## **Supplementary Information**

Title: A Critica	l Domain of	Ebolavirus	Envelope	Glycoprotein	Determines
Glycoform and Infectivity					
Authors and Affi	liations				
Haruhiko Fujihira	<sup>1,2,3*</sup> , Katsuaki U	Jsami <sup>1</sup> , Keita	Matsuno <sup>4,5</sup> ,	Hideyuki Takeu	ıchi <sup>1,6</sup> , Kaori
Denda-Nagai <sup>1,2</sup> , Ju	ın-ichi Furukaw	a <sup>7,8</sup> , Yasuro S	hinohara <sup>7,9</sup> ,	Ayato Takada <sup>5,1</sup>	<sup>0</sup> , Yoshihiro

7 Kawaoka<sup>11,12,13</sup>, and Tatsuro Irimura<sup>1,2\*</sup>

1

 $\mathbf{2}$ 

3

4

 $\mathbf{5}$ 

6

- 8 <sup>1</sup>Laboratory of Cancer Biology and Molecular Immunology, Graduate School of
- 9 Pharmaceutical Sciences, The University of Tokyo, Tokyo 113-0033, Japan
- <sup>10</sup> <sup>2</sup>Division of Glycobiologics, Intractable Disease Research Center, Juntendo University
- 11 Graduate School of Medicine, Tokyo 113-8421, Japan
- <sup>12</sup> <sup>3</sup>Glycometabolome Team, Systems Glycobiology Research Group, Global Research
- 13 Cluster, RIKEN, Saitama 351-0198, Japan
- <sup>14</sup> <sup>4</sup>Laboratory of Microbiology, Faculty of Veterinary Medicine, Hokkaido University,
- 15 Sapporo 060-0818, Japan
- <sup>5</sup>Division of International Services, Global Institution for Collaborative Research and
- 17 Education (GI-CoRE), Hokkaido University, Sapporo 001-0020, Japan
- <sup>18</sup> <sup>6</sup>Department of Molecular Biochemistry, Nagoya University School of Medicine,
- 19 Nagoya 4668550, Japan
- 20 <sup>7</sup>Laboratory of Medical and Functional Glycomics, Graduate School of Advanced Life
- 21 Science, Hokkaido University, Sapporo 001-0021, Japan
- <sup>8</sup>Department of Advanced clinical glycobiology, Faculty of Medicine and Graduate
- 23 School of Medicine, Hokkaido University, Sapporo 0010021, Japan
- <sup>9</sup>Department of Pharmacy, Kinjo Gakuin University, Nagoya 4638521, Japan

26	Control, Sapporo 001-0020, Japan
27	<sup>11</sup> CREST, Japan Science and Technology Agency, Saitama 332-0012, Japan
28	<sup>12</sup> Division of Virology, Department of Microbiology and Immunology, Institute of
29	Medical Science, The University of Tokyo, Tokyo 108-8639, Japan
30	<sup>13</sup> Department of Pathobiological Sciences, University of Wisconsin, Madison, WI
31	53706, USA
32	
33	Supplementary Figure Legends
34	Supplementary Figure 1. Amino acid sequences of ZGP and RGP. Amino acid
35	sequences of ZGP and RGP are shown. Red box: 33-50 amino acid residue, green box:
36	mucin-like domain, orange line: border of GP1 and GP2.
37	Supplemental Figure 2. MALDI-TOF MS spectra of <i>N</i> -glycans released from VLPs.
38	Representative MS spectra of <i>N</i> -glycans released from VLPs bearing (a) ZGP, (b) RGP,
39	(c) Z33-50R, (d) Z33-186, (e) R33-50, or (f) R33-186 are shown. Magnified spectra
40	between $m/z=3800$ and $m/z=4200$ are shown in the upper right window of each
41	spectrum.
42	Supplementary Figure 3. Biosynthetic rate and intracellular localization of RGP

<sup>10</sup>Division of Global Epidemiology, Hokkaido University Research Center for Zoonosis

25

43 and ZGP were not different. (a) Results of pulse chase experiments of [<sup>35</sup>S]-labeled

