500 µg/ml, and applied on a nitrocellulose membrane (Hi-Flow Plus HF135; Millipore, Billerica, MA, USA) using immunoliner 200 (System Biotics, Kanagawa, Japan) at 0.8 µl/cm. The antibodies on nitrocellulose membranes were dried at 50°C for 30 min, dipped in 0.05 M boric acid with 0.5% casein (pH 8.5) for 30 min at room temperature, dipped in 50 mM Tris-HCl (pH 7.4) with 0.5% sucrose and 0.05% sodium cholate for 30 min at room temperature, and dried at room temperature overnight.

The cellulose fiber membrane (Cellulose Absorbent Sample Pads; Millipore), which was cut to a width of 1 cm, dipped in 100 mM Tris-HCl (pH 8.0) with 1.6 mM ethylenediaminetetraacetic acid (EDTA) for blocking against non-specific reactions due to contamination of saliva or hemolytic fluid and dried overnight at room temperature, was used as an absorption pad, and stuck with adhesive tape on the downstream side of the nitrocellulose membrane, overlapping by 5 mm (ARcare 7815; Adhesives Research, Glen Rock, PA, USA). A cotton membrane (GE Healthcare Life Sciences, Little Chalfont, UK) 2 cm in width was used as an absorption pad and stuck with adhesive tape on the upstream side of the nitrocellulose membrane, overlapping by 5 mm. Constructed membranes were then cut at a width of 0.5 cm, and the strips were stored at room temperature under low humidity.

Monoclonal antibody-based sandwich direct ELISA

MSD-ELISA was established and described in our previous reports [8, 9], and the protocol is described in detail [8]. In the present study, OD values over 0.1 were deemed positive.

Indirect sandwich ELISA

Indirect sandwich (IS) ELISA [23], which is a method described in the OIE manual [5], was carried out according to the relevant instruction manuals. For the purposes of the present study, OD values over 0.1 were considered positive.

Statistics

The Pearson's chi-square test was used to analyze the statistical significance of the differences in virus detection rates between IS-ELISA and the FMDV serotyping strip. To analyze correlations between the OD values of MSD-ELISA and the mABS values of the FMDV serotyping strip, Pearson's correlation coefficient was used.

RT-PCR

For the RT-PCR for the detection of FMDV nucleic acid, the SuperScript III One-Step RT-PCR System with Platinum Taq DNA Polymerase (Invitrogen, Waltham, MA, USA) and primers for the 3D region were used [15].

Lateral flow antigen detection commercial kit

A Svanodip FMDV-Ag kit (Boehringer Ingelheim Svanova, Uppsala, Sweden) was used following the manufacturer's instructions. A total of 170 μ l of samples diluted with sample buffer were added to the device and, after 10 min, mABS was measured using the Immunochromato-Reader C10066.

Clinical samples

A total of 16 samples of vesicular fluids collected from pigs inoculated with FMDV O/JPN/ 2000 and A15 TAI 1/60 were used for each antigen detection method. A total of 53 samples of vesicular epithelial tissues collected from pigs and cattle inoculated with FMDV O/JPN/2010 were used for each antigen detection method.

In addition to the experimental samples, a total of 49 RT-PCR-positive samples (41 oral and/or nasal swabs soaked in approximately 10-times volumes of PBS (about 2 ml) and 12 samples of 10% emulsion of homogenized epithelial tissues) collected from cattle and pigs involved in the 2010 type O FMD outbreak in Japan caused by O/JPN/2010 (SEA topotype) [14] were used for comparative studies.

Field samples (vesicular epithelium and vesicular lesion's swab) were submitted from Miyazaki Prefecture, Japan, for diagnosis of the 2010 FMD in Japan.

Ethics statement

The animal experiments carried out in the present study were approved by the Ethics Committee of the National Institute of Animal Health, Japan (approvals #09–029, 14–060, and 14– 080). The field samples used in this study were submitted from Miyazaki Prefecture, Japan, for diagnosis of the 2010 FMD in Japan. These vesicular epithelial tissues were collected by veterinarians in accordance with the guidelines specified in the Act on Domestic Animal Infectious Diseases Control.

