



Title	Development of an Immunochromatography Assay (QuickNavi-Ebola) to Detect Multiple Species of Ebolaviruses
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1 **Development of an immunochromatography assay (QuickNavi™-Ebola) to detect**
2 **multiple species of ebolaviruses**

3

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37 **Abstract**

38 The latest outbreak of Ebola virus disease (EVD) in West Africa has highlighted the
39 urgent need for the development of rapid and reliable diagnostic assays. We utilized
40 monoclonal antibodies specific to the ebolavirus nucleoprotein to develop an
41 immunochromatography (IC) assay (QuickNavi™-Ebola) for rapid diagnosis of EVD.
42 The IC assay was first evaluated with tissue culture supernatants of infected Vero E6 cells
43 and found to be capable of detecting 10^3 - 10^4 focus-forming units/ml of ebolaviruses.
44 Using serum samples from experimentally infected nonhuman primates, we confirmed
45 that the assay could detect the viral antigen shortly after disease onset. It was also noted
46 that multiple species of ebolaviruses could be detected by the IC assay. Owing to the
47 simplicity of the assay procedure and absence of requirements for special equipment and
48 training, QuickNavi™-Ebola is expected to be a useful tool for rapid diagnosis of EVD.
49

50 **Keywords:** Ebola virus; ebolavirus; EVD; filovirus; nucleoprotein; monoclonal
51 antibody; immunochromatography assay; diagnosis

52 INTRODUCTION

53 Ebolaviruses and marburgviruses are enveloped, nonsegmented, negative-
54 stranded RNA viruses belonging to the family *Filoviridae*. These filoviruses are known
55 to cause severe hemorrhagic fever in humans and nonhuman primates with human case
56 fatality rates of up to 90% [1]. In contrast to the genus *Marburgvirus*, which contains only
57 one known species, *Marburg marburgvirus* consisting of Marburg virus (MARV) and
58 Ravn virus, five distinct species are known in the genus *Ebolavirus*, *Zaire*
59 *ebolavirus*, *Sudan ebolavirus*, *Tai Forest ebolavirus*, *Bundibugyo ebolavirus*, and *Reston*
60 *ebolavirus*, represented by Ebola virus (EBOV), Sudan virus (SUDV), Tai Forest virus
61 (TAFV), Bundibugyo virus (BDBV), and Reston virus (RESTV), respectively [2].

62 Ebola virus disease (EVD) poses a significant public health threat as implicated
63 by the increase in the incidence of EVD outbreaks in Africa over the past two decades
64 with some occurring in previously unaffected areas [3]. The most recent epidemic of EVD
65 severely affected Sierra Leone, Guinea, and Liberia, and emphasizes the need for rapid,
66 sensitive, reliable and virus-specific diagnostic tests to control the spread of the virus.
67 Rapid and simple antigen-detection tests such as immunochromatography (IC) assays
68 utilizing filovirus-specific monoclonal antibodies (mAbs) are likely one of the options for
69 early diagnosis of filovirus infections in the field setting.

70 EBOV particles consist of seven structural proteins, the nucleoprotein (NP), viral
71 protein (VP) 35, VP40, glycoprotein (GP), VP30, VP24, and polymerase [1, 4]. The NP
72 appears to be an ideal target antigen for IC assays because of its abundance in filovirus
73 particles, its strong antigenicity, and the presence of common epitopes among ebolavirus
74 species [5]. The average EBOV particle contains about 3,200 NP molecules [6]. The NP
75 plays an important role in the replication of the viral genome and is essential for formation

76 of the ribonucleocapsid [7]. Coexpression of NP VP40 and GP in cultured cells leads to
77 efficient production of virus-like particles (VLPs) containing NP in the core [6, 8].

