



Title	Enzyme-Linked Immunosorbent Assay for Detection of Filovirus Species-Specific Antibodies
Author(s)	Nakayama, Eri; Yokoyama, Ayaka; Miyamoto, Hiroko; Igarashi, Manabu; Kishida, Noriko; Matsuno, Keita; Marzi, Andrea; Feldmann, Heinz; Ito, Kimihito; Saijo, Masayuki; Takada, Ayato
Citation	Clinical and Vaccine Immunology, 17(11), 1723-1728 https://doi.org/10.1128/CVI.00170-10
Issue Date	2010-11
Doc URL	http://hdl.handle.net/2115/45376
Rights	Copyright © 2010, American Society for Microbiology
Type	article (author version)
File Information	CVI17-11_1723-1728.pdf



[Instructions for use](#)

1 **Enzyme-linked immunosorbent assay for the detection of filovirus species-specific**
2 **antibodies**

3
4
5 Eri Nakayama¹, Ayaka Yokoyama¹, Hiroko Miyamoto¹, Noriko Kishida², Keita
6 Matsuno¹, Andrea Marzi³, Heinz Feldmann³, Kimihito Ito¹, Ayato Takada^{1*}.

7
8
9 ¹Department of Global Epidemiology, Hokkaido University Research Center for
10 Zoonosis Control, Sapporo, Hokkaido, Japan

11 ²Laboratory of Influenza Virus Surveillance, Center for Influenza Virus Research,
12 National Institute of Infectious Diseases, Tokyo, Japan

13 ³Laboratory of Virology, Division of Intramural Research, National Institute of Allergy
14 and Infectious Diseases. National Institutes of Health, Rocky Mountain Laboratories,
15 NIAID, NIH, Hamilton, MT, USA

16
17
18 Running title: **Filovirus species-specific ELISA**

19
20
21 *Corresponding author. Mailing address: Department of Global Epidemiology,
22 Hokkaido University Research Center for Zoonosis Control, Kita-20, Nishi-10,
23 Kita-ku, Sapporo 001-0020, Japan. Phone: +81-11-706-9502. Fax: +81-11-706-7310.
24 E-mail: atakada@czc.hokudai.ac.jp

25 **Abstract**

26 Several enzyme-linked immunosorbent assays (ELISA) for the detection of
27 filovirus-specific antibodies have been developed. However, diagnostic methods to
28 distinguish antibodies specific to the respective species of filoviruses, which provide the
29 basis for serological classification, are not readily available. We established an ELISA
30 using His-tagged secreted forms of the transmembrane glycoproteins (GPs) of five
31 different Ebola viruses (EBOV) species and one Marburg virus (MARV) strain as
32 antigens for the detection of filovirus species-specific antibodies. The GP-based
33 ELISA was evaluated by testing antisera collected from mice immunized with virus-like
34 particles, as well as humans and nonhuman primates infected with EBOV or MARV.
35 In our ELISA, little cross-reactivity of IgG antibodies was observed in most of the
36 mouse antisera. Although sera and plasma from some patients and monkeys showed
37 notable cross-reactivity with the GPs from multiple filovirus species, the highest
38 reactions of IgG were uniformly detected against the GP antigen homologous to the
39 virus species that infected individuals. We further confirmed that MARV-specific IgM
40 antibodies were specifically detected in specimens collected from patients during the
41 acute phase of infection. These results demonstrate the usefulness of our ELISA for
42 diagnostics as well as ecological and serosurvey studies.

43 **Introduction**

44 Ebola virus (EBOV) and Marburg virus (MARV) belong to the family *Filoviridae*,
45 and cause severe hemorrhagic fever in primates (18). While MARV consists of a
46 single species, *Lake Victoria marburgvirus*, four distinct EBOV species are known:
47 *Zaire ebolavirus* (ZEBOV), *Sudan ebolavirus* (SEBOV), *Côte d'Ivoire ebolavirus*
48 (CIEBOV), and *Reston ebolavirus* (REBOV). The phylogenetically distinct
49 *Bundibugyo ebolavirus* (BEBOV) was recently identified in Uganda and was proposed
50 as a new species of EBOV (Fig. 1) (29).

51 EBOV and MARV are filamentous, enveloped, single-stranded, negative-sense
52 RNA viruses. The virus genome encodes seven structural proteins, nucleoprotein (NP),
53 polymerase cofactor (VP35), matrix protein (VP40), glycoprotein (GP),
54 replication-transcription protein (VP30), minor matrix protein (VP24), and
55 RNA-dependent RNA polymerase (L). EBOV also express at least one secreted
56 non-structural glycoprotein (sGP) (18). GP is responsible for receptor binding and
57 fusion of the viral envelope with host cell membranes (9, 20, 32) and has an important
58 role in the pathogenesis of filovirus infection (3, 21, 33). GP is the main target of
59 neutralizing antibodies, and most of the known ZEBOV-specific monoclonal antibodies
60 (MAbs) show little cross-reactivity to other filovirus species (22, 25, 31).

61 Serological diagnostic methods based on enzyme-linked immunosorbent assays
62 (ELISA) using the recombinant EBOV and MARV NP antigens have been developed to
63 detect filovirus-specific antibodies (4, 15). Using ZEBOV NP antigen, NP-specific
64 antibodies were broadly detected in animals infected with ZEBOV, SEBOV, CIEBOV,
65 or REBOV (15) indicating strong cross-reactivity among EBOV species. It is
66 predicted, however, that the antibody response to GP is more species-specific due to the

67 larger genetic variability with this protein which is supposed to be the main target of the
68 host humoral immune response. Therefore, in this study we developed a filovirus
69 species-specific ELISA using recombinant GP antigens to serologically distinguish
70 filovirus species.

