

HOKKAIDO UNIVERSITY

Title	Valosin-containing protein (VCP/p97) plays a role in the replication of West Nile virus
Author(s)	Phongphaew, Wallaya; Kobayashi, Shintaro; Sasaki, Michihito; Carr, Michael; Hall, William W.; Orba, Yasuko; Sawa, Hirofumi
Citation	Virus Research, 228, 114-123 https://doi.org/10.1016/j.virusres.2016.11.029
Issue Date	2017-01-15
Doc URL	http://hdl.handle.net/2115/67761
Rights	© 2016. This manuscript version is made available under the CC-BY-NC-ND 4.0 license http://creativecommons.org/licenses/by-nc-nd/4.0/
Rights(URL)	https://creativecommons.org/licenses/by-nc-nd/4.0/
Туре	article (author version)
File Information	(77370) Valosin-containing protein and WNV.pdf



1	Valosin-containing protein (VCP/p97) plays a role in the replication of
2	West Nile virus
3	
4	Wallaya Phongphaew ¹ , Shintaro Kobayashi ^{1,2} , Michihito Sasaki ¹ , Michael Carr ^{3,4} , William W.
5	Hall ^{3,5,6} , Yasuko Orba ¹ and Hirofumi Sawa ^{1,3,6*}
6	
7	¹ Division of Molecular Pathobiology, Research Center for Zoonosis Control, Hokkaido University,
8	N20, W10, Kita-ku, Sapporo, 001-0020, Japan
9	² Laboratory of Public Health, Graduate School of Veterinary Medicine, Hokkaido University, N18,
10	W9, Kita-ku, Sapporo, 001-0020, Japan
11	³ Global Institution for Collaborative Researches and Education (GI-CoRE), Global Station for
12	Zoonosis Control, Hokkaido University, N20, W10, Kita-ku, Sapporo, 001-0020, Japan
13	⁴ National Virus Reference Laboratory, University College Dublin, Belfield, Dublin 4, Ireland
14	⁵ Center for Research in Infectious Diseases, University College of Dublin, Belfield, Dublin 4,
15	Dublin, Ireland
16	⁶ Global Virus Network (GVN), The Institute of Human Virology, University of Maryland, 22 S.
17	Greene Street, Baltimore, MD 21201, USA
18	*Corresponding author: h-sawa@czc.hokudai.ac.jp
19	
20	
21	
22	
23	
24	
25	

26 Abstract

Valosin-containing protein (VCP) is classified as a member of the type II AAA⁺ ATPase protein family. VCP functions in several cellular processes, including protein degradation, membrane fusion, vesicular trafficking and disassembly of stress granules. Moreover, VCP is considered to play a role in the replication of several viruses, albeit through different mechanisms. In the present study, we have investigated the role of VCP in West Nile virus (WNV) infection. Endogenous VCP expression was inhibited using either VCP inhibitors or by siRNA knockdown. It could be shown that the inhibition of endogenous VCP expression significantly inhibited WNV infection. The entry assay revealed that silencing of endogenous VCP caused a significant reduction in the expression levels of WNV-RNA compared to control siRNA-treated cells. This indicates that VCP may play a role in early steps either the binding or entry steps of the WNV life cycle. Using WNV virus like particles and WNV-DNA-based replicon, it could be demonstrated that perturbation of VCP expression decreased levels of newly synthesized WNV genomic RNA. These findings suggest that VCP is involved in early steps and during genome replication of the WNV life cycle.

41 Keywords: VCP, WNV replication, early steps, genome replication

52 **1. Introduction**

53 West Nile virus (WNV) belongs to genus the Flavivirus, family Flaviviridae and has an 54 approximately 11 kb positive sense, single-stranded genomic RNA [(+)ssRNA]. The WNV genome encodes three structural proteins (C, prM and E) and seven non-structural proteins (NS1, NS2a, 55 56 NS2b, NS3, NS4a, NS4b and NS5) (Brinton, 2014; Fields et al., 2013). Many species of mammals and birds can be infected by WNV (Dauphin et al., 2004; Egberink et al., 2015; Fields et al., 2013; 57 58 Gamino et al., 2016; Kramer and Bernard, 2001; Lichtensteiger et al., 2003; Read et al., 2005) and 59 infection causes West Nile fever and encephalitis in human and horses (Dauphin et al., 2004; 60 Samuel and Diamond, 2006). WNV was firstly isolated from a Ugandan woman in 1937 (C. et al., 61 1940; Fields et al., 2013), but has now spread widely to many countries (Fields et al., 2013; Paz, 2015; Troupin and Colpitts, 2016). In the United States, approximately 44,000 cases of WNV 62 infection were reported between 1999 and 2015 (CDC, 2016). 63

WNV attaches to host cells through the interaction of the viral E protein and cellular receptors on 64 65 the surface of host cells (Fields et al., 2013). Several attachment receptors of WNV have been reported and include the laminin receptor (Bogachek et al., 2010; Perera-Lecoin et al., 2014; 66 67 Zaitsev et al., 2014; Zidane et al., 2013), TIM (T cell/transmembrane, immunoglobulin and mucin) and TAM (Tyro3, Axl and Mer) families (Carnec et al., 2016; Morizono and Chen, 2014; Perera-68 69 Lecoin et al., 2014), DC-SIGN/L-SIGN (dendritic cell-specific intercellular adhesion molecule-3-70 grabbing non-integrin) (Davis et al., 2006; Denizot et al., 2012; Martina et al., 2008; Shimojima et 71 al., 2014) and integrin αvβ3 (Bogachek et al., 2010; Fields et al., 2013; Perera-Lecoin et al., 2014; 72 Smit et al., 2011; Zaitsev et al., 2014). Following attachment, the virus is then internalized into the 73 cytoplasm via clathrin-mediated endocytosis (Brinton, 2014; Chu and Ng, 2004; Fields et al., 2013). WNV particles are delivered to early or intermediate endosomes, which mature into late 74 75 endosomes, following a conformational change of the viral E protein dimer triggered by the acidic 76 environment in late endosomes. Membrane fusion between viral particles and endosomal membranes then occurs and, thereafter, WNV genomic RNA is released into the cytosol, with 77