78 Previously, we generated a panel of mouse monoclonal antibodies (mAbs)
79 recognizing the EBOV NP and identified their epitopes, some of which are shared among
80 multiple ebolavirus species, whereas others are species-specific [5]. Using these mAbs,
81 we generated an IC assay (QuickNaviTM-Ebola) and evaluated its ability to detect the NP
82 antigen in culture supernatants of infected Vero E6 cells and in sera collected from EBOV-
83 infected nonhuman primates (NHPs).

84

85

86 **MATERIALS AND METHODS**

87 **Viruses and Cells**

88 EBOV (Mayinga, Kikwit, Makona C05, and C07), SUDV (Boniface), TAFV
89 (Pauléoula), BDBV (Butalya), RESTV (Pennsylvania), and MARV (Angola) were
90 propagated in African green monkey kidney Vero E6 cells and stored at -80°C. Virus titers
91 were determined as focus-forming units (FFU) by immunoplaque assays. All experiments
92 involving the use of infectious filoviruses were performed in the Biocontainment Level 4
93 (BSL4) laboratories of the Integrated Research Facility at the Rocky Mountain
94 Laboratories (RML), Division of Intramural Research, National Institute of Allergy and
95 Infectious Diseases, National Institutes of Health, Hamilton, Montana, USA. Standard
96 operating procedures for infectious work were approved by the RML Biosafety
97 Committee.

98

99 **Preparation of VLPs and purified recombinant NPs (rNPs)**

100 Plasmids encoding NP, VP40, and GP of EBOV (Mayinga and Makona C07),
101 SUDV, TAFV, BDBV, and RESTV were constructed as described previously [5, 9]. VLPs
102 were produced by transfection of 293T cells with plasmids expressing ebolavirus NP,
103 VP40, and GP as described previously [6, 8]. Forty-eight hours after transfection,
104 supernatants containing VLPs were harvested and used for kit evaluation assays. For the
105 preparation of purified NPs, 293T cells were transfected with the plasmids expressing the
106 NPs of the respective ebolaviruses. Forty-eight hours later, the cells were lysed, and the
107 NP fraction was collected by discontinuous CsCl gradient centrifugation as described
108 previously [6, 8].

109

110 **ELISA**

111 Enzyme-linked immunosorbent assay (ELISA) was performed to determine the
112 reactivities of mAbs with rNPs as previously described [5]. Briefly, 96-well ELISA plates
113 (Nunc, Maxisorp) were coated with purified ebolavirus rNPs (2 µg/ml in PBS, 50 µl/well)
114 overnight at 4°C, followed by blocking with 1% bovine serum albumin, and purified
115 mAbs (1 µg/ml or 4-fold serial dilutions from 10 µg/ml) were added. Bound antibodies
116 were visualized with horseradish peroxidase-conjugated goat anti-mouse IgG (H+L)
117 (Jackson ImmunoResearch) and 3,3',5,5'-tetramethylbenzidine (Sigma).

118

119 **Evaluation of the IC assay**

120 Tissue culture supernatants from Vero E6 cells infected with filoviruses (EBOV,
121 SUDV, TAFV, BDBV, RESTV, or MARV), sera collected from NHPs (rhesus and
122 cynomolgus macaques) experimentally infected with EBOV, and EBOV-infected patients
123 were used to determine the sensitivity and specificity of the IC assay. For each assay, 30
124 µl (serum/plasma and culture supernatant samples) or 10-20µl (whole blood samples)
125 were used. Other human pathogens were used to test for cross-reactivity of the assay
126 (Supplementary Table 1).

127

128 **Ethical statement**

129 NHP serum samples [10-13] were collected in compliance with the Animal
130 Welfare Act and other federal statutes and regulations relating to animals and experiments
131 involving animals, and adhered to the principles stated in the Guide for the Care and Use
132 of Laboratory Animals, National Research Council, 2011. EBOV-infected human blood
133 samples were collected from patients during the 2014 EBOV outbreak in DR Congo.

134 Blood collection during outbreak investigations was approved under a special response
135 protocol established between the World Health Organization and national authorities. The
136 experiments involving the use of human samples were approved by the institutional ethics
137 committee, (Denka Seiken Co., LTD, Niigata, Japan) in accordance with the Declaration
138 of Helsinki.