71 **Materials and Methods**

72 **Plasmids.** Viral RNA extracted from the supernatant of Vero E6 cells infected
73 with ZEBOV, SEBOV, CIEBOV, BEBOV, REBOV, or MARV strain Angola, was used
74 for cloning of the respective GP cDNAs lacking the transmembrane domain and
75 cytoplasmic tail. The cDNAs of truncated EBOV and MARV GPs with a C-terminal
76 histidine (His) tag (His-EBOV-GP, His-MARV-GP) were cloned into a pATX vector.
77 Finally, the cDNA fragments of His-EBOV-GP and His-MARV-GP were inserted into
78 the mammalian expression vector pCAGGS/MCS, which contains the chicken β -actin
79 promoter (11). All clones were sequence confirmed prior to expression.

80 **Monoclonal antibodies (MAb).** The hybridoma cells producing EBOV
81 GP-specific MAb ZGP42/3.7 (IgG1) (22, 24), which recognizes a linear epitope on GP
82 comprising the sequence GEWAFWENKKN, and MARV GP-specific MAb
83 AGP127-8 (IgG1) were grown in DMEM (Sigma) and RPMI (Sigma), respectively,
84 supplemented with fetal calf serum (FCS) and antibiotics. Mouse ascites were
85 obtained by a standard procedure and MAbs were purified from ascites fluid using
86 protein A agarose columns (Bio-Rad). The S139/1 monoclonal antibody (IgG2a),
87 which binds to the hemagglutinin of influenza A viruses (34), was used as a negative
88 control.

89 **Sera and plasma.** Five-week-old female BALB/c mice were twice immunized
90 intraperitoneally with 100 μ g virus-like particles (VLPs) (12, 19) in a 3-week interval,
91 and the serum samples were collected 7-10 days after the second immunization.
92 Convalescent phase plasma samples were collected from cynomolgus macaques
93 vaccinated and/or infected with EBOV as described previously (25). ZEBOV
94 convalescent human plasma (patients 2 to 7) and serum (patients 1 and 8) samples were

95 obtained 51 to 135 days after the onset of ZEBOV infection during the 1995 outbreak in
96 Kikwit, Democratic Republic of the Congo (23). SEBOV convalescent patient serum
97 samples (patients 9 and 10) were collected about two months after onset during the
98 Ebola hemorrhagic fever 2000 outbreaks in Uganda associated with SEBOV in 2000 (2).
99 These EBOV-infected human samples were kindly provided by Dr. T.G. Ksiazek
100 (Centers for Disease Control and Prevention). MARV-infected human blood samples
101 (patients 11 to 21) were collected within a few days after the onset of symptoms from
102 patients admitted of the 2004/05 outbreak in Angola (27). Blood collection during
103 outbreak investigations were approved under special response protocol established
104 between the World Health Organization and national authorities.

105 **Expression and purification of His-EBOV-GP and His-MARV-GP.** Human
106 epithelial kidney 293T cells cultured in high glucose DMEM containing 10% FCS and
107 antibiotics were transfected with pCAGGS expressing His-EBOV-GP
108 (pCHis-ZEBOV-GP, pCHis-SEBOV-GP, pCHis-CIEBOV-GP, pCHis-BEBOV-GP, or
109 pCHis-REBOV-GP) or His-MARV-GP (pCHis-MARV-GP) using TransIT LT1 (Mirus).
110 Forty-eight hours after transfection, the supernatants were collected and the
111 recombinant GPs were purified using the Ni-NTA Purification System (Invitrogen)
112 according to the manufacturer's instructions. The majority of contaminant protein was
113 removed with wash buffer containing 15mM imidazole. Finally, bound proteins were
114 collected with elution buffer containing 250mM imidazole. To monitor inevitable
115 nonspecific reactions (i.e., nonspecific antibodies) to FCS-derived impurities in each GP
116 preparation, control antigens (FCS-derived proteins non-specifically bound to the
117 Ni-beads) were prepared using the Ni-NTA column under the same conditions. The
118 eluted protein was concentrated using Amicon Ultra 4 spin columns (Millipore) and

119 dialyzed against PBS at 4°C overnight. Purified His-EBOV-GP and His-MARV-GP
120 were analyzed by sodium dodecyl sulfate-polyacrylamide gel electrophoresis
121 (SDS-PAGE) and stained with Coomassie brilliant blue. Western blotting was
122 performed using ZGP42/3.7, AGP127-8, and anti-His MAbs (COVANCE).

123 **ELISA.** ELISA plates (Nunc Maxisorp) were coated with the purified GPs or
124 control antigens (100ng/50µl/well) in PBS at 4°C overnight, and then washed with PBS
125 containing 0.05% Tween 20 (PBST). Unspecific binding of the antibodies was
126 avoided by blocking with 3% skim milk (150µl/well) for 2 hours at room temperature.
127 Monkey plasma samples were pre-incubated with 2% FCS to absorb antibodies to FCS
128 components, since they were exposed to FCS by injection of the vaccines or viruses
129 diluted in DMEM containing FCS. After washing three times with PBST, 50µl of
130 appropriately diluted serum/plasma samples or the GP-specific MAb in PBST
131 containing 1% skim milk was added and incubated for 1 hour at room temperature.
132 After washing three times with PBST, the bound antibodies were detected using the
133 following secondary antibodies conjugated with horseradish peroxidase diluted in 1%
134 skim milk in PBST: goat anti-mouse IgG (Jackson ImmunoResearch), goat anti-monkey
135 IgG (ROCKLAND), goat anti-human IgG (Jackson ImmunoResearch), or donkey
136 anti-human IgM (Jackson Immuno Research). After incubation for 1 hour at room
137 temperature and three PBST washes, 50 µl of 3,3',5,5'- tetramethylbenzidine were added
138 to each well and incubated for 15 minutes at room temperature. Reaction was stopped
139 by adding 1N sulfuric acid and the optical density (OD) at 450 nm was measured.