78 subsequent translation and replication (Chu et al., 2006; Chu and Ng, 2004; Fields et al., 2013; 79 Heinz and Allison, 2000; Smit et al., 2011). Host cell membrane rearrangements are induced during 80 replication of flaviviruses, including WNV, to coordinate the processes of genomic RNA replication and virus assembly. Viral genomic RNA replication is thought to occur in endoplasmic 81 82 reticulum (ER) membrane-derived vesicles (in structures termed vesicle packets) (Gillespie et al., 2010; Kaufusi et al., 2014; Welsch et al., 2009). Encapsidation of nascent viral genomic RNA is 83 achieved by the capsid protein and budding into the ER vielding a viral envelope coated with prM 84 85 and E proteins (Brinton, 2014; Fields et al., 2013; Suthar et al., 2013; Welsch et al., 2009). The 86 immature virions are transported via the host secretory pathway and virion maturation then occurs 87 in the acidic compartments of the Golgi by cleavage of the prM protein by a furin-like protease (Plevka et al., 2014; Roby et al., 2015; Yu et al., 2008). Mature virions are then released from the 88 89 infected cells through exocytosis (Fields et al., 2013; Samuel and Diamond, 2006). It has been 90 reported that several cellular pathways and host factors are involved in WNV infection (Ambrose 91 and Mackenzie, 2011; Brinton, 2014; Chahar et al., 2013; Chu and Ng, 2004; Courtney et al., 2012; Fernandez-Garcia et al., 2011; Fields et al., 2013; Gilfoy et al., 2009; Kobayashi et al., 2016a; 92 93 Krishnan et al., 2008; Ma et al., 2015); however, the role of valosin-containing protein (VCP) has 94 remained controversial.

95 VCP, also known as CDC48 in Saccharomyces cerevisiae, is well conserved among eukaryotes 96 with orthologues in archaea, protozoa, insects and plants (Meyer et al., 2012; Wolf and Stolz, 97 2012), and is classified as a member of the type II AAA⁺ ATPase (adenosine triphosphatase-98 associated with diverse cellular activities) family (Koller and Brownstein, 1987; Pye et al., 2006; 99 Stolz et al., 2011; Wolf and Stolz, 2012; Xia et al., 2016). VCP is a homohexameric complex 100 composed of six protomers organized as two concentric-rings with a central pore. VCP 101 conformational changes, driven by adenosine triphosphate hydrolysis, acts as a chaperone in protein 102 homeostasis systems, which include ER-associated degradation (ERAD) (Wolf and Stolz, 2012; Xia 103 et al., 2016; Zhong and Pittman, 2006) and mitochondria-associated degradation and autophagy

104 (Bug and Meyer, 2012; Dargemont and Ossareh-Nazari, 2012; Xia et al., 2016; Yamanaka et al., 105 2012) to prevent accumulation of misfolded-proteins and turnover of certain proteins. Recently, a 106 role of VCP in the disassembly of stress granules (SGs) has also been reported (Buchan et al., 2013; 107 Seguin et al., 2014). Generally, after removal of stress stimuli, SGs are disassembled by VCP and 108 mRNA in the SGs could be restored allowing mRNA translation to proceed. Otherwise, depletion 109 of VCP causes persistence of SGs leading to blockage of mRNA restoration and an arrest of mRNA 110 translation (Buchan et al., 2013). It has also been reported that VCP is involved in chromatin-111 associated degradation and several nuclear substrates of VCP have been described (Maric et al., 112 2014; Verma et al., 2011; Wilcox and Laney, 2009). Furthermore, VCP also participates in 113 membrane fusion and vesicular trafficking events (Bug and Meyer, 2012; Meyer et al., 2012; 114 Ramanathan and Ye, 2012; Ritz et al., 2011; Xia et al., 2016). VCP binds to endocytic components, 115 and silencing of VCP leads to a failure of maturation and enlargement of the early endosome 116 (Ramanathan and Ye, 2012). 117 Interestingly, VCP has also been implicated in the life cycle of several (+)ssRNA viruses. It has 118 been previously reported that VCP facilitates the replication of poliovirus (PV) (Arita et al., 2012). 119 Depletion of VCP caused a reduction of PV infection, whereas, a mutant PV, which has a secretion inhibition-negative phenotype, increases the affinity of binding to VCP and resists VCP-knockdown 120 121 compared to wild-type PV, suggesting that VCP may play a role in PV replication through cellular 122 secretion pathways (Arita et al., 2012). In addition, other roles for VCP have been described in 123 other picornaviruses (Arita et al., 2012). Although VCP knockdown strongly inhibits PV infection, 124 inhibition of VCP does not affect the replication of Coxsackievirus B3 (Arita et al., 2012), which is

also a member of the same genus *Enterovirus*. In contrast, replication of Aichivirus A, genus

126 *Kobuvirus*, another member of family *Picornaviridae*, is enhanced when VCP is depleted (Arita et

127 al., 2012).

128 A relationship between VCP and Sindbis virus (SINV) replication has also been reported (Panda

129 et al., 2013). VCP is involved in trafficking of the entry receptor of SINV, which is the natural

resistance-associated macrophage protein 2 (NRAMP2). Deficiency of VCP suppresses SINV
replication through alteration of trafficking routes of NRAMP2 leading to degradation of NRAMP2
by lysosomes.

Studies of infectious bronchitis virus (IBV), family *Coronaviridae*, have suggested that VCP is
engaged in the internalization steps of IBV (Wong et al., 2015). Depletion of VCP using siRNA
knockdown, resulted in accumulation of IBV particles in early endosomes as maturation of the
endosome and acidification was disrupted. Failure of the acidification of virus-containing
endosomes inhibited fusion between the virus envelope and endosomal membrane and prevented
IBV exit from the endosomes to the cytosol (Wong et al., 2015)
In the present study, we investigated whether VCP is involved in WNV infection. Specifically,

we employed VCP inhibitors and siRNA knockdown to elucidate a potential role of VCP in WNVreplication.