139

140

141 **RESULTS**

142 **Selection of mAbs for the IC assay**

143 We first analyzed the binding affinities of 10 different clones of NP-specific
144 mAbs in ELISA with purified rNPs of the representative isolates of all known ebolavirus
145 species (EBOV, SUDV, TAFV, BDBV, and RESTV), including the 2014 EBOV Makona
146 isolate, as antigens (Table 1 and Supplementary Figure 1). To select two mAbs employed
147 in the IC assay (i.e., one labeled mAb and the other capturing mAb), we produced a test
148 device based on a lateral-flow IC assay and assessed all possible combinations of mAbs
149 suitable for labeled and capturing mAbs to detect EBOV VLPs. We found two
150 combinations of mAbs giving the highest sensitivity (ZNP105-7/ZNP108-2-5 and
151 ZNP105-7/ZNP62-7) (data not shown). Based on their cross-reactivity profiles, we
152 selected the ZNP105-7 (labeled)/ZNP108-2-5 (capturing) mAbs since this combination
153 was expected to have the potential to detect ebolaviruses from the different species (i.e.,
154 EBOV, TAFV, and BDBV), whereas the other mAb combination was EBOV-specific. The
155 binding assay with the chimeric rNPs between EBOV and SUDV indicated that both
156 mAbs ZNP105-7 and ZNP108-2-5 recognized amino acids at positions 632-739 of the
157 NP molecule (Supplementary Figure 2). These amino acids are located in the C-terminal
158 region of NP, which has been reported to have strong antigenicity [14].

159

160 **Sensitivity and specificity of the IC assay to detect ebolaviruses in tissue culture** 161 **supernatants**

162 Using ZNP105-7 and ZNP108-2-5 as labeled and capturing mAbs, respectively,
163 we produced an IC device (QuickNaviTM-Ebola) (Fig. 1). For one test, 10-30 µl of
164 samples can be applied onto the sample pad of this IC assay, followed by the addition of

165 2 drops (approximately 40 μ l) of the sample buffer (saline-based reagent) supplied with
166 the kit. The results can be interpreted 10 min later as positive by the appearance of both
167 control and test lines and negative if only the control line appears. We first assessed the
168 specificity and sensitivity of the IC assay by analyzing 10-fold serially diluted infectious
169 filoviruses (10^1 - 10^6 FFU/ml) and found that the assay was reactive for EBOV Mayinga
170 (10^3 FFU/ml), EBOV Makona (10^4 FFU/ml), TAFV (10^3 FFU/ml), and BDBV (10^3
171 FFU/ml), but not SUDV, RESTV, and MARV (Table 2). The spectrum of virus detection
172 was consistent with the reactivity profile of the mAbs chosen for the IC assay. To further
173 evaluate the sensitivity of the assay, we used purified rNPs and estimated that the limit of
174 the detection (LOD) ranged from 0.03 to 0.3 μ g/ml depending on the isolates (Table 2).
175 Other human pathogens, including viruses and bacteria, some of which potentially cause
176 hemorrhagic fever, were also tested (Supplementary Table 1) but did not react in the assay.
177 In addition, we confirmed that the presence of these pathogens in human plasma did not
178 interfere with the reactivity of the IC assay to detect EBOV VLPs.

179

180 **Application of the IC assay for EVD diagnosis**

181 We next evaluated the utility of the IC assay using serum samples from
182 experimentally infected NHPs (Table 3). Serum samples were collected on the indicated
183 days after infection, and virus titers were determined. For the IC assay, undiluted serum
184 samples (30 μ l) were directly applied and the results were obtained 10 min later. We found
185 that the IC assay was able to detect EBOV and BDBV antigens in most of the samples
186 containing infectious viruses that were detectable in 50% tissue culture infectious dose
187 (TCID₅₀) assays (Table 3). Consistent with the data of the tissue culture experiment, the
188 LOD of the IC assay seemed to be 10^3 - 10^4 FFU/ml. It was noted that EBOV in the serum