140 **Phylogenetic analysis.** Phylogenetic analysis was based on whole amino acid
141 sequences of filovirus GPs. The sequences were analyzed using GENETYX (Genetyx
142 Corp., Japan) for Windows Version 7 software. A phylogenetic tree was constructed

143 using the neighbor-joining bootstrap method (1,000 replicates) in MEGA 4.0 software
144 (26). Amino acid sequences of ZEBOV (strain Mayinga-76), ZEBOV (strain
145 Kikwit-95), SEBOV (strain Boniface-76), SEBOV (strain Maleo-79), CIEBOV (strain
146 Côte d'Ivoire-94), BEBOV, REBOV (strain Reston-89), REBOV (strain Siena
147 Philippine-92), MARV (strain Musoke-80), and MARV (strain Angola/2005) used in
148 phylogenetic analyses were obtained from GenBank under accession numbers Q05320,
149 P87666, Q66814, Q66798, Q66810, ACI28624, Q66799, Q89853, P35253, and
150 Q1PD50, respectively.

151 **Results**

152 **Expression and purification of recombinant EBOV and MARV GPs.**

153 Expression and secretion of His-EBOV-GP and His-MARV-GP in the supernatants of
154 293T cells transfected with a plasmid encoding His-GP was confirmed by
155 immunoblotting using anti-GP and anti-His MAbs (data not shown). These
156 recombinant GPs were purified as described in Materials and Methods. All purified
157 His-GPs were detected by SDS-PAGE and immunoblotting, using anti-GP and anti-His
158 MAbs as prominent protein bands of the predicted size of the transmembrane
159 anchor-minus EBOV and MARV GPs (Fig. 2). These purified GPs were used as
160 antigens for the ELISA described in the following experiments.

161 **Sensitivity of the GP-based ELISA.** The sensitivity of the purified GP-based
162 ELISA was tested using anti-EBOV-GP MAb ZGP42/3.7 and anti-MARV-GP MAb
163 AGP127-8. Serial 10-fold dilutions of the antibodies (10^{-5} to 10^2 $\mu\text{g/ml}$) were
164 prepared and the reactivity to each GP antigen was examined (Fig. 3). The negative
165 control MAb, S139/1, did not bind to any His-GPs in the ELISA. At concentrations
166 ranging from 0.1 $\mu\text{g/ml}$ to 100 $\mu\text{g/ml}$, ZGP42/3.7 reacted with all His-EBOV-GPs but not
167 His-MARV-GP, whereas AGP127-8 reacted specifically with His-MARV-GP but not
168 any of the His-EBOV-GP. The detection limit for specific antibodies using this assay
169 was approximately 0.01-0.1 $\mu\text{g/ml}$.

170 **Specificity of the GP-based ELISA.** Next, the species-specificity of the ELISA
171 was assessed by testing the antisera of mice immunized with VLP containing the
172 respective EBOV and MARV GPs. We found that species-specific IgG antibodies
173 were clearly detected in these mouse antisera (Fig. 4). All the anti-EBOV IgG
174 antibodies in the sera showed low reactivity to heterologous EBOV GPs, and no

175 cross-reactivity to MARV GP was found (Figs. 4a-e). Similarly, anti-MARV VLP
176 serum antibodies reacted to MARV GP but not to EBOV GPs (Fig. 4f). These results
177 indicated that this GP-based ELISA sufficiently detected filovirus species-specific
178 antibodies.

179 **Analysis of clinical samples in GP-based ELISA.** To further confirm the
180 specificity of our ELISA, we used convalescent-phase plasma samples obtained from
181 monkeys experimentally infected with ZEBOV or SEBOV (Fig. 5). We detected IgG
182 antibodies in the ZEBOV-infected monkey plasma with higher reactivity against
183 His-ZEBOV-GP than against any heterologous GP antigens. Although IgG antibodies
184 in the SEBOV-infected monkey plasma showed binding to all His-EBOV-GPs, the
185 highest reactivity was observed with the homologous antigen, His-SEBOV-GP.
186 Neither of these plasma antibodies reacted with MARV GP.

187 We then examined IgG antibody levels in serum/plasma derived from ZEBOV-,
188 SEBOV- and MARV-infected patients (Fig. 6a). In most of the samples tested, IgG
189 antibodies to homologous GP antigens were detected with highest reactivity (Fig 5a).
190 All of the samples derived from ZEBOV-infected patients cross-reacted with
191 His-CIEBOV-GP and His-BEBOV-GP antigens, whereas only one of the
192 SEBOV-infected human samples (no. 9) showed cross-reactivity with His-REBOV-GP.
193 Overall, the level of cross-reactivity was consistent with the phylogenetic relationship
194 among EBOV species (Fig. 1). On the other hand, in most of the samples from
195 patients infected with Angola MARV, IgG antibodies to His-MARV-GP were
196 specifically detected, except for specimens 17 and 21, which showed no IgG response to
197 any GP. Interestingly, IgG antibodies detected in specimen 11 showed remarkable
198 cross-reactivity with the heterologous antigens, His-CIEBOV-GP and His-BEBOV-GP.

199 We next evaluated whether GP-specific IgM antibodies could be detected in the
200 patient serum/plasma samples using the GP-based ELISA (Fig. 6b). ZEBOV- or
201 SEBOV-specific IgM antibodies were only detected in patients 2 and 10. In contrast,
202 MARV-specific IgM antibodies were detected in 8 out of the 11 specimens derived
203 from MARV Angola-infected patients. No obvious IgM cross-reactivity to
204 heterologous GP antigens was found in these samples (data not shown).