44	RGP and [ <sup>35</sup> S]-labeled ZGP are shown. (b) Results of immunofluorescent staining of
45	VLP-producing HEK293T cells. Red: GP (ZGP or RGP), Green: Golgi marker (GM130
46	or TGN46). Representative results are shown.
47	Supplementary Figure 4. Sialylation and fucosylation of N-glycans found in GPs
48	were not different. Relative ratios of sialylated (a) or fucosylated (b) N-glycans are
49	shown.
50	Supplementary Figure 5. Uncropped images of blots shown in the figures (Fig. 1c,
51	1d, 2b, 3b, 3d, 5a).
52	Supplementary Figure 6. GPs that lack 18 amino acid residues (33 through 50) is
52 53	Supplementary Figure 6. GPs that lack 18 amino acid residues (33 through 50) is expressed on the VSV pseudotyped viruses and VLPs.
53	expressed on the VSV pseudotyped viruses and VLPs.
53 54	<ul><li>expressed on the VSV pseudotyped viruses and VLPs.</li><li>(a-b) Results of immunoblotting using VSV pseudotyped viruses or VLPs containing</li></ul>
53 54 55	<ul> <li>expressed on the VSV pseudotyped viruses and VLPs.</li> <li>(a-b) Results of immunoblotting using VSV pseudotyped viruses or VLPs containing</li> <li>ZGPΔ33-50 or RGPΔ33-50 are shown. (c) Results of virus titration experiment of VSV</li> </ul>
53 54 55 56	expressed on the VSV pseudotyped viruses and VLPs. (a-b) Results of immunoblotting using VSV pseudotyped viruses or VLPs containing ZGPΔ33-50 or RGPΔ33-50 are shown. (c) Results of virus titration experiment of VSV pseudotyped ZGP, RGP, ZGPΔ33-50, and RGPΔ33-50 using Vero E6 cells. VSV

## 60 Supplementary Materials and Methods

**Pulse chase experiment.** HEK293T cells  $(6 \times 10^5)$  were seeded on a 6-well culture plate 61 one day before transfection. Cells were transfected with pCAGGS-GP and 62 pCAGGS-VP40 by using Trans-IT LT1 in the same way as in VLP production. The cells 63 were washed with PBS 24 hours after transfection and subsequently cultured with 64 DMEM-HG (10% FCS, without Met and Cys; Gibco) for 1 hour. Then the medium was 65changed to DMEM-HG (10% FCS, containing 150 µCi/mL of [35S]-labeled Met and 66 Cys) and cultured for 30 min. The medium was changed to DMED-HG (10% FCS) after 67 washing with DMEM-HG (10% FCS) and cultured for 0, 1, 2, or 3 hours. Cells were 68 collected after washing with PBS and solubilized with RIPA buffer containing protease 69 inhibitor cocktail (Calbiochem) by shaking at 4°C for 2 hours. The solution was 70 centrifuged at 21,500g for 15 min to remove cellular debris. The supernatant was 71collected, and the radioactivity of [<sup>35</sup>S] in the supernatant was measured by a liquid 72 scintillation counter (LS6000SC, BECKMAN). GPs were immunoprecipitated using 73mAb 42/3.7 from lysates containing equal amounts of [<sup>35</sup>S] radioactivity. 74Immunoprecipitated GPs were separated on an 8% SDS-PAGE gel. The gel was then 75

fixed and stained with CBB (BioRad). The gel was dried (RapiDry mini; ATTO) and
 [<sup>35</sup>S]-labeled GPs were detected with Typhoon FLA9000 (GE Healthcare).