- 142
- 143 **2. Methods**

144 **2.1 Cell and viruses**

145 Human cervical adenocarcinoma cells, HeLa, were grown in Dulbecco's Modified Eagle's Medium 146 (DMEM) supplemented with 10% fetal bovine serum (FBS) and 2 mM L-glutamine. Human embryonic kidney cells, HEK293T, were grown in high glucose DMEM supplemented with 110 147 148 mg/L sodium pyruvate, 2 mM L-glutamine and 5% FBS. African green monkey kidney cells, Vero, 149 were grown with Minimum Essential Media (MEM) supplemented with 5% FBS and 2 mM L-150 glutamine. Human neuroblastoma cells, SK-N-SH, were grown with Minimum Essential Media 151 (MEM) supplemented with 10% FBS and 2 mM L-glutamine. Cells were grown at 37 °C with 5% 152 supplemented CO₂. The mosquito cell line, Aedes albopictus clone C6/36, were grown in MEM 153 supplemented with 10% FBS, 1% non-essential amino acid and 2 mM L-glutamine at 28 °C. WNV 154 New York strain (NY99 6-LP) was propagated in C6/36 at 28 °C. WNV-NY99 6-LP was kindly provided by Dr. Takashima (Laboratory of Public Health, Graduate school of Veterinary Medicine, 155

Hokkaido University, Sapporo, Japan) (Hasebe et al., 2010; Shirato et al., 2004a; Shirato et al.,
2004b). Viral titer was measured by plaque assay and stock of viruses were stored at -80 °C until
use. All experiments with WNV were performed in the Biosafety level-3 facility at the Research
Center for Zoonosis Control, Hokkaido University in accordance with institutional guidelines.
Pseudotyped vesicular stomatitis virus (VSV) was provided by Dr. Takada (Research Center for
Zoonosis Control, Hokkaido University) (Takada et al., 2007).

162

163 2.2 MTT assay

164 HeLa cells were treated with 5, 10 and 20 µM of Eevarestatin I (EerI) (Sigma Aldrich, St. Louis, 165 MO) (Wang et al., 2008; Wang et al., 2010) or 12.5, 25 and 50 μM of 3,4-Methylenedioxy-β-166 nitrostyrene (MDBN) (Abcam, Cambridge, UK) (Chou and Deshaies, 2011) and incubated at 37 °C for 24 h. Thereafter, the treated cells were examined by MTT assay following addition of 3-(4,5-167 dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) and incubated at 37 °C for 1 h, then 168 169 addition of solubilization solution [10% Triton-X 100 in acidic isopropanol (0.1N HCl)] and 170 incubation at room temperature with shaking for 30 min. Absorbance values were measured at 570 171 nm and 630 nm using the Model 680 microplate reader (Bio-rad, Hercules, CA).

172

173 **2.3 Plaque assay**

174 Virus suspensions were diluted in a series of 10-fold dilutions and inoculated onto monolayers of 175 Vero cells. The WNV-inoculated Vero cells were grown with MEM containing 1.25% methyl 176 cellulose, 5% FBS and 2 mM L-glutamine at 37 °C for 4 days. Fixation was done after 4 days using 177 10% formalin for 10 min at room temperature. The fixed cells were stained with 1% crystal violet 178 in 70% ethanol for 30 min. The number of plaques was counted and the virus titer was determined 179 in plaque forming unit per milliliter (PFU/ml).

180

181 **2.4 Immunofluorescence assay (IFA)**

182 The WNV-infected cells grown on coverslips were fixed at various times after infection. The 183 infected cells were fixed in 4% paraformaldehyde for 10 min and permeabilized using 0.1% Triton 184 X-100 for 5 min at room temperature. Blocking was performed with 1% bovine serum albumin (BSA) for 30 min before incubation with primary antibody. The cells were incubated with primary 185 186 antibody (rabbit anti-JEV serum; 1:1,500) (Kimura et al., 1994; Kobayashi et al., 2012) that have 187 cross-reactivity with the WNV antigens at 4 °C overnight, followed by incubation with Alexa Fluor 188 488-conjugated secondary antibody against rabbit IgG (1:2,000; Life technologies, Rockville, MD) 189 for 1 h at room temperature. The cells were washed three times with phosphate buffered saline 190 (PBS) before fluorescence microscopy examination. The cells were visualized using an inverted 191 fluorescence microscope (IX70, Olympus, Tokyo, Japan) and images were processed using DP 192 manager software (Olympus).

193

194 **2.5 Immunoblotting analysis**

195 Cell samples were harvested at the indicated time points using TNE lysis buffer [1% Triton X-100, 196 10 mM Tris-HCl (pH 7.5), 150 mM NaCl, 5 mM EDTA, 10% glycerol] and centrifuged at 17,800 x g at 4 °C for 20 min. Only supernatants were collected and mixed with SDS-PAGE sample buffer 197 198 [0.1 M Tris-HCl (pH 6.8), 3.3% SDS, 11% glycerol]. The samples were separated by 8% 199 polyacrylamide gel electrophoresis and then transferred to a polyvinylidene difluoride filter (Merck 200 Millipore, Billerica, MA). The filters were blocked with 5% skimmed milk for 30 min. The filters 201 were incubated with each primary antibody, including mouse anti-VCP antibody (1:2,000; Abcam), 202 mouse anti-envelope protein of West Nile/Kunjin virus (Merck 1:1,000; Millipore), rabbit anti-NS3 203 serum (1:1,000), which was prepared from a rabbit immunized by twice intravenous inoculations of 204 synthetic peptides of NS3 (CEREKVYTMDGEYRLRGEER), and mouse anti-actin (1:1,000; 205 Merck Millipore). Thereafter, the filters were incubated with secondary antibody, goat anti-mouse 206 IgG antibody conjugated with horseradish peroxidase at 1:10,000 dilution (Biosource International, 207 Camarillo, CA) and washed with TBS [50 mM Tris-HCl (pH 7.5), 150 mM NaCl,] containing

0.05% Tween 20 (TBS-T) three times. Chemiluminescence was detected by Immobilon Western
HRP Substrate (Merck Millipore) and visualized with VersaDoc 5000MP (Bio-Rad), and images
were analyzed using Quantity One software (Bio-Rad).