205 **Discussion**

206 In this study, we established a GP-based ELISA to detect filovirus species-specific
207 antibodies. To date, lysates from Vero E6 cells infected with live EBOV and MARV
208 or recombinant EBOV and MARV NPs have been used as antigens in ELISA for
209 detection of filovirus-specific antibodies (4, 6, 15). Since the NPs of EBOV and
210 MARV contain similar amino acid sequences (16), common antibody epitopes seem to
211 be present (10). Indeed, cross-reactivity was to be expected among all EBOV species
212 (14, 15). Therefore, NP antigens may be useful for the detection of genus-specific
213 antibodies but not for the detection of species-specific humoral responses (6, 14, 15)

214 Heterogeneity of EBOV and MARV GPs has been demonstrated at the genetic
215 level through sequence analyses (15, 17). An ELISA using recombinant ZEBOV GP
216 expressed in a baculovirus-insect cell expression system was reported previously (14),
217 but it is known that the protein glycosylation pathways in insect cells differ from those
218 in mammalian cells (5). This may significantly affect the antigenic properties of
219 filovirus GPs, since large amounts of both N- and O-linked carbohydrate chains are
220 present in GP molecules. To overcome this difficulty, we used mammalian 293T cells
221 for the expression of GP antigens, and verified the sensitivity and specificity of
222 GP-based ELISAs. Our results were consistent with a previous study suggesting that
223 anti-EBOV GP antibodies were highly species-specific and showed little
224 cross-reactivity to GPs of other EBOV species (25). These findings indicated that
225 most antibodies induced against filovirus GPs recognized epitopes in the variable
226 regions of the protein. Expectedly, the serological classification mirrors the
227 phylogenetic relationship of the different GPs (Fig. 1). Interestingly, serological
228 characterization of anti-BEBOV antibodies clearly supports the molecular

229 investigations (29) suggesting that BEBOV represents a new species within the EBOV
230 genus. (Fig. 7).

231 IgG antibodies in some of the serum and plasma samples collected from infected
232 monkeys and humans showed appreciable cross-reactivity to heterologous antigens,
233 whereas antibodies in the mouse sera produced by immunization with VLPs specifically
234 reacted to the homologous antigens. This result led us to conjecture that VLP
235 immunization and live virus infection induce a distinct antibody repertoire or that the
236 antibody repertoire of mice differs from that of primates. Interestingly, the serum of
237 patient 11 infected with Angola MARV contained IgG, but not IgM antibodies,
238 cross-reactive to His-CIEBOV-GP and His-BEBOV-GP. It might be possible that
239 prior to infection with Angola MARV, this patient was infected with CIEBOV, BEBOV,
240 or another unknown filovirus whose GP has epitopes shared among CIEBOV and
241 BEBOV. In the sera of patients 17 and 21, neither IgG nor IgM antibodies were
242 readily detected. An explanation for this observation might be the difference of
243 immunological conditions in individuals. Or, alternatively, the serum samples have
244 been collected before a detectable antibody response was induced.

245 Notably, our GP-based ELISA detected Angola MARV-specific IgM antibodies in
246 most of the sera collected during the acute or subacute phase of infection, although it
247 has been reported that detection of antibodies is only of limited use to for acute case
248 diagnosis due to a lack of detectable antibody response (7). The present study suggests
249 that if proper antigen and sensitive assays is available IgM antibodies can be useful for
250 the diagnosis of acute EBOV and MARV infections, and support the use of antigen
251 capture ELISA and reverse transcription-PCR, the most commonly used technologies .

252 Despite the more recent discovery of REBOV in domestic pigs in the Philippines

253 (1) and the discovery of fruit bat species as potential reservoirs for EBOV and MARV
254 (8, 13, 28, 30), the search for the reservoirs and potential amplifying hosts remains
255 ongoing. Advanced diagnostic technologies are welcome here and our new GP-based
256 species-specific antibody detection ELISA may be useful tools for future ecological and
257 seroepidemiological studies in endemic areas of Central Africa and part of Asia.

258 **Acknowledgments**

259 We thank Aiko Ohnuma for technical assistance and Kim Barrymore for editing
260 the manuscript. This work was supported by a grant-in-aid from the Ministry of
261 Health, Labor and Welfare of Japan, and in part by the Takeda Science Foundation and
262 the Program of Founding Research Centers for Emerging and Reemerging Infectious
263 Diseases and Global COE Program "Establishment of International Collaboration
264 Centers for Zoonosis Control" from the Ministry of Education, Culture, Sports, Science
265 and Technology, Japan. The work was further supported by the Division of Intramural
266 Research of the National Institute of Allergy and Infectious Diseases, National Institutes
267 of Health.

268 **Reference**

- 269 1. **Barrette R. W., Metwally S. A., Rowland J. M., Xu L., Zaki S. R., Nichol S. T.,**
270 **Rollin P. E., Towner J. S., Shieh W. J., Batten B., Sealy T. K., Carrillo C.,**
271 **Moran K. E., Bracht A. J., Mayr G. A., Sirios-Cruz M., Catbagan D. P.,**
272 **Lautner E. A., Ksiazek T. G., White W. R., McIntosh M. T.** 2009. Discovery of
273 swine as a host for the Reston ebolavirus. *Science*. Jul 10;325(5937):204-6.
- 274 2. **CDC.** 2001. Outbreak of Ebola hemorrhagic fever Uganda, August 2000–January
275 2001. *M.M.W.R. Morb. Mortal. Wkly. Rep.* 50, 73–77.
- 276 3. **Hoenen T., Groseth A., Falzarano D., Feldmann H.** 2006. Ebola virus:
277 unravelling pathogenesis to combat a deadly disease. *Trends Mol Med.*
278 12(5):206-15.
- 279 4. **Ikegami T., Saijo M., Niikura M., Miranda M. E., Calaor A. B., Hernandez M.,**
280 **Manalo D. L., Kurane I., Yoshikawa Y., Morikawa S.** 2003. Immunoglobulin G
281 enzyme-linked immunosorbent assay using truncated nucleoproteins of Reston
282 Ebola virus. *Epidemiol Infect.* Jun;130(3):533-9.
- 283 5. **Jarvis D. L., Kawar Z. S., Hollister J. R.** 1998. Engineering N-glycosylation
284 pathways in the baculovirus-insect cell system. *Curr Opin Biotechnol.*
285 Oct;9(5):528-33.
- 286 6. **Ksiazek T. G., Cynthia P. W., Pierre E. R., Peter B. J., Peters C. J.** 1999a.
287 ELISA for the Detection of Antibodies to Ebola Viruses. *J Infect Dis.* 179, S192-8.
- 288 7. **Ksiazek T. G., Rollin P. E., Williams A. J., Bressler D. S., Martin M. L.,**
289 **Swanepoel R., Burt F. J., Leman P. A., Khan A. S., Rowe A. K., Mukunu R.,**
290 **Sanchez A., Peters C. J.** 1999b. Clinical virology of Ebola hemorrhagic fever
291 (EHF): virus, virus antigen, and IgG and IgM antibody findings among EHF