The FMD serotyping strip

One hundred μ L of samples diluted with PBS were added to the tube containing the freezedried G-MAb, and the FMDV serotyping strip was then dipped into the tube. After 12 min, the lines were interpreted by visual observation and/or the milli absorbance (mABS) of the lines was measured using an Immunochromato-Reader C10066 (Hamamatsu Photonics, Shizuoka, Japan). With respect to visual interpretation, mABS "from 10 to 15" and "over 15" were deemed equivalent to "weak positive" and "positive," respectively.

Antigen detection for all seven serotypes and serotyping (O, A, C and Asia1). A total of 17 viruses, 16 strains of FMDV and the SVDV J1 strain, were adjusted to almost 10^6 of 50% tissue culture infectious dose (TCID₅₀)/100 µL with PBS, and those of 100 µL were applied for the FMD serotyping strips.

Detection limits of the FMD serotyping strip. Ten-hold serial dilutions were prepared with PBS or running buffer for Svanodip against the 16 FMDV strains as follows: FMDV O/ JPN/2000, O/JPN/2010, O1 BFS 1860, O1 Manisa (TUR 8/69), O/TUR/5/2009, O/TAW/97, A15 TAI 1/60, A22 IRQ 24/64, A/IRN/01/2011, A/TAI/10/2011, C PHI 7/84, Asia1 Shamir

(ISR 3/89), Asia1/TUR/49/2011, SAT1/KEN/117/2009, SAT2/SAU/6/2000 and SAT3/ZIM/3/83, and applied to the FMD serotyping strip (100 μ l) and Svanodip (170 μ l). The total of applied virus antigens of Svanodip were 1.7 times as much as that of the FMD serotyping strip. Only those for type SAT3 were applied from 10⁶ TCID₅₀.

Evaluation of the FMD serotyping strip using clinical samples. The FMD serotyping strips were evaluated using 10 µl of vesicular fluid from pigs inoculated with O/JPN/2000 or A15/TAI/1/60, or almost 10% (1 to 10%) vesicular epithelial emulsions of pigs and cattle inoculated with O/JPN/2010 and of animals infected in the 2010 outbreak in Japan caused by O/JPN/2010, and the results were compared with those obtained using other antigen detection methods, specifically MSD-ELISA, IS-ELISA and RT-PCR. IS-ELISA was not performed for vesicular fluids because of insufficient sample volume. In the present study, mABS values over 15 were judged positive samples, and this almost correlated with visual inspections.

In this evaluation, the positions of the serotype-specific antibodies were rearranged from O, A, Asia1, C (from upstream to downstream) to C, O, A, Asia1 to inspect non-specific reactions due to contamination of hemolytic fluid or other contamination.

Results

Antigen detection for all seven serotypes and serotyping (O, A, C and Asia1)

The FMDV serotyping strips were able to detect all 16 strains of the 7 serotypes of FMDVs including recent epidemic strains, and were able to distinguish serotypes O, A, C and Asia1, but did not react to SVDV. However, when a high titer of SAT2 was applied, a weak non-specific line was observed on the serotype A line (Fig 1).

Sensitivities of the strip

The antigen detection limits of each serotyping line of the FMDV serotyping strip were found to be almost equal to 10^3 to 10^4 TCID₅₀ of each virus stain (Fig 2). For O/JPN/2010 (SEA), the FMDV serotyping strip showed a strong positive line at 10^4 TCID₅₀ and a doubtful line at 10^3



Fig 1. FMDV serotyping on one strip. The result of reactivity of the FMD serotyping strip against seven serotypes of FMDVs and SVDV. One hundred μ L of a total of 17 viruses of almost 10⁶ TCID₅₀ were added to the tube containing the freeze-dried G-MAb, and the FMDV serotyping strip was then dipped into the tube. After 12 min, the lines were interpreted.

doi:10.1371/journal.pone.0134931.g001



Fig 2. The sensitivities of the FMD serotyping strips. The sensitivities of the FMD serotyping strip and Svanodip against each FMDV strain were compared. Only type SAT3 was applied from 10⁶ TCID₅₀. Closed bars show the mABS of each serotype-specific line of the FMD serotyping strip. Shaded bars show the mABS of Svanodip. The dashed lines on the graphs show 15 mABS lines judged positive by visual inspection. The numbers on the right edge of the graph show mABS over 150.