211

212 **2.6 WNV inoculation in the presence of VCP inhibitors**

213 Based on the results of MTT assays, we determined the optimal concentration of EerI (2.5 and 5 214 uM) and MDBN (6.25 and 12.5 uM) without cytotoxicity in HeLa cells. Multiplicity of infection 215 (MOI) of 1 of WNV NY99 6-LP strain was inoculated into HeLa cells. After 1 h of incubation at 37 216 °C with rocking, the inocula were removed. Suspensions of either EerI or MDBN diluted in normal 217 cultured medium were added to the WNV-inoculated cells and incubated at 37 °C for 24 h. 218 Thereafter, the inoculated-cells and the supernatants were prepared for IFA, plaque assay and 219 immunoblotting analysis to measure the number of WNV-infected cells, production of infectious 220 WNV and expression of WNV proteins, respectively.

221

222 2.7 VCP knockdown

- 223 The endogenous VCP was inhibited using siRNA. HeLa cells (2×10^4 cells in 250 µl medium per
- well) in 48-well plates were transfected with 5 nM of each siRNA targeting VCP, no. (1), (2) and

225 (3), which have the sequences 5'-GAAUAGAGUUGUUCGGAAUTT-3',

- 226 5'-GAACCGUCCCAAUCGGUUATT-3', and 5'-GGCUCGUGGAGGUAACAUUTT-3' (Thermo
- 227 Fisher Scientific, Waltham, MA), respectively, using lipofectamine RNAiMAX transfection reagent
- 228 (Thermo Fisher Scientific) according to the manufacturer's instructions. The transfected cells were
- incubated at 37 °C for 48 h. The expression level of VCP was evaluated using immunoblotting.
- 230

231 2.8 WNV inoculation in siRNA-treated cells

- At 48 h post transfection with either siRNA against VCP or control siRNA (Catalogue No: 439084,
- 233 Thermo Fisher Scientific), HeLa cells were inoculated with WNV NY99 6-LP strain (MOI=1) and

234 incubated at 37 °C for 1 h with rotation. Thereafter, supernatants of the cells were removed, and 235 normal HeLa culture medium was added to the cells and incubated at 37 °C for 12, 24 and 48 h. 236 The inoculated cells and the supernatants were prepared for IFA, plaque assay and immunoblotting 237 analysis, respectively. To investigate the role of VCP in the early stages of the WNV replicative 238 cycle, the siRNA-transfected cells were inoculated with WNV as described above. After inoculation 239 of WNV, cells were incubated on ice for 1 h, and then washed with PBS five times. The cells were then placed at 37 °C and incubated for 1 h. Thereafter, the inoculated-cells were harvested using 240 241 trypsin. The detached cells from cell culture plates were centrifuged at 1,500 x g for 3 min. After 242 removal of supernatants, total RNA was extracted from cell pellets using Trizol (Thermo Fisher 243 Scientific). Extracted total RNAs were analyzed for WNV genome using real-time reverse 244 transcription PCR (qRT-PCR) analysis.

245

246 2.9 Pseudotyped VSV inoculation in siRNA-treated cells

247 It has been previously reported that VCP knockdown did not affect VSV infection (Panda et al., 248 2013). Therefore, control experiments using pseudotyped VSV were performed. The pseudotyped 249 VSV encoding GFP was kindly provided by Dr. Takada (Hokkaido University) (Takada et al., 250 2007). Approximately 30% infectivity of pseudotyped VSV was inoculated into HeLa cells transfected with either siRNA against VCP or control siRNA. The cells were incubated at 37 °C 251 252 with rotation for 1 h. Thereafter, supernatants of the cells were removed, normal growth media was 253 added to the cells and incubated at 37 °C for 8 h. The percentage positivity following pseudotyped VSV infection was measured by counting the number of GFP-positive cells using an inverted 254 255 fluorescence microscope (IX70, Olympus, Tokyo, Japan).

256

257 2.10 Production of WNV virus-like particles (WNV-VLPs)

258 WNV-VLPs with reporter DsRed protein were produced following transfection. Three plasmid 259 vectors carrying WNV sequences: pCMV-WNrep-DsRed, pCMV-SVP and pCSXN-C were 260 transfected into HEK293T cells using lipofectamine 2000 (Thermo Fisher Scientific). These 261 plasmids were constructed as follows. A plasmid encoding WNV replicon cDNA, pCMV-WNrep-262 DsRed encodes the WNV non-structural (NS1-NS5) proteins. Almost all of the sequences encoding structural proteins, including C, prM and E, were deleted and replaced by the gene encoding the 263 264 DsRed protein. The 3'-terminus of the WNV genome was accomplished by containing sequences 265 enabling ribozyme-mediated post-transcriptional cleavage of the RNA (Kobayashi et al., 2016b). 266 The fragment of prM-E was amplified by PCR from pCAGGS-C-prM-E, which was a gift from Dr. 267 Takashima (Hokkaido University) (Takahashi et al., 2009) as a template, and subcloned into the 268 pCMV vector, and the plasmid was named pCMV-SVP. For the pCSXN-C, the C fragment with 269 restriction sequences of Xho I and Not I (Takara Bio, Kyoto, Japan) were amplified by PCR and 270 inserted into pCSXN-flag which was generated from pCMV-myc (Clontech Laboratories, Mountain View, CA) as previously described (Kobayashi et al., 2013), using Xho I and Not I restriction sites. 271 272 The plasmid was named as pCXSN-C. The transfected cells were incubated at 37 °C for 72 h. The 273 supernatants from transfected cells were collected and filtered through a 0.45 µm filter (Sigma 274 Aldrich). The WNV-VLPs were concentrated by ultracentifugation at 4 °C, 68,000 x g for 2 h. The 275 supernatant was discarded and only the pellet was collected after ultracentrifugation. The pellet was resuspended with normal HeLa culture medium and the titers of WNV-VLPs were measured by 276 277 hemagglutination assay as previously described (Makino et al., 2014). The titer of VLPs was 278 calculated as hemagglutination units (HAU)/50 µl based on the highest dilution of VLP suspension 279 causing agglutination of chicken red blood cells.

280

281 2.11 Inoculation of WNV-VLPs in VCP knockdown HeLa cells

WNV-VLPs (16 HAU/50 µl) were inoculated into siRNA-treated HeLa cells 48 h post transfection.
After 1 h incubation at 37 °C, the inocula were discarded and normal HeLa culture medium was
added and incubated for 72 h. Comparison of the quantity of WNV-RNA between VCP knockdown
and control was determined using aRT-PCR.