- 292 patients in Kikwit, Democratic Republic of the Congo, 1995. *J Infect Dis.* Feb;179
293 Suppl 1:S177-87.
- 294 8. **Leroy E. M., Kumulungui B., Pourrut X., Rouquet P., Hassanin A., Yaba P.,**
295 **Délicat A., Paweska J. T., Gonzalez J. P., Swanepoel R.** 2005. Fruit bats as
296 reservoirs of Ebola virus. *Nature.* Dec 1;438(7068):575-6.
- 297 9. **Manicassamy B., Wang J., Rumschlag E., Tymen S., Volchkova V., Volchkov**
298 **V., Rong L.** 2007. Characterization of Marburg virus glycoprotein in viral entry.
299 *Virology.* Feb 5;358(1):79-88. Epub 2006 Sep 20.
- 300 10. **Niikura M., Ikegami T., Saijo M., Kurata T., Kurane I., Morikawa S.** 2003.
301 Analysis of linear B-cell epitopes of the nucleoprotein of ebola virus that
302 distinguish ebola virus subtypes. *Clin Diagn Lab Immunol.* Jan;10(1):83-7.
- 303 11. **Niwa H., Yamamura K., Miyazaki J.** 1991. Efficient selection for
304 high-expression transfectants with a novel eukaryotic vector. *Gene.* Dec
305 15;108(2):193-9.
- 306 12. **Noda T., Sagara H., Suzuki E., Takada A., Kida H., Kawaoka Y.** 2002. Ebola
307 virus VP40 drives the formation of virus-like filamentous particles along with GP. *J*
308 *Virol.* May;76(10):4855-65.
- 309 13. **Pourrut X., Souris M., Towner JS., Rollin P. E., Nichol S. T., Gonzalez J. P.,**
310 **Leroy E.** 2009. Large serological survey showing cocirculation of Ebola and
311 Marburg viruses in Gabonese bat populations, and a high seroprevalence of both
312 viruses in *Rousettus aegyptiacus*. *BMC Infect Dis.* Sep 28;9:159.
- 313 14. **Prehaud C., Hellebrand E., Coudrier D., Volchkov V. E., Volchkova V. A.,**
314 **Feldmann H., Le Guenno B., Bouloy M.** 1998. Recombinant Ebola virus
315 nucleoprotein and glycoprotein (Gabon 94 strain) provide new tools for the

- 316 detection of human infections. *J Gen Virol.* 79. 2565-72.
- 317 15. **Saijo M., Niikura M., Morikawa S., Ksiazek T. G., Meyer R. F., Peters C. J.,**
318 **Kurane I.** 2001. Enzyme-Linked Immunosorbent Assays for Detection of
319 Antibodies to Ebola and Marburg Viruses Using Recombinant Nucleoproteins. *J*
320 *Clin Microbiol.* 39, 1-7
- 321 16. **Sanchez A., Kiley M. P., Klenk H. D., Feldmann H.** 1992. Sequence analysis of
322 the Marburg virus nucleoprotein gene: comparison to Ebola virus and other
323 non-segmented negative-strand RNA viruses. *J Gen Virol.* Feb;73 (Pt 2):347-57.
- 324 17. **Sanchez A., Trappier S. G., Ströher U., Nichol S. T., Bowen M. D., Feldmann**
325 **H.** 1998. Variation in the glycoprotein and VP35 genes of Marburg virus strains.
326 *Virology.* Jan 5;240(1):138-46.
- 327 18. **Sanchez A., T. W. Geisbert, and H. Feldmann.** 2006. Filoviridae: Marburg and
328 Ebola viruses, p. 1409–1448. *In* D. M. Knipe, P. M. Howley, D. E. Griffin et al.
329 (ed.), *Fields virology*, 5th ed. Lippincott Williams & Wilkins, Philadelphia, PA.
- 330 19. **Swenson D. L., Warfield K. L., Kuehl K., Larsen T., Hevey M. C., Schmaljohn**
331 **A., Bavari S., Aman M. J.** 2004. Generation of Marburg virus-like particles by
332 co-expression of glycoprotein and matrix protein. *FEMS Immunol Med Microbiol.*
333 *Jan 15;40(1):27-31.*
- 334 20. **Takada A, Robison C, Goto H, Sanchez A, Murti K. G, Whitt M. A, Kawaoka**
335 **Y.** 1997. A system for functional analysis of Ebola virus glycoprotein. *Proc Natl*
336 *Acad Sci U S A.* Dec 23;94(26):14764-9.
- 337 21. **Takada A., and Kawaoka Y.** 2001. The pathogenesis of Ebola hemorrhagic fever.
338 *Trends Microbiol.* Oct;9(10):506-11.
- 339 22. **Takada A., Feldmann H., Stroehel U., Bray M., Watanabe S., Ito H.** 2003a.