189 became detectable simultaneously in TCID₅₀ and the IC assay on day 3 or 4 after infection,
190 at a time before significant hematologic changes (e.g., a rapid increase of ALT and
191 decrease of platelet counts) appeared (Figure 2), suggesting that our IC assay was capable
192 of detecting the viral antigen shortly after disease onset. We further confirmed that the IC
193 assay could specifically detect the EBOV NP antigen in plasma or blood samples
194 collected from EBOV-infected humans during the 2014 outbreak in DR Congo.
195 Unfortunately, we were only able to obtain a limited number of samples without
196 information about disease onset of the person; therefore a thorough evaluation of the IC
197 assay sensitivity could not be performed.

198

199 **Performance evaluation of the IC assay**

200 To evaluate the stability of the IC assay, we compared its performance at different
201 temperatures and humidity conditions mimicking mild, tropical, and severe conditions,
202 and noted that it was not significantly decreased even under tropical conditions
203 (Supplementary Tables 2-4). The influence of potentially interfering substances on the
204 sensitivity and specificity of the IC assay was also evaluated (Supplementary Table 5).
205 No interference was observed for hemoglobin, conjugated bilirubin, intrafat, triglyceride,
206 ibuprofen, ribavirin, promethazine, ampicillin, acetaminophen, or quinine. The following
207 substances also exhibited no significant effect on the test results up to the specified
208 concentrations: free bilirubin 25 mg/dl; doxycycline 195 µM; ciprofloxacin 10 mg/ml;
209 and aspirin 15 mg/ml. Rheumatoid factor negatively impacted the test results in a
210 concentration-dependent manner, causing false-positive and/or false-negative reactions at
211 a concentration higher than 200 IU/ml.

212

213 **DISCUSSION**

214 The 2014 EVD outbreak has raised many serious concerns regarding the
215 development of effective strategies to control this deadly infectious disease. One of the
216 key components to minimize the spread of EVD is early diagnosis. However, clinical
217 signs and symptoms of EVD are mainly non-specific, often leading to misdiagnosis as
218 other more frequent infectious diseases that are endemic to the area present with the same
219 symptoms [15]. Consequently, there was usually a delay between the initial case
220 identification and the laboratory-confirmed diagnosis of EVD in most of the past
221 outbreaks. In addition, the initial cases and the subsequent spread of disease at the
222 beginning of an outbreak often occur in remote, sometimes inaccessible areas in the
223 affected countries, thereby making the collection of samples for laboratory diagnosis
224 difficult and resulting in a long time-lapse prior to awareness of the EVD outbreak [16].
225 These facts further underline the need for diagnostic methods that can be used on-site,
226 enabling immediate commencement of appropriate intervention measures upon
227 recognition of suspected EVD cases.

228 For early diagnosis of EVD in suspected cases, detection of viral RNA by
229 quantitative real-time RT-PCR (qRT-PCR) and viral antigens by ELISA are the
230 recommended methods [17, 18]. Although these methods are highly sensitive, specific,
231 and relatively rapid, they require a special laboratory setup, equipment and personnel
232 training. Furthermore, the samples need to be transported to the main diagnostic
233 laboratory, often leading to a delay in diagnosis and subsequent commencement of
234 measures to control the spread of the disease. On the other hand, IC assays are
235 successfully used for rapid diagnosis of various diseases such as influenza, human
236 adenoviral infection, streptococcal infection, etc. [19]. The advantages of these assays are

237 their simplicity of use, lack of need for special training, and no electricity requirement,
238 with significant reliability. In this study, we successfully developed an IC assay for easy
239 and rapid diagnosis of EVD using mAbs specific to ebolavirus NPs.