286

287 2.12 WNV-VLP production and transfection of pCMV-WNrep-DsRed to siRNA-treated

288 HeLa cells

289 Plasmid transfection to generate WNV-VLPs has been reported to investigate the role of VCP in the 290 late steps of viral life cycle, from genome replication to virus release (Kobayashi et al., 2016a). The 291 results obtained employing the plasmid-encoded VLPs would not be attributable to early steps of 292 WNV infection, including attachment and entry. WNV-VLPs were used to investigate the role of 293 VCP in distinct steps (early and genome replication steps) of WNV infection cycle. At 24 h after 294 siRNA transfection, the plasmid set (pCMV-WNrep-DsRed, pCMV-SVP and pCSXN-C) was 295 transfected into siRNA-treated HeLa cells using FuGENE HD (Promega, Madison, WI). The plasmid transfected-cells were incubated at 37 °C for 72 h. Thereafter, the supernatants from 296 297 transfected cells were collected and inoculated onto Vero cells monolayers in 10-fold serial 298 dilutions. WNV-VLP titer was calculated as infectious units (IFU)/ml based on the total number of 299 DsRed-positive cells (Figure 3C).

To investigate the role of VCP in WNV genomic RNA replication, only pCMV-WNrep-DsRed (Kobayashi et al., 2016b) was transfected into the siRNA-treated cells. The procedure and time of transfection are similar to plasmid transfection for VLP production. After 72 h incubation, the total RNAs were extracted and prepared for qRT-PCR.

304

305 2.13 Real-time reverse transcription-PCR (qRT-PCR)

306 Total RNA was isolated by Trizol and chloroform according to the manufacturer's protocol. The

307 RNA samples were treated with DNase I (Thermo Fisher Scientific) to remove genomic DNA.

308 qRT-PCR was performed with a Brilliant III Ultra-Fast qRT-PCR master mix (Agilent

309 Technologies, Santa Clara, CA) following the manufacturer's protocol. The oligonucleotide primers

and fluorescent probe targeting the 3'UTR of WNV, 5'-AAGTTGAGTAGACGGTGCTG-3' and 5'-

311 AGACGGTTCTGAGGGCTTAC-3', WNV probe, FAM-5'-GCTCAACCCCAGGAGGACTGG-3'-

- 312 BHQ, were used for detection of WNV-RNA. A TaqMan Gene expression assays kit corresponding
- 313 to human β-actin (Thermo Fisher Scientific) was used as an endogenous control. The expression
- 314 level of viral RNA was normalized to the expression of human β -actin.
- 315

316 2.14 Statistical analysis

- 317 The statistical significance was calculated using one-way ANOVA.
- 318

319 3. Results

320 **3.1 WNV infection is inhibited in the presence of VCP inhibitors.**

321 To determine if VCP is involved in WNV infection, the effect of VCP inhibitors, both EerI (Wang 322 et al., 2008; Wang et al., 2010) and MDBN (Chou and Deshaies, 2011) at concentrations without 323 cytotoxicity were assayed in WNV infection. The cytotoxicity of either EerI or MDBN treatment in 324 HeLa cells was examined using a MTT assay (Supplementary Fig. 1). Each VCP inhibitor was 325 added to HeLa cells at 1 h.p.i. with WNV. WNV-inoculated cells and cultured supernatants were 326 harvested at 24 h.p.i. The number of WNV-infected cells was examined by IFA, and this revealed 327 that the number of WNV-infected cells was significantly decreased in a dose-dependent manner in 328 the presence of either EerI or MDBN (Fig. 1A and 1B). Viral titers of supernatants from WNV-329 inoculated cells were also measured by plaque assay. Consistently, this demonstrated that viral titers 330 of supernatants from WNV-inoculated cells were significantly decreased in a dose-dependent 331 manner in the presence of either EerI or MDBN (Fig. 1C). We also confirmed the inhibitory effects 332 of EerI in WNV infection in a different cell line, human neuroblastoma SK-N-SH cells. Inhibition 333 of VCP by EerI both decreased the percentage of WNV-infected cells and viral titer in SK-N-SH 334 cells (Supplementary Fig. 2). These findings suggest that VCP may play a role in WNV infection. 335

333

336 **3.2 WNV infection is inhibited by knockdown of VCP**

337 To confirm that the inhibition of WNV infection was caused by perturbation of VCP activity, small 338 interfering RNAs (siRNAs) were employed to deplete endogenous VCP. HeLa cells were 339 transfected with either of three siRNAs targeting three different regions of the VCP gene [siVCP 340 (1), (2) and (3)] or a control siRNA (siCont) and then inoculated with WNV 48 h post transfection 341 and incubated for 24 h. The expression level of VCP after silencing was confirmed by 342 immunoblotting. Reverse transfection of siVCP (1) and (3) for 48 h strongly decreased expression levels of endogenous VCP in HeLa cells (Fig. 2A). Furthermore, depletion of endogenous VCP 343 344 reduced expression levels of WNV-E protein at 24 h.p.i. of WNV (Fig. 2A). Thereafter, we 345 examined siRNA targeting of VCP [(1) and (3)] which significantly reduced the percentage of 346 WNV-infected cells (Fig. 2B and 2C). However, siRNA (2) failed to knockdown endogenous VCP 347 as shown in the immunoblotting and IFA results (Fig. 2A, 2B and 2C). We further measured viral 348 titers in supernatants of WNV-inoculated HeLa cells treated by the siRNAs against VCP. Plaque 349 assays revealed that the viral release was significantly inhibited by siRNA treatment [siCVP (1) and 350 (3)] (Fig. 2D). We also examined the effect of siRNA against VCP on WNV infection by IFA at 351 different time points (12, 24 and 48 h). A decrease in the immunofluorescence signals between cells 352 transfected with control and VCP siRNAs was detected (Supplementary Fig. 3). These results 353 indicate that a depletion of VCP significantly inhibits WNV infection. In contrast, VCP-knockdown 354 did not affect infection by pseudotyped VSV (Supplementary Fig. 4).