- 340 Identification of protective epitopes on Ebola virus glycoprotein at the single amino
341 acid level using recombinant vesicular stomatitis viruses. *J Virol.* 77:1069–74.
- 342 23. **Takada A., Feldmann H., Ksiazek T. G., Kawaoka Y.** 2003b.
343 Antibody-dependent enhancement of Ebola virus infection. *J Virol.*
344 Jul;77(13):7539-44.
- 345 24. **Takada A., Ebihara H., Jones S., Feldmann H., Kawaoka Y.** 2007. Protective
346 efficacy of neutralizing antibodies against Ebola virus infection. *Vaccine.* Jan
347 22;25(6):993-9.
- 348 25. **Takada A., Ebihara H., Feldmann H., Geisbert TW., Kawaoka Y.** 2007.
349 Epitopes required for antibody-dependent enhancement of Ebola virus infection. *J*
350 *Infect Dis.* Nov 15;196 Suppl 2:S347-56.
- 351 26. **Tamura K., Dudley J., Nei M., Kumar S.** 2007. MEGA4: Molecular
352 Evolutionary Genetics Analysis (MEGA) software version 4.0. *Molecular Biology*
353 *and Evolution* 10.1093/molbev/msm092.
- 354 27. **Towner J. S., Khristova M. L., Sealy T. K., Vincent M. J, Erickson B. R, Bawiec**
355 **D. A, Hartman A. L., Comer J. A., Zaki S. R, Ströher U., Gomes da Silva F.,**
356 **del Castillo F., Rollin P. E., Ksiazek T. G., Nichol S. T.** 2006. Marburgvirus
357 genomics and association with a large hemorrhagic fever outbreak in Angola. *J*
358 *Virol.* Jul;80(13):6497-516.
- 359 28. **Towner J. S., Pourrut X., Albariño C. G., Nkogue C. N., Bird B. H., Grard G.,**
360 **Ksiazek T. G., Gonzalez J. P., Nichol S. T., Leroy E. M.** 2007. Marburg virus
361 infection detected in a common African bat. *PLoS One.* Aug 22;2(1):e764.
- 362 29. **Towner J. S., Sealy T. K., Khristova M. L., Albariño C. G., Conlan S., Reeder**
363 **S. A., Quan P. L., Lipkin W. I., Downing R., Tappero J. W., Okware S.,**

- 364 **Lutwama J., Bakamutumaho B., Kayiwa J., Comer J. A., Rollin P. E., Ksiazek**
365 **T. G., Nichol S. T.** 2008. Newly discovered ebola virus associated with
366 hemorrhagic fever outbreak in Uganda. *PLoS Pathog.* Nov;4(11):e1000212. Epub
367 2008 Nov 21.
- 368 30. **Towner J. S., Amman B. R., Sealy T. K., Carroll S. A., Comer J. A., Kemp A.,**
369 **Swanepoel R., Paddock C. D., Balinandi S., Khristova M. L., Formenty P. B.,**
370 **Albarino C. G., Miller D. M., Reed Z. D., Kayiwa J. T., Mills J. N., Cannon D.**
371 **L., Greer P. W., Byaruhanga E., Farnon E. C., Atimmedi P., Okware S.,**
372 **Katongole-Mbidde E., Downing R., Tappero J. W., Zaki S. R, Ksiazek T. G.,**
373 **Nichol S. T., Rollin P. E.** 2009. Isolation of genetically diverse Marburg viruses
374 from Egyptian fruit bats. *PLoS Pathog.* Jul;5(7):e1000536. Epub 2009 Jul 31.
- 375 31. **Wilson J. A., Hevey M., Bakken R., Guest S., Bray M., Schmaljohn A. L., Hart**
376 **M. K.** 2000. Epitopes involved in antibody-mediated protection from Ebola virus.
377 *Science.* 287(5458):1664-6.
- 378 32. **Wool-Lewis R. J., and Bates P.** 1998. Characterization of Ebola virus entry by
379 using pseudotyped viruses: identification of receptor-deficient cell lines. *J Virol.*
380 Apr;72(4):3155-60.
- 381 33. **Yang Z. Y., Duckers H. J., Sullivan N. J., Sanchez A., Nabel E. G., Nabel G. J.**
382 2000. Identification of the Ebola virus glycoprotein as the main viral determinant of
383 vascular cell cytotoxicity and injury. *Nat Med.* Aug;6(8):886-9.
- 384 34. **Yoshida R., Igarashi M., Ozaki H., Kishida N., Tomabechi D., Kida H., Ito K.,**
385 **Takada A.** 2009. Cross-protective potential of a novel monoclonal antibody
386 directed against antigenic site B of the hemagglutinin of influenza A viruses. *PLoS*
387 *Pathog.* Mar;5(3):e1000350. Epub 2009 Mar 20.

388

389 **Figure Legends**

390

391 **Fig. 1.** Phylogenetic analysis of filovirus GP amino acid sequences. The
392 phylogenetic tree was constructed using the neighbor-joining method. For
393 construction of this tree, we used 10 GP amino acid sequences, each comprising a
394 whole GP amino acid sequence. Numbers at branch points indicate bootstrap values
395 (1,000 replicates).

396

397 **Fig. 2.** Identification and characterization of purified His-GPs. His-EBOV-GP and
398 His-MARV-GP were analyzed on 8% SDS-PAGE and stained with Coomassie brilliant
399 blue (a). Immunoblotting of purified His-GPs were performed using MAbs to EBOV
400 (ZGP42/3.7), MARV GPs (AGP127-8) (b) and His-tag (c). Arrows indicate the
401 location of the His-GPs. The protein bands represent His-ZEBOV-GP (lane 1),
402 His-SEBOV-GP (lane 2), His-CIEBOV-GP (lane 3), His-BEBOV-GP (lane 4),
403 His-REBOV-GP (lane 5), and His-MARV-GP (lane 6). Lane 7 shows FCS-derived
404 proteins used as a control antigen (see Materials and Methods).

405

406 **Fig. 3.** Sensitivity of GP-based ELISA. Serial 10-fold dilutions of MAbs to EBOV
407 (a) and MARV (b) were prepared and tested. S139/1 (specific to influenza virus
408 hemagglutinin) was used as a negative control antibody (c).

409

410 **Fig. 4.** IgG antibodies detected in mouse antisera. Serial 10-fold dilutions of the
411 anti-ZEBOV (a), anti-SEBOV (b), anti-CIEBOV (c), anti-BEBOV (d), anti-REBOV (e),

412 and anti-MARV (f) sera obtained from mice immunized with EBOV and MARV VLPs
413 were tested for IgG antibodies reacting with His-GPs.