doi:10.1371/journal.pone.0134931.g002

TCID₅₀, while the Svanodip showed a very weak positive at 10^4 TCID₅₀. The detection limits of both the FMDV serotyping strip and Svanodip for O/TAW/97 and O1 BFS1860 were almost 10^3 TCID₅₀. The detection limits for the ME-SA topotype (O/JPN/2000, O1 Manisa, O/TUR/ 5/2009) of the FMDV serotyping strip and Svanodip were 10^4 and 10^5 TCID₅₀, respectively. In the case of serotype A strains, the detection limits of the FMDV serotyping strip and Svanodip were around 10^4 TCID₅₀. The detection limits of the FMDV serotyping strip and Svanodip for Asia1 Shamir (ISR 3/84) and Asia1/TUR/2011 were 10^4 and 10^5 TCID₅₀, respectively. The detection limits of both the FMDV serotyping strip and Svanodip for C/PHI/7/84, SAT1/KEN/ 2009 and SAT2/SAU/2000 were 10^4 TCID₅₀. The FMDV serotyping strip showed high sensitivity to the serotype O ME-SA topotype compared to Svanodip. On the other hand, the FMDV serotyping strip was not sensitive to SAT3, at a detection limit of 10^5 TCID₅₀, while Svanodip were able to detect 10^4 TCID₅₀.

Evaluation of the FMD serotyping strip using clinical samples

As shown in Table 1, the FMDV serotyping strip was able to detect FMD antigens on the target serotype lines against all 16 samples of vesicular fluid (9 samples for A15 TAI 1/60 and 7 samples for O/JPN/2000), 90.2% (55/61) samples of 10% vesicular epithelial emulsions collected from experimental infection of O/JPN/2010, and 68.3% (28/41) of oral and or nasal swabs from the 2010 outbreak in Japan caused by O/JPN/2010. The FMDV serotyping strip was found to be more sensitive than IS-ELISA against vesicular epithelial emulsions and oral and/ or nasal swabs (p<0.01). The correlation coefficient between mABS and the OD of MSD-E-LISA was significantly high (R = 0.67).

PLOS ONE

virus strain	Experimental or field	animals	samples	FMDV Serotyping Strip (O or A)	MSD-ELISA ^a (O or A)	IS-ELISA ^b (O)
A15 TAI 1/60	experimental	pig	VF ^{c,d}	100% (9/9) ^e	100% (9/9) ^e	Not tested
O/JPN/2000	experimental	pig	VF	100% (7/7)	100% (7/7)	Not tested
O/JPN/2010	experimental	pig	VEE ^f	100% (37/37)	100% (37/37)	67.6% (25/37) ^e
O/JPN/2010	field	pig	VEE	100% (4/4)	100% (4/4)	100% (4/4)
O/JPN/2010	field	pig	SW ^g	55.6% (5/9)	100% (9/9)	0.0% (0/9)
		subtotal		93.9% (62/66)	100.0% (66/66)	58.0% (29/50)
O/JPN/2010	experimental	cattle	VEE	68.8% (11/16)	87.5% (14/16)	25.0% (4/16)
O/JPN/2010	field	cattle	VEE	75.0% (3/4)	75.0% (3/4)	0.0% (0/4)
O/JPN/2010	field	cattle	SW	71.9% (23/32)	100% (32/32)	9.4% (3/32)
		subtotal		71.2% (37/52)	94.2% (49/52)	13.5% (7/52)
		total		83.9% (99/118)	97.5% (115/118)	35.3% (36/102)

Table 1. Comparison of the results of antigen detection using clinical samples.

^a MSD-ELISA is MAb-based antigen detection ELISA

^b IS-ELISA is indirect sandwich ELISA supplied by the World Reference Laboratory

^c Vesicular fluid

^d10 µl of vesicular fluid were added to 90 µl of running buffer

^eThe number of positive samples by each antigen detection method/the number of positive samples by RT-PCR

^f Vesicular epithelial emulsion

^g Oral and/or nasal vesicular lesion swabs.

doi:10.1371/journal.pone.0134931.t001

Discussion

In the present study, we developed an FMD serotyping strip that enables rapid antigen detection for all 7 serotypes and serotyping for O, A, C and Asia1, including recent epidemic FMDVs. This FMD serotyping strip allows for rapid antigen detection and 4 types of serotyping of FMDV without extra devices.