355

356 **3.3 VCP** participates in the early and genome replication steps during the WNV life cycle

We next investigated the specific role of VCP in the life cycle of WNV. The intracellular life cycle of WNV is divided into two major steps, early and late (Fernandez-Garcia et al., 2011; Kaufmann and Rossmann, 2011). The early step consists of viral attachment, entry and uncoating (Jiang et al., 2010; Kaufmann and Rossmann, 2011), while the late step involves genome translation, genome replication, viral assembly and release (Kobayashi et al., 2016a). WNV-VLPs were employed to determine whether VCP participates in the early or late replication steps (Kobayashi et al., 2014). 363 WNV-VLPs are unable to produce progeny virions because of the absence of WNV structural 364 protein coding sequences in their genome. Therefore, our results are independent of the assembly 365 and virion-releasing steps (Hasebe et al., 2010; Scholle et al., 2004). Thus, WNV- VLPs allow the 366 determination of whether VCP plays a role in either an early step or during the genomic replication 367 of the WNV life cycle. WNV-VLPs were inoculated into both VCP and control siRNA-transfected 368 cells and monitored by expression of DsRed-encoded in the WNV-VLP replicon. qRT-PCR 369 demonstrated that the quantity of WNV-RNA in VCP-knockdown (siVCP (1)) cells was 370 significantly lower than that in control siRNA-treated cells (Fig. 3A). These results suggest that 371 VCP knockdown significantly inhibits infection of WNV-VLPs through an inhibition of early step 372 and/or genome replication steps of the WNV life cycle. 373 To confirm the role of VCP in the early stages of WNV replication, the siRNA-treated cells

were inoculated with WNV and viral RNA was investigated at an early time point of WNV
infection, at 2 h post infection. The result revealed that silencing of VCP (siVCP (1)) significantly
decreased the quantity of WNV-RNA at 2 h post inoculation of WNV compared to control siRNA
treated-cells (Fig. 3B). This suggests that VCP plays a role in the early step of WNV replication
cycle, including attachment or entry into cells.

379 To examine whether VCP plays a role in the late step of the WNV life cycle, three plasmids 380 encoding WNV sequences (pCMV-C, pCMV-SVP and pCMV-WNrep-DsRed) were co-transfected 381 into either VCP-knockdown (siVCP (1)) or control siRNA-treated cells. The plasmid transfection 382 protocol for VLP production has been previously employed to investigate the role of host factor(s) 383 in the late stages of viral infection, including genome replication and virus release (Kobayashi et al., 384 2016a). The components of the VLP are transfected into cells, therefore, the results obtained 385 employing the plasmid-encoded VLPs were not associated with the early steps of WNV replication 386 (Kobayashi et al., 2016a). At 72 h after plasmid transfection, we examined the WNV-VLP titers in 387 the supernatants from plasmid transfected-cells. The titer of WNV-VLPs was significantly decreased in VCP-knockdown (siVCP (1)) cells compared with control siRNA-treated cells (Fig. 388

389 3C). These data suggest that VCP is also involved in the genome replication and/or assembly andrelease steps of the WNV life cycle.

391 To confirm that VCP is important for WNV genome replication, the WNV DNA-based replicon 392 pCMV-WNrep-DsRed was introduced into both VCP knockdown and control siRNA-treated cells. 393 This replicon consists of coding sequences of WNV non-structural proteins, while almost all of the 394 sequences encoding WNV structural proteins were deleted and replaced by sequences encoding 395 DsRed. The replicon can replicate and translate to synthesize viral genomic RNA and the non-396 structural proteins of WNV, respectively. However, it is not capable of producing viral progeny 397 because of the absence of structural protein sequences of WNV. The results of qRT-PCR of the 398 plasmid-transfected cells at 72 h post transfection demonstrated a 10-fold reduction of synthesized 399 WNV-RNA in the VCP-knockdown cells (siVCP (1)) compared to control siRNA-treated cells 400 (Fig. 3D). Taken together, these data indicate that VCP plays a role in the genome replication of the 401 WNV life cycle.

402

403 4. Discussion

404 In the present study, we have investigated the roles of VCP during WNV infection. We found that 405 perturbation of endogenous VCP using a potent VCP inhibitor or siRNA targeting VCP 406 significantly inhibited WNV infection. We could confirm that the expression of endogenous VCP in 407 HeLa cells is depleted after 48 h siRNA transfection [with siVCP (1) and (3)], while another siRNA 408 siVCP (2) did not silence expression of endogenous VCP. Silencing of endogenous VCP, by siVCP 409 (1) and (3), demonstrated that depletion of VCP significantly suppressed WNV infection. This finding suggests that VCP is required for WNV infection and, thereafter, we employed the most 410 411 potent siRNA [siVCP (1)] to investigate the roles of VCP in WNV replication. A previous report 412 indicated that silencing of endogenous VCP and nuclear protein localization 4 (NPL4), a VCP 413 cofactor, did not inhibit WNV infection, whereas depletion of ubiquitin fusion degradation 1-like 414 (UFD1L) and p47, other cofactors of VCP, suppressed WNV infection (Krishnan et al., 2008). We suggest that the differences between this previous report and the present results may be related to the use of different strains of WNV or could be related to the silencing efficiency of the siRNA employed in the experiments. In addition, control experiments from the present study using pseudotyped VSV demonstrated that depletion of VCP did not suppress pseudotyped VSV infection. These results suggest that VCP plays a role in WNV infection specifically and inhibition of WNV infection in VCP knockdown cells is not a consequence of cellular cytotoxicity.

421 Employing VLPs, we demonstrated that perturbation of VCP suppressed the infectivity of 422 WNV-VLPs (Fig. 3A). This indicates that VCP is potentially involved in either the early steps or 423 during genome replication of WNV. Using plasmid transfection to generate WNV-VLPs, to bypass 424 early steps of the WNV life cycle, we next examined whether VCP plays a role in the late steps 425 (from genome replication until virus release) of the viral life cycle. Knockdown of VCP reduced the 426 yield of WNV-VLPs compared to control siRNA-treated cells (Fig. 3C) and this finding suggests 427 that VCP also participates in the late steps of WNV life cycle. Taken together, we hypothesized that 428 VCP may be implicated in the genome replication steps of WNV. Therefore, a WNV DNA-based 429 replicon was employed to clarify whether VCP is required for genome replication of WNV. 430 Depletion of VCP significantly decreased expression levels of synthesized WNV-RNA (Fig. 3D) 431 and this finding indicates that VCP is engaged in WNV genomic RNA replication.