240 Currently, there are three rapid diagnosis assays approved by the FDA and/or
241 WHO, ReEBOV (Corgenix), OraQuick Ebola (OraSure Technologies, Inc.), and SD Q
242 Line Ebola Zaire Ag (SD Biosensor Inc.) [20]. ReEBOV uses whole blood samples or
243 plasma samples and has been most widely used in both point-of-care and reference
244 laboratory facilities in West Africa. This assay uses caprine polyclonal antibodies against
245 EBOV VP40. Two independent studies reported that the LOD of ReEBOV was 10^6
246 plaque-forming units/ml or 211 million RNA copies/ml (maximum CT value of 26.3).
247 The OraQuick Ebola, which is also based on the detection of VP40, can detect EBOV,
248 SUDV and BDBV. The LOD of this assay, according to the manufacturer, is 1.64×10^6
249 TCID₅₀/ml. The SD Q Line Ebola Zaire Ag has three capture lines coated with mouse
250 mAbs specific for EBOV GP, NP, or VP40 and one control line. The LODs of GP, NP,
251 and VP40 are 31.3 ng/ml, 3.9 ng/ml and 62.5 ng/ml, respectively. On the other hand, the
252 LODs of QuickNavi™-Ebola were 10^3 - 10^4 FFU/ml for infectious ebolaviruses and 33
253 ng/ml (1 ng/30 µl/test) for purified EBOV rNP, suggesting that our IC assay has at least
254 comparable or even better sensitivity than the above-mentioned rapid diagnostic assays.

255 It has been reported that viremia is often not detectable during the first few days
256 after EBOV infection but increases steeply with severe disease onset in many EBOV-
257 infected patients, and the viral load eventually reaches 10^6 - 10^9 plaque-forming units/ml
258 in the blood [21, 22]. In a NHP model of EVD, a similar correlation between the disease
259 onset and increased viremia was observed [10, 23, 24]. QuickNavi™-Ebola could detect
260 5.6×10^2 and 1.8×10^3 TCID₅₀/ml of EBOV (Kikwit) in the sera of the infected NHPs

261 before a significant increase of ALT and a decrease in platelet count were observed
262 (Figure 2). While the virus was not detected in the blood or sera of these NHPs on day 2
263 after infection by our IC assay, qRT-PCR, or TCID₅₀ assay, samples collected on day 4
264 after EBOV infection became positive both in our IC assay and qRT-PCR. It should be
265 noted that this correlation between increased viremia and onset of clinical symptoms was
266 commonly seen in both Kikwit- and Makona- infected NHPs and there was not a
267 remarkable difference in the timing of detectable viremia between these NHPs, although
268 Makona-infected NHPs eventually showed a little longer survival [23]. Indeed,
269 QuickNavi™-Ebola could detect EBOV (Makona) in the sera of infected NHPs
270 concomitantly with a significant increase of viremia (Table 3). These results suggest that
271 the sensitivity of QuickNavi™-Ebola is reliable and high enough for the practical use in
272 EVD diagnosis, particularly for the detection of an initial EVD case in remote areas and
273 also for the first screening of suspected patients in Ebola treatment units.

274 Moreover, the absence of cross-reactivity and interference with other tested
275 pathogens such as hemorrhagic fever viruses (e.g., Lassa virus and Crimean-Congo
276 hemorrhagic fever virus), bacteria causing acute diarrhea or dysentery (e.g., *Salmonella*
277 *Typhimurium*, *Shigella sonnei*, and *Vibrio cholerae*), and *Plasmodium falciparum*
278 suggests the high specificity and reliability of QuickNavi™-Ebola. Accumulatively, these
279 findings support the suitability of our IC assay for clinical application for the diagnosis
280 of EVD in endemic settings in Africa. Since the idea of developing IC assays comes from
281 the on-site utility and performance under rough and tough field conditions, we confirmed
282 its stability to be satisfactory in both high temperature and high humidity conditions. An
283 ongoing study suggests a shelf-life time of at least 8 months at room temperature. These
284 data indicate that QuickNavi™-Ebola does not require refrigeration for transportation and

285 storage, and can be used under severe conditions without special instruments in remote
286 areas in Africa.