414

415 **Fig. 5.** IgG antibodies detected in experimentally infected monkey sera. C105, C332,
416 C508, and C725 were infected with ZEBOV, whereas C0287 and C0436 were infected
417 with SEBOV. All sera were diluted 1:1,000. Naïve monkey serum was used as a
418 negative control.

419

420 **Fig. 6.** IgG and IgM antibodies detected in human sera. OD values for specific IgG
421 (a) and IgM (b) antibodies in the patient sera are shown. Sera from 21 individuals
422 were analyzed in 1:1,000 dilutions. Naïve human serum (no. 22) was used as a
423 negative control.

424

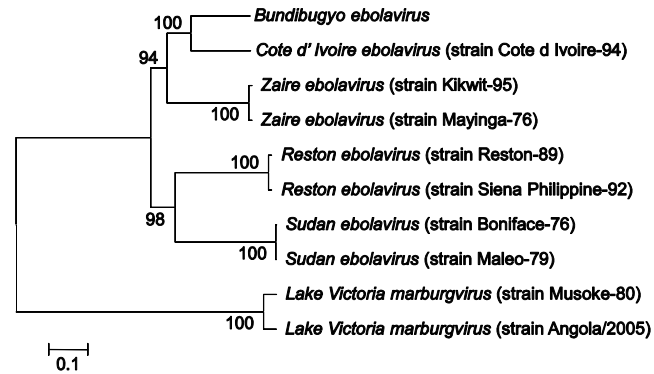


Fig. 1. Phylogenetic analysis of filovirus GP amino acid sequences. The phylogenetic tree was constructed using the neighbor-joining method. For construction of this tree, we used 10 GP amino acid sequences, each comprising a whole GP amino acid sequence. Numbers at branch points indicate bootstrap values (1,000 replicates).



Fig. 2. Identification and characterization of purified His-GPs. His-EBOV-GP and His-MARV-GP were analyzed on 8% SDS-PAGE and stained with Coomassie brilliant blue (a). Immunoblotting of purified His-GPs were performed using MAbs to EBOV (ZGP42/3.7), MARV GPs (AGP127-8) (b) and His-tag (c). Arrows indicate the location of the His-GPs. The protein bands represent His-ZEBOV-GP (lane 1), His-SEBOV-GP (lane 2), His-CIEBOV-GP (lane 3), His-BEBOV-GP (lane 4), His-REBOV-GP (lane 5), and His-MARV-GP (lane 6). Lane 7 shows FCS-derived proteins used as a control antigen (see Materials and Methods).

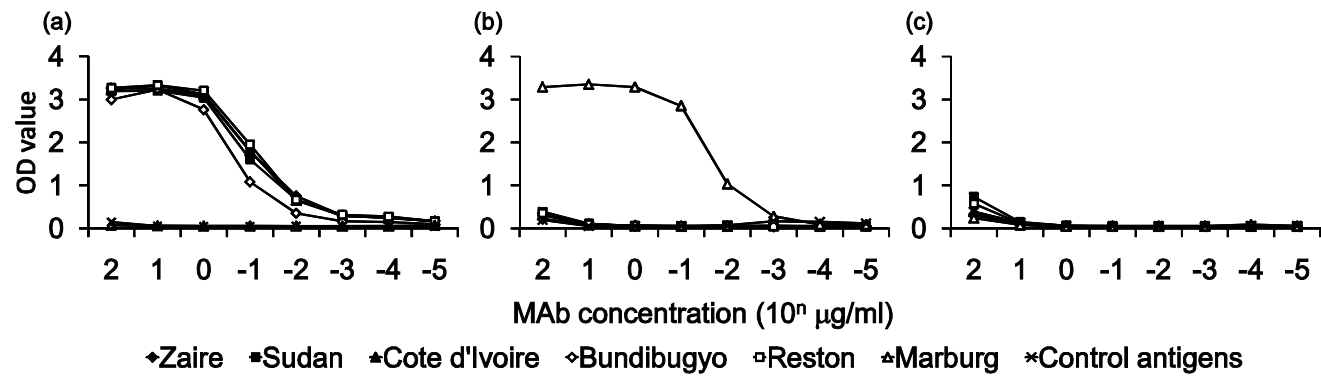


Fig. 3. Sensitivity of GP-based ELISA. Serial 10-fold dilutions of MAbs to EBOV (a) and MARV (b) were prepared and tested. S139/1 (specific to influenza virus hemagglutinin) was used as a negative control antibody (c).

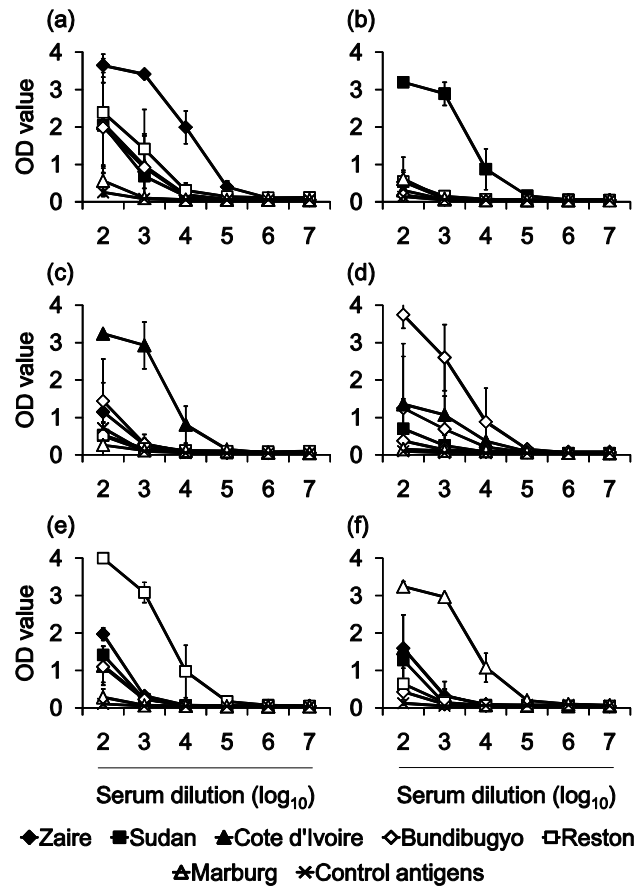


Fig. 4. IgG antibodies detected in mouse antisera. Serial 10-fold dilutions of the anti-ZEBOV (a), anti-SEBOV (b), anti-CIEBOV (c), anti-BEBOV (d), anti-REBOV (e), and anti-MARV (f) sera obtained from mice immunized with EBOV and MARV VLPs were tested for IgG antibodies reacting with His-GPs.