78	<b>Immunofluorescent staining.</b> HEK293T cells $(4 \times 10^6)$ were seeded on a 10 cm culture
79	dish one day before transfection. Cells were transfected with pCAGGS-GP and
80	pCAGGS-VP40 by using Trans-IT LT1 in the same way as in VLP production. 24 hours
81	after transfection, cells were washed with PBS three times and collected. Collected cells
82	were fixed with 20 mM sodium phosphate buffer (pH 7.0) containing 4 (w/v) $\%$
83	paraformaldehyde at 4°C for 1 hour. Fixed cells were washed with 10 mM glycine in
84	PBS, and permeabilized by 0.1 (w/v) % Triton X-100 in PBS (at room temperature for 5
85	min). The cells were blocked with 4 (w/v) % BSA in PBS at room temperature for 30
86	min, and incubated with anti-EBOV GP1 mAb 42/3.7 (10 $\mu g/mL)$ and rabbit
87	anti-GM130 (10 $\mu$ g/mL; Abcam) or rabbit anti-TGN46 antibody (10 $\mu$ g/mL; Abcam),
88	followed by Alexa 568 goat anti-mouse IgG(H+L) (10 $\mu\text{g/mL};$ Molecular Probes) and
89	Alexa 488 goat anti-rabbit IgG(H+L) (10 $\mu$ g/mL; Molecular Probes) and DAPI
90	(Boehringer Mannheim). Analysis was performed using a Leica TSC SP5 confocal
91	microscope (Leica).

Chimeric GPs		Primer Sequences
Z311-462R	forward	CCGCTCGAGGTCTCGATAGAACTGTGAAAGACAACTCTTC
	reverse	CGGAATTCGGTCTCCAACACTCATCACCAAGATACC
Z297-462R	forward	CCGAATTCGGTCTCGAAAAGTTTTTTTTAGTTTCCCAGAAGGC
	reverse	CGGAATTCGGTCTCCAACACTCATCACCAAGATACC
Z260-462R	forward	CCGAATTCGGTCTCCGAAGTGTCTCATTCAGCTGGAGC
	reverse	CGGAATTCGGTCTCCAACACTCATCACCAAGATACC
Z187-462R	forward	CCGAATTCGGTCTCTCTGACAGTATCAGAAATGCAACGAC
	reverse	CGGAATTCGGTCTCCAACACTCATCACCAAGATACC
Z33-462R	forward	CGCGGTCTCGGCATGGAAAATGTTCTTTGGAAAAGG
	reverse	CGGAATTCGGTCTCCAACACTCATCACCAAGATACC
Z1-462R	forward	CCGCTCGAGGTCTCCCCATTGTGTTGTTGGATCCTC
	reverse	CGGAATTCGGTCTCCAACACTCATCACCAAGATACC
Z33-186R	forward	CGCGGTCTCCATGCCGCTTGGTATAGTG
	reverse	CGCGGTCTCCGGGGCAGAATTAAAAAAGCTACGACACC
Z33-50R	forward	CGCGGTCTCCATGCCGCTTGGTATAGTG
	reverse	CGCGGTCTCACTAGTTGATCAATTTCTGTTGCTTTG
R33-186Z	forward	CGCGGTCTCCATCCCACTTGGAGTCATC
	reverse	CGCGGTCTCTCTGACAGTATCAGAAATGCAACGACA
R33-50R	forward	CGCGGTCTCACTAGTTTGTCGTGACAAAC
	reverse	CGCGGTCTCGGCATGGAAAATGTTCTTTGGAAAAGG