287 Although EBOV has caused the majority of human infections, other ebolaviruses
288 (e.g., SUDV and BDBV) have also repeatedly caused outbreaks in Central Africa. Since
289 it is not predictable which ebolavirus species will cause EVD in future, the first priority
290 for the initial case identification should be to quickly clarify EVD or other viral diseases,
291 enabling us to act immediately in response to suspected outbreaks, followed by ebolavirus
292 species-specific countermeasures once it is confirmed. In summary, QuickNaviTM-Ebola
293 showed high sensitivity and high specificity for EBOV, TAFV, and BDBV. We have
294 already produced anti-SUDV NP mAbs and determined the optimal mAb combination
295 that could detect SUDV (10^3 FFU/ml) (data not shown). With the addition of these SUDV
296 NP-specific mAbs, our IC assay will have the potential to readily detect all known African
297 ebolaviruses. Although further analyses are needed to assess the clinical applicability of
298 QuickNaviTM-Ebola using human samples, our findings suggest that this IC assay based
299 on the detection of NP is a valuable tool for the rapid diagnosis of EVD.

300

301 **Notes**

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315

316 **Potential conflicts of interest.** All authors: No reported conflicts.

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320

321

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- 391

392

393 **Figure legends**

394

395 **Figure 1.** Design of the IC assay. QuickNavi™-Ebola is a lateral flow-based IC kit
396 (85mm x 21mm x 6.9mm) for the direct detection of ebolavirus NP antigens. For each
397 assay, 30 µl of serum/plasma or 10-20µl of whole blood samples can be applied. When a
398 sample is added to the sample window of the kit, it migrates through the reagent pads via
399 capillary action. One of the reagent pads is a conjugate pad on which latex conjugated
400 with an anti-ebolavirus NP mAb has been dried. The ebolavirus NP antigens present in
401 the sample bind to the latex-conjugated mAbs on the pad. Another ebolavirus NP-specific
402 mAb is also immobilized on a nitrocellulose membrane at the Test line position to capture
403 the complexes of NP antigens and mAbs conjugated with latex. As the assay develops,
404 those complexes deposit a visible blue line.

405

406 **Figure 2.** Hematological changes and TCID₅₀ of NHPs infected with EBOV. Virus titers
407 in the blood samples were determined as TCID₅₀ in Vero E6 cells (Please also see #1 and
408 #2 in Table 3). Platelet counts were determined from whole blood samples. Alanine
409 aminotransferase levels (ALT) were determined from serum. The arrows indicate the time
410 at which the NHPs tested were positive in the IC assay.

Table 1. Binding profiles of anti-NP mAbs.

mAb	Isotype	EBOV (Mayinga)	EBOV (Makona C07)	SUDV	TAFV	BDBV	RESTV
ZNP31-1-8	IgG1	++	++	++	++	++	++
ZNP41-2-4	IgG1	++	++	++	++	++	++
ZNP74-7	IgG1	++	++	++	++	++	++
ZNP24-4-2	IgG1	++	++	++	++	++	++
ZNP106-9	IgG1	++	++	+	++	++	-
ZNP108-2-5	IgG1	++	++	-	++	++	-
ZNP105-7	IgG1	++	++	-	+	++	++
ZNP98-7	IgG2a	++	++	-	-	++	-
ZNP35-16-3-5	IgG1	++	++	-	-	-	-
ZNP62-7	IgG2b	++	++	-	-	-	-

Antibody binding was evaluated based on ELISA OD₄₅₀ values at a mAb concentration of 1 µg/ml. ++, OD ≥ 1.0, +, 0.2 < OD < 1.0; -, OD ≤ 0.2.

Table 2. Detection of infectious ebolaviruses and rNP by QuickNavi™-Ebola.