It has been previously reported that VCP was found to be localized in the cytosol, ER and 432 433 nucleus, and can play a role in several cellular processes (Arita et al., 2012; Meyer et al., 2012; 434 Meyer and Weihl, 2014; Yamanaka et al., 2012). The possible mechanism(s) of VCP involvement 435 in WNV infection may be based on the localization and physiological function of VCP. Functional 436 roles of VCP in the replication of the WNV-related virus in the family *Flaviviridae*, hepatitis C virus (HCV) have been reported (Yi et al., 2016). VCP knockdown significantly decreased 437 438 expression of HCV RNA levels and VCP was found to be colocalized with the HCV replication 439 complex. It is thus possible that this function of VCP in the HCV life cycle is also required for WNV genome replication, however, no direct evidence currently exists for an interaction between
WNV replicase components and VCP and further investigations are required.

442 Apart from during the genome replication of the WNV life cycle, VCP might potentially be involved in other steps. The present study demonstrates that VCP may also function in the early 443 444 steps, during either attachment or entry, of the viral life cycle (Fig. 3B). The results of an entry 445 assay revealed that silencing of endogenous VCP caused a significant reduction in the expression 446 levels of WNV-RNA compared to control siRNA-treated cells. This suggests that VCP may also 447 play a role in either the binding or entry steps of the WNV life cycle. A role for VCP in early stages 448 of viral infection has previously been reported for coronavirus and Sindbis virus (Panda et al., 2013; 449 Wong et al., 2015). Depletion of VCP inhibited coronavirus infection through a failure in the 450 maturation of virus-loaded endosomes leading to accumulation of coronavirus particles in the early 451 endosomal compartment (Wong et al., 2015). Studies on Sindbis virus indicated that VCP 452 functioned as a regulator of viral entry as knockdown of VCP caused an alteration of trafficking and 453 resulted in the degradation of Sindbis virus entry receptor (Panda et al., 2013). However, the 454 possible function of VCP on early stages of WNV replication has not been investigated and will 455 require further study.

In conclusion, our findings suggest that VCP is required for replication of WNV at a number of different stages of the viral life cycle, thus, VCP potentially represents a candidate for the therapeutic inhibition of WNV infection.

459

460 **Reference**

461 Ambrose, R.L., Mackenzie, J.M., 2011. West Nile virus differentially modulates the unfolded
462 protein response to facilitate replication and immune evasion. J Virol 85(6), 2723-2732.

Arita, M., Wakita, T., Shimizu, H., 2012. Valosin-containing protein (VCP/p97) is required for
poliovirus replication and is involved in cellular protein secretion pathway in poliovirus
infection. J Virol 86(10), 5541-5553.

466	Bogachek, M.V., Zaitsev, B.N., Sekatskii, S.K., Protopopova, E.V., Ternovoi, V.A., Ivanova, A.V.,
467	Kachko, A.V., Ivanisenko, V.A., Dietler, G., Loktev, V.B., 2010. Characterization of
468	glycoprotein E C-end of West Nile virus and evaluation of its interaction force with
469	alphaVbeta3 integrin as putative cellular receptor. Biochemistry (Mosc) 75(4), 472-480.
470	Brinton, M.A., 2014. Replication cycle and molecular biology of the West Nile virus. Viruses 6(1),
471	13-53.
472	Buchan, J.R., Kolaitis, R.M., Taylor, J.P., Parker, R., 2013. Eukaryotic stress granules are cleared
473	by autophagy and Cdc48/VCP function. Cell 153(7), 1461-1474.
474	Bug, M., Meyer, H., 2012. Expanding into new marketsVCP/p97 in endocytosis and autophagy. J
475	Struct Biol 179(2), 78-82.
476	C., S.K., P., H.T., W., B.A., J., P.H., 1940. A Neurotropic Virus Isolated from the Blood of a Native
477	of Uganda. s1-20(20), 471-492.
478	Carnec, X., Meertens, L., Dejarnac, O., Perera-Lecoin, M., Hafirassou, M.L., Kitaura, J., Ramdasi,
479	R., Schwartz, O., Amara, A., 2016. The Phosphatidylserine and Phosphatidylethanolamine
480	Receptor CD300a Binds Dengue Virus and Enhances Infection. J Virol 90(1), 92-102.
481	Chahar, H.S., Chen, S., Manjunath, N., 2013. P-body components LSM1, GW182, DDX3, DDX6
482	and XRN1 are recruited to WNV replication sites and positively regulate viral replication.
483	Virology 436(1), 1-7.
484	Chou, T.F., Deshaies, R.J., 2011. Quantitative cell-based protein degradation assays to identify and
485	classify drugs that target the ubiquitin-proteasome system. J Biol Chem 286(19), 16546-
486	16554.
487	Chu, J.J., Leong, P.W., Ng, M.L., 2006. Analysis of the endocytic pathway mediating the infectious
488	entry of mosquito-borne flavivirus West Nile into Aedes albopictus mosquito (C6/36) cells.
489	Virology 349(2), 463-475.
490	Chu, J.J., Ng, M.L., 2004. Infectious entry of West Nile virus occurs through a clathrin-mediated
491	endocytic pathway. J Virol 78(19), 10543-10555.

- 492 Courtney, S.C., Scherbik, S.V., Stockman, B.M., Brinton, M.A., 2012. West nile virus infections
 493 suppress early viral RNA synthesis and avoid inducing the cell stress granule response. J
 494 Virol 86(7), 3647-3657.
- 495 Dargemont, C., Ossareh-Nazari, B., 2012. Cdc48/p97, a key actor in the interplay between
 496 autophagy and ubiquitin/proteasome catabolic pathways. Biochim Biophys Acta 1823(1),
 497 138-144.
- Dauphin, G., Zientara, S., Zeller, H., Murgue, B., 2004. West Nile: worldwide current situation in
 animals and humans. Comp Immunol Microbiol Infect Dis 27(5), 343-355.
- Davis, C.W., Nguyen, H.Y., Hanna, S.L., Sánchez, M.D., Doms, R.W., Pierson, T.C., 2006. West
 Nile virus discriminates between DC-SIGN and DC-SIGNR for cellular attachment and
 infection. J Virol 80(3), 1290-1301.
- 503 Denizot, M., Neal, J.W., Gasque, P., 2012. Encephalitis due to emerging viruses: CNS innate
 504 immunity and potential therapeutic targets. J Infect 65(1), 1-16.
- 505 Egberink, H., Addie, D.D., Boucraut-Baralon, C., Frymus, T., Gruffydd-Jones, T., Hartmann, K.,