## 92 Supplemental Table 1. Primer sequences that used to make chimeric GPs

Figure S1

i iguio c						
ZGP	1	MGVT-GILOL	PRDRFKRTSF	FLWVIILFOR	TF <mark>S</mark> IPLGVIH	NSTIQVSDVD
RGP	1	MGSGYQLLQL	PRERFRKTSF	LVWVIILFQR	AI <mark>S</mark> MPLGIVT	NSTLKATEID
ZGP	50	KLVCRDKLSS	TNOLRSVGLN	LEGNG <mark>V</mark> ATDV	PSATKRWGFR	SGVPPKVVNY
RGP	50	QLVCRDKLSS	TSQLKSVGLN	LEGNGIATDV	PSATKRWGFR	SGVPPKVVSY
ZGP	100	EAGEWAENCY	NLEIKKPDGS	ECLPAAPDGI	RGFPRCRYVH ******	KVSGTGPCAG
RGP	100	EAGEWAENCY	NLEIKK DGS	ECLP <mark>LP</mark> PDGV	RGFPRCRYVH	KV <mark>Q</mark> GTGPC <mark>P</mark> G
ZGP	150	DFAFHKEGAF	FLYDRLASTV	IYRGTTFAEG	VVAFLIL *****	KKDFFSSHPL
RGP	150	DLAFHKNGAF	FLYDRLASTV	IYRGTTFAEG	<b>VVAFLIL</b> SEP	<b>Ŕĸ</b> Ħ <b>F</b> ₩KATPA
ZGP	200	R <mark>EPVN</mark> ATEDP	SSGYYSTTIR	YQATG <mark>FG</mark> T <mark>NE</mark>	TEYLFEVDNL	TYVOL * * * * *
RGP	200	HEPVNTTDDS	TSYYMTLTLS	Ŷ EMSNFGGNÊ	SNTLFKVDNH	<b>TYVQL</b> DRPHT
ZGP	250	POFL ****	IYTSGKR <mark>SN</mark> T	TGKLIWKVNP	EIDTTIGEWA	FWETKKNLTR
RGP	250	**** PQFL <mark>VQLNET</mark>	LRRNNRL SNS	<b>ŤĠ</b> ŖĹ <b>ĬŴ</b> TLDP	KIEPDVGEWA	****** FWETKKN <mark>F</mark> SQ
ZGP	300	KIRSEELSET	VVSNGAKNIS	G <mark>QSPA</mark> R <mark>T</mark> SSD	PGTNTTTEDH	KIMASEN <mark>S</mark> SA
RGP	300	QLHG <mark>ÊNL</mark> H <mark>Ê</mark> Q	ILŜTHTN <mark>N</mark> SŜ	D <mark>QSPA</mark> GTVQG	KISYHPPANN	SELVPTD <sup>Ŝ</sup> PP
ZGP	350	MVQVHSQGRE	AAVSHLTTLA	TIS <mark>T</mark> SPQSLT	TKPGPDNSTH	NTPVYKLDIS
RGP	350	VVSVLTAGRT	EEMST-QGLT	NGETIT-GFT	ANPMTTTIAP	-SPTMTSEVD
ZGP	400	EATQVEQHHR	RTD-N <mark>DS</mark> TA <mark>S</mark>	DTPSATTA	ag <mark>p</mark> pkae <mark>n</mark> tn	TSKSTDFLD
RGP	400	NNVPS <mark>EQ</mark> PNN	TASIE <mark>DS</mark> PPS	ASNETIYHSE	MDPIQGS <mark>N</mark> NS	AQSPQTKTTP
ZGP	447	ATTTSP ****	ETAGNNNTHH	QDTGEES <mark>A</mark> SS	GKL <mark>GI</mark> IT <mark>NT</mark> I	AG <mark>VA</mark> GLITGG
RGP	447	AP <mark>TTSP</mark> MTQD	PQETA <mark>N</mark> SSKP	GTSPGSA <mark>A</mark> GP	SQP <mark>GL</mark> TI <mark>NT</mark> V	SK <mark>VA</mark> DSLSPT
ZGP	497	RRTRR EAI	VNAQPKCNPN		AAIGLAWIPY	FGPAAEGIYI
RGP	497	RKQKR SVR	QNTAN <mark>KCNP</mark> D	LYYWTAVDEG	AA <mark>V</mark> GLAWIPY	FGPAAEGIYI
ZGP	545	EGLMHNODGL	ICGLRQLANE	TTOALOLFLR	ATTELRTFSI	LNRKAIDFLL
RGP	545	ÊĜ <mark>V</mark> MHNQ <mark>N</mark> ĜĹ	ICGLRQLANE	TTQALQLFLR	<b>ATTELRT</b> YSL	LNRKAIDFLL
ZGP	595	QRWGGTCHIL	GPDCCIEPHD	WTKNITD *****	QI **	LPDQGDNDNW
RGP	595	QRWGGTCRIL	GPSCCIEPHD	WTKNITD <b>E</b> IN	QIKHDFIDNP	Ĺ₽ĎĦĠĎDĽŇL
ZGP	645	WTGWROWIPA	GIG <mark>VT</mark> GVIIA	VIALFCICKF	VF	
RGP	645	WTGWRQWIPA	ĜIĜ <mark>II</mark> ĜVIIA	IIALLCICK	LC	

Figure S2