Virus	Limit of detection (LOD)	
	Virus titer (FFU/ml)	Purified rNP (µg/ml)
EBOV (Mayinga)	1×10^3	0.033
EBOV (Makona C07)	1×10^4	0.15
SUDV	$>1 \times 10^5$	>30
TAFV	1×10^4	0.33
BDBV	1×10^3	0.15
RESTV	$>2 \times 10^5$	>30
MARV	$>2 \times 10^5$	>30

Serial dilutions of the samples (30 µl) were applied to the sample pad of the device, and LOD are expressed as the lowest titers (virus) and concentrations (rNP) that were positive in the IC assay.

Table 3. Detection of ebolaviruses in the sera of infected monkeys.

NHP	Virus	Days post infection	TCID ₅₀ /ml	CT value ^b	IC assay
#1 (rhesus) ^a	EBOV (Kikwit)	0	- ^c	-	-
		2	-	-	-
		4	1.8x10 ³	34.94	+
		6	5.6x10 ⁶	20.66	+
		8	3.2x10 ⁷	17.6	+
#2 (rhesus) ^a	EBOV (Kikwit)	0	-	-	-
		2	-	-	-
		4	5.6x10 ²	35.1	+
		6	5.6x10 ³	25.32	+
		8	3.2x10 ⁶	18.98	+
#3 (rhesus) ^d	EBOV (Makona C05)	0	-	ND ^e	-
		2	-	ND	-
		4	-	ND	-
		6	1.8x10 ⁶	ND	+
		8	5.6x10 ⁴	ND	+
#4 (rhesus) ^d	EBOV (Makona C05)	0	-	ND	-
		2	-	ND	-
		4	-	ND	-
		6	3.2x10 ⁶	ND	+
		8	5.6x10 ⁵	ND	+
#5 (rhesus) ^d	EBOV (Makona C05)	0	-	ND	-

		2	-	ND	-
		4	1.8x10 ⁴	ND	+
		6	5.6x10 ⁶	ND	+
		8	1.8x10 ⁴	ND	+
#6 (cynomolgus) ^d	EBOV (Makona C07)	0	-	ND	-
		3	3.2x10 ²	ND	-
		6	3.2x10 ⁶	ND	+
#7 (cynomolgus) ^d	EBOV (Makona C07)	0	-	ND	-
		3	3.2x10 ³	ND	-
		5	1.8x10 ⁷	ND	+
#8 (cynomolgus) ^a	BDBV	3	ND	ND	-
		7	ND	ND	+
		10	ND	ND	+
#9 (cynomolgus) ^d	MARV	0	-	ND	-
		4	1.8x10 ⁵	ND	-
		7	3.2x10 ⁷	ND	-

^a Gamma-irradiated serum samples were used.

^b Real-time RT-PCR cycle threshold values.

^c Not detected.

^d Intact serum samples were used.

^e Not determined.