Although antigen detection ELISA requires that samples of each serotype are added to detecting wells, this FMD serotyping strip is able to perform simultaneous detection and sero-typing of FMDV on a single strip. Furthermore, our lateral flow system detects not only FMDV but also type-specific FMDV. This not only increases sensitivity but also counteracts the short-comings of MAb, which recognizes a single epitope that depends on broad intra-type reactivity.

In Southeast Asia, there have been many cases in which the causative FMDV serotype was unknown [3]. The present method will be useful not only in FMD-free countries but also in FMD-infected countries, especially where laboratory diagnosis is not possible. Thus, the FMD serotyping strip will prevent delay of serotyping reports to the OIE and we will be able to share more precise information about FMD epidemics.

The advantage of using MAbs is that it is possible to select highly efficient MAbs for high specificity and sensitivity. However, a MAb recognizes a single epitope, and thus the broad intra- and intertype reactivity of the MAbs should be evaluated. To confirm the relativities of FMDV epidemic strains and our MAbs, we introduced the recent epidemic FMDV serotype O, A and Asia1 strains O/TUR/5/2009, A/IRN/01/2011, A/TAI/10/2011, Asia1/TUR/49/2011, and serotypes SAT1, SAT2 and SAT3 as follows: SAT1/KEN/117/2009, SAT2/SAU/6/2000 and SAT3/ZIM/3/83. MAbs 70C4 and 12C7 were able to react to the recent epidemic FMDV strains of serotypes O and Asia1, respectively. MAb 1H5 was able to react to the recent epidemic FMDV strains including serotype SAT1, 2 and 3. Therefore, the multi-serotype

detection line is able to be warranty for SAT1, 2 and 3, and in view of mismatch of each serotype specific MAbs. In our previous study, we used MAb 16C6 as a detecting antibody for MSD-ELISA for type A FMDV, however it did not react to A/IRN/01/2011, although it react to A/TAI/10/2011. The strain A/IRN/01/2011 belongs to the A-Iran-05 lineage, which appeared in 2005 as a new lineage of FMDV serotype A [24, 25]. That strains spread throughout Iran and moved westward into Saudi Arabia and Turkey, and were also reported in 2006 in Pakistan, and in 2007 in Jordan [25]. To solve this problem, we produced MAb 2A1 by A/IRN/01/ 2011 as the antigen, and this MAb was able to react to A/IRN/01/2011 as well as to A22 IRQ 24/64. Therefore, in the present study, we used both MAb 16C6 and 2A1 as cocktails of the antigen detection MAb on the membrane.

We used oral swabs for samples for MSD-ELISA in our previous study [9], however, we found that some oral and/or nasal vesicular lesion swabs from 2010 inhibited chromatography flow in the present study (data not shown). Saliva is expected to be a useful sample for detecting antigens, especially from preclinical infected animals [26] and subclinical infected animals such as those that are vaccinated-infected or have low pathogenic FMDV. For instance, the cattle in the 2000 FMD outbreak in Japan did not show any distinct vesicles on the feet, or in or around the oral and nasal cavities. [11–13]. In addition, even in typical clinical cases such as those in 2010, almost all samples were oral and/or nasal vesicular lesion swabs from the outbreak area because the vesicles readily ruptured and it was difficult to collect epithelial tissues.

On the other hand, using vesicular fluids and/or epithelial emulsions as a sample has the advantage that the sample would include a high titer of virus, and antigens can be detected a long time after the appearance of lesions. The present FMDV serotyping strip was able to detect antigens from 4 to 13 days (the end of the experiment) post-inoculation when the lesions were almost healed (<u>Table 1</u>; sampling date not shown).

In this study, the correlation coefficient between mABS of the FMDV serotyping strip and the OD of MSD-ELISA was significantly high (R = 0.67) and this strip was found to be more sensitive than IS-ELISA (p<0.01). As described in our previous study [9], the difference of sensitivity was probably due to the ability of MAb that we used in this system, much more than those of rabbit and guinea pig anti-sera in IS-ELISA. In addition, concerning specificity, the FMD serotyping strip showed negative results against a total of 21 of RT-PCR negative samples (14 of oral swabs or saliva and 7 of epithelial emulsions) (data not shown).