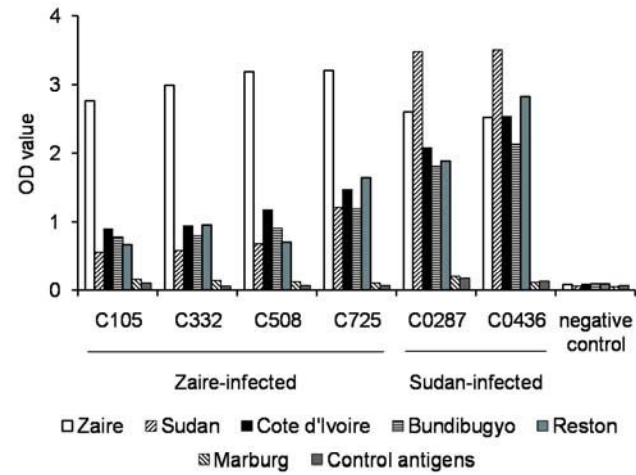


Fig. 5. IgG antibodies detected in experimentally infected monkey sera. C105, C332, C508, and C725 were infected with ZEBOV, whereas C0287 and C0436 were infected with SEBOV. All sera were diluted 1:1,000. Naïve monkey serum was used as a negative control.

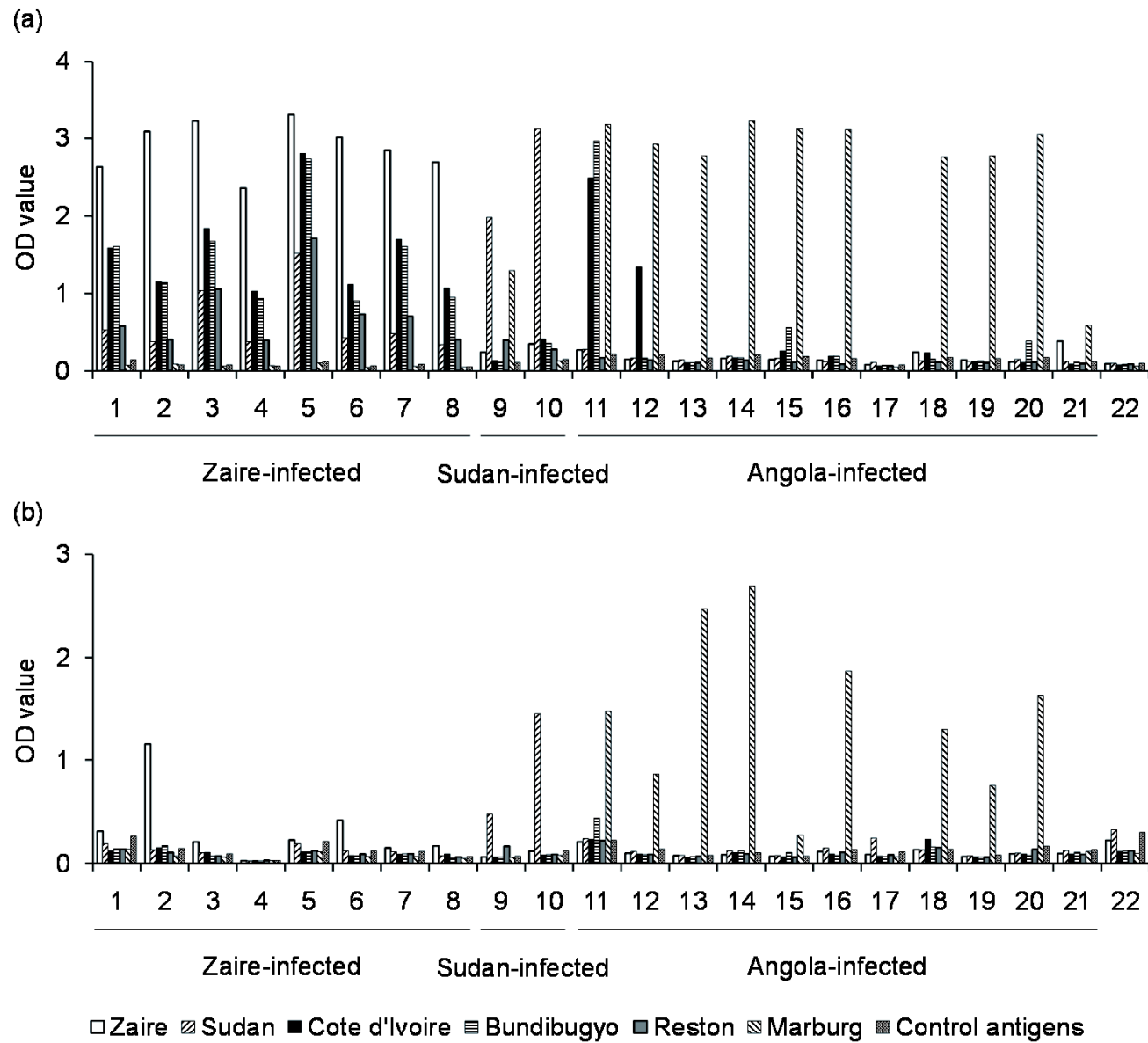


Fig. 6. IgG and IgM antibodies detected in human sera. OD values for specific IgG (a) and IgM (b) antibodies in the patient sera are shown. Sera from 21 individuals were analyzed in 1:1,000 dilutions. Naïve human serum (no. 22) was used as a negative control.