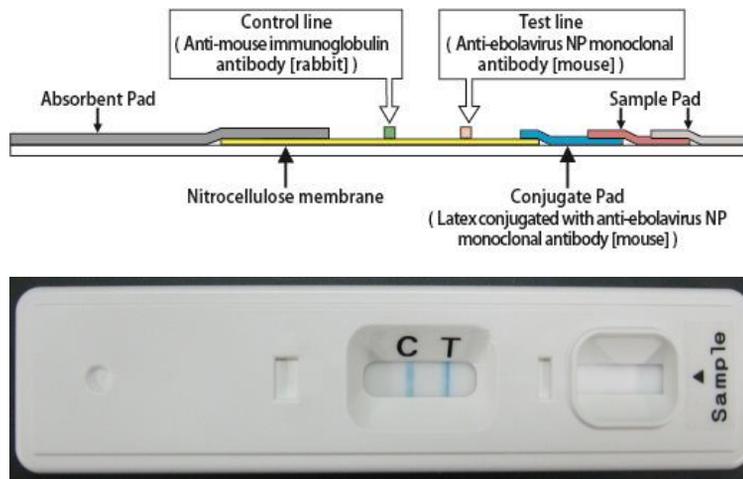


Fig. 1

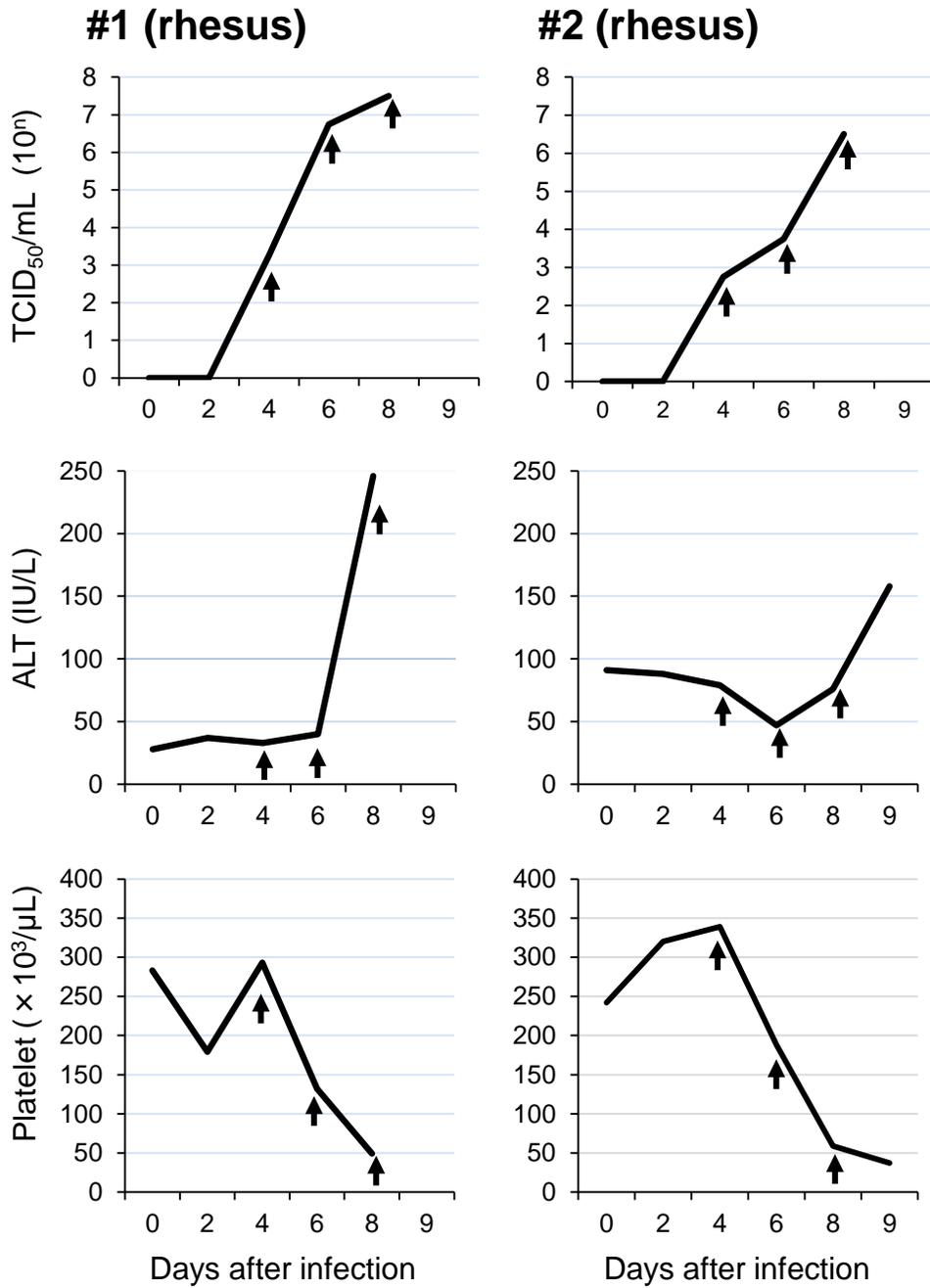


Fig. 2