Further study is ongoing, and we are trying to establish serotype-specific MAbs against SAT1, 2 and 3, and apply them in the lateral flow assay for the detection and serotyping of all 7 serotypes of FMDVs.

Acknowledgments

We are grateful to the Pirbright institute (UK) for the supply of recent epidemic foot-andmouth disease viruses and Dr. Hiroshi Iseki (NIAH, Japan) for helpful technical advice. We would also like to thank Mr. Hiroki Kimura, Mr. Masayuki Kanda, Mr. Shinya Sato, Mr. Kenichi Ishii, Mr. Tatsuo Nakamura, and Mr. Shigeo Mizumura for care of the animals.

Author Contributions

Conceived and designed the experiments: KM KY TK. Performed the experiments: KM RK RY. Analyzed the data: KM RK. Contributed reagents/materials/analysis tools: KM KF MY TN. Wrote the paper: KM KF KY MY TN TK.

References

- Knowles NJ, Samuel AR. Molecular epidemiology of foot-and-mouth disease virus. Virus Res. 2003; 91: 65–80. PMID: <u>12527438</u>
- Kitching RP. Global epidemiology and prospects for control of foot-and-mouth disease. Curr Top Microbiol Immunol. 2005; 288: 133–148. PMID: <u>15648177</u>
- 3. OIE SEAFMD home page, Monthly FMD maps and country reports. [cited 2010–2015]. Available: http://www.arahis.oie.int/reports.php?site = seafmd
- Hall MD, Knowles NJ, Wadsworth J, Rambaut A, Woolhouse MEJ. Reconstructing geographical movements and host species transitions of foot-and-mouth disease virus serotype SAT 2. mBio Sep-Oct. 2013; 4(5): e00591–13. doi: <u>10.1128/mBio.00591-13</u> PMID: <u>24149511</u>
- 5. Office International des Epizooties. Foot and mouth disease. In: OIE manual of diagnostic tests and vaccines for terrestrial animals. Paris, France: OIE; 2012. pp. 190–216.
- Mohapatra JK, Subramaniam S, Tosh C, Hemadri D, Sanyal A, Periyasamy TR, et al. Genotype differentiating RT-PCR and sandwich ELISA: handy tools in epidemiological investigation of foot and mouth disease. J Virol Methods. 2007; 143: 117–121. PMID: <u>17400298</u>
- Reid SM, Forsyth MA, Hutchings GH, Ferris NP. Comparison of reverse transcription polymerase chain reaction, enzyme linked immunosorbent assay and virus isolation for the routine diagnosis of foot-andmouth disease. J Virol Methods. 1998; 70: 213–217. PMID: <u>9562415</u>
- Morioka K, Fukai K, Yoshida K, Yamazoe R, Onozato H, Ohashi S, et al. Foot-and-mouth disease virus antigen detection enzyme-linked immunosorbent assay using multiserotype-reactive monoclonal antibodies. J Clin Microbiol. 2009; 47: 3663–3668. doi: 10.1128/JCM.00695-09 PMID: 19759230
- Morioka K, Fukai K, Sakamoto K, Yoshida K, Kanno T. Evaluation of monoclonal antibody-based sandwich direct ELISA (MSD-ELISA) for antigen detection of foot-and-mouth disease virus using clinical samples. PLoS ONE. 2014; 9(4): e94143. doi: <u>10.1371/journal.pone.0094143</u> PMID: <u>24736560</u>
- Longjam N, Deb R, Sarmah AK, Tayo T, Awachat VB, Saxena VK. A Brief Review on Diagnosis of Foot-and-Mouth Disease of Livestock: Conventional to Molecular Tools. *Vet. Med Int.* 2011; 2011:905768. doi: 10.4061/2011/905768
- 11. Kanno T, Yamakawa M, Yoshida K, Sakamoto K. The complete nucleotide sequence of the PanAsia strain of foot-and-mouth disease virus isolated in Japan. Virus Gene. 2002; 25: 119–125.
- Morioka K, Fukai K, Ohashi S, Sakamoto K, Tsuda T, Yoshida K, et al. Comparison of the characters of the plaque-purified viruses from foot-and-mouth disease virus O/JPN/2000. J Vet Med Sci. 2007; 70: 653–658.
- Sakamoto K, Kanno T, Yamakawa M, Yoshida K, Yamazoe R, Murakami Y. Isolation of foot-and-mouth disease virus from Japanese black cattle in Miyazaki Prefecture, Japan, 2000. J Vet Med Sci. 2002; 64: 91–94. PMID: 11853156
- 14. Fukai K, Morioka K, Yoshida K. An experimental infection in pigs using a foot-and-mouth disease virus isolated from the 2010 epidemic in Japan. J Vet Med Sci. 2011; 73: 1207–1210. PMID: 21532260
- Chen BJ, Sung WH, Shieh HK. Managing an animal health emergency in Taipei China: foot and mouth disease. Rev Sci Tech. 1999; 18: 186–192. PMID: <u>10190214</u>
- Huang CC, Jong MH, Lin SY. Characteristics of foot-and-mouth disease virus in Taiwan. J Vet Med Sci. 2000; 62: 677–679. PMID: <u>10945282</u>
- 17. Jamal SM, Ferrari G, Ahmed S, Normann P, Belsham GJ. Molecular characterization of serotype Asia-1 foot-and-mouth disease viruses in Pakistan and Afghanistan; emergence of a new genetic group and evidence for a novel recombinant virus. Infect, Genet Evol. 2011; 11: 2049–2062.
- Knight-Jones TJ, Bulut AN, Gubbins S, Stärk KD, Pfeiffer DU, Sumption KJ, et al. Retrospective evaluation of foot-and-mouth disease vaccine effectiveness in Turkey. Vaccine. 2014; 32: 1848–1855. doi: <u>10.1016/j.vaccine.2014.01.071</u> PMID: <u>24530150</u>
- Inoue T, Yamaguchi S, Kanno T, Sugita S, Saeki T. The complete nucleotide sequence of a pathogenic swine vesicular disease virus isolated in Japan (J1_73) and phylogenetic analysis. Nucleic Acids Res. 1993; 21: 3896. PMID: <u>8367308</u>
- 20. Tokui T, Kono M, Tokuda G, Kumagai T, Sasahara J. Outbreaks of swine vesicular disease in Japan: virus isolation and epizootiological survey. Natl Inst Anim Health Q. 1975; 15: 165–173.
- 21. Chapman WG, Ramshaw IA. Growth of the IB-RS-2 pig kidney cell line in suspension culture and its susceptibility to foot-and-mouth disease virus. Appl. Environ. Microbiol. 1971: 22:1–5.
- Stoker M, Macpherson I. Syrian Hamster Fibroblast Cell Line BHK21 and its Derivatives, Nature 1964: 203: 1355–1357. PMID: <u>14207308</u>

- Roeder PL, Le Blanc Smith PM. Detection and typing of foot-and-mouth disease virus by enzymelinked immunosorbent assay: a sensitive, rapid and reliable technique for primary diagnosis. Res Vet Sci. 1987; 43: 225–232. PMID: <u>2825310</u>
- Klein J, Hussain M, Ahmad M, Normann P, Afzal M, Alexandersen S. Genetic characterization of the recent foot-and-mouth disease virus subtype A/IRN/2005. Virol J. 2007; 4: 122. PMID: <u>18001482</u>
- 25. Knowles NJ, Nazem Shirazi MH, Wadsworth J, Swabey KG, Stirling JM, Statham RJ, et al. Recent spread of a new strain (A-Iran-05) of foot-and-mouth disease virus type A in the Middle East. Transbound Emerg Dis. 2009; 56:157–69. doi: <u>10.1111/j.1865-1682.2009.01074.x</u> PMID: <u>19432637</u>
- 26. Fukai K, Yamada M, Morioka K, Ohashi S, Yoshida K, Kitano R, et al. Dose-dependent responses of pigs infected with foot-and-mouth disease virus O/JPN/2010 by the intranasal and intraoral routes. Arch Virol. 2015; 160:129–139. doi: 10.1007/s00705-014-2239-4 PMID: 25281431