506 Horzinek, M.C., Hosie, M.J., Marsilio, F., Lloret, A., Lutz, H., Pennisi, M.G., Radford,

- 507A.D., Thiry, E., Truyen, U., Möstl, K., Diseases, E.A.B.o.C., 2015. West Nile virus
- infection in cats: ABCD guidelines on prevention and management. J Feline Med Surg17(7), 617-619.
- 510 Fernandez-Garcia, M.D., Meertens, L., Bonazzi, M., Cossart, P., Arenzana-Seisdedos, F., Amara,
- A., 2011. Appraising the roles of CBLL1 and the ubiquitin/proteasome system for flavivirus
 entry and replication. J Virol 85(6), 2980-2989.
- 513 Fields, B.N., Knipe, D.M., Howley, P.M., 2013. Fields virology. 6th ed. 2 vols. Wolters Kluwer
 514 Health/Lippincott Williams & Wilkins, Philadelphia.
- 515 Gamino, V., Escribano-Romero, E., Blázquez, A.B., Gutiérrez-Guzmán, A.V., Martín-Acebes, M.,
- 516 Saiz, J.C., Höfle, U., 2016. Experimental North American West Nile Virus Infection in the
- 517 Red-legged Partridge (Alectoris rufa). Vet Pathol 53(3), 585-593.

518	Gilfoy, F., Fayzulin, R., Mason, P.W., 2009. West Nile virus genome amplification requires the
519	functional activities of the proteasome. Virology 385(1), 74-84.
520	Gillespie, L.K., Hoenen, A., Morgan, G., Mackenzie, J.M., 2010. The endoplasmic reticulum
521	provides the membrane platform for biogenesis of the flavivirus replication complex. J Virol
522	84(20), 10438-10447.
523	Hasebe, R., Suzuki, T., Makino, Y., Igarashi, M., Yamanouchi, S., Maeda, A., Horiuchi, M., Sawa,
524	H., Kimura, T., 2010. Transcellular transport of West Nile virus-like particles across human
525	endothelial cells depends on residues 156 and 159 of envelope protein. BMC Microbiol 10,
526	165.
527	Heinz, F.X., Allison, S.L., 2000. Structures and mechanisms in flavivirus fusion. Adv Virus Res 55,
528	231-269.
529	Jiang, D., Weidner, J.M., Qing, M., Pan, X.B., Guo, H., Xu, C., Zhang, X., Birk, A., Chang, J., Shi,
530	P.Y., Block, T.M., Guo, J.T., 2010. Identification of five interferon-induced cellular proteins
531	that inhibit west nile virus and dengue virus infections. J Virol 84(16), 8332-8341.
532	Kaufmann, B., Rossmann, M.G., 2011. Molecular mechanisms involved in the early steps of
533	flavivirus cell entry. Microbes Infect 13(1), 1-9.
534	Kaufusi, P.H., Kelley, J.F., Yanagihara, R., Nerurkar, V.R., 2014. Induction of endoplasmic
535	reticulum-derived replication-competent membrane structures by West Nile virus non-
536	structural protein 4B. PLoS One 9(1), e84040.
537	Kimura, T., Kimura-Kuroda, J., Nagashima, K., Yasui, K., 1994. Analysis of virus-cell binding
538	characteristics on the determination of Japanese encephalitis virus susceptibility. Arch Virol
539	139(3-4), 239-251.
540	Kobayashi, S., Orba, Y., Yamaguchi, H., Kimura, T., Sawa, H., 2012. Accumulation of
541	ubiquitinated proteins is related to West Nile virus-induced neuronal apoptosis.
542	Neuropathology 32(4), 398-405.

- Kobayashi, S., Orba, Y., Yamaguchi, H., Takahashi, K., Sasaki, M., Hasebe, R., Kimura, T., Sawa,
 H., 2014. Autophagy inhibits viral genome replication and gene expression stages in West
 Nile virus infection. Virus Res 191, 83-91.
- 546 Kobayashi, S., Suzuki, T., Igarashi, M., Orba, Y., Ohtake, N., Nagakawa, K., Niikura, K., Kimura,
- T., Kasamatsu, H., Sawa, H., 2013. Cysteine residues in the major capsid protein, Vp1, of
 the JC virus are important for protein stability and oligomer formation. PLoS One 8(10),
 e76668.
- Kobayashi, S., Suzuki, T., Kawaguchi, A., Phongphaew, W., Yoshii, K., Iwano, T., Harada, A.,
 Kariwa, H., Orba, Y., Sawa, H., 2016a. Rab8b Regulates Transport of West Nile Virus
 Particles from Recycling Endosomes. J Biol Chem 291(12), 6559-6568.
- Kobayashi, S., Yoshii, K., Hirano, M., Muto, M., Kariwa, H., 2016b. A Novel Reverse Genetics
 System for Production of Infectious West Nile Virus using Homologous Recombination in
 Mammalian Cells. J Virol Methods.
- Koller, K.J., Brownstein, M.J., 1987. Use of a cDNA clone to identify a supposed precursor protein
 containing valosin. Nature 325(6104), 542-545.
- Kramer, L.D., Bernard, K.A., 2001. West Nile virus infection in birds and mammals. Ann N Y
 Acad Sci 951, 84-93.
- 560 Krishnan, M.N., Ng, A., Sukumaran, B., Gilfoy, F.D., Uchil, P.D., Sultana, H., Brass, A.L.,
- 561 Adametz, R., Tsui, M., Qian, F., Montgomery, R.R., Lev, S., Mason, P.W., Koski, R.A.,
- 562 Elledge, S.J., Xavier, R.J., Agaisse, H., Fikrig, E., 2008. RNA interference screen for human
 563 genes associated with West Nile virus infection. Nature 455(7210), 242-245.
- Lichtensteiger, C.A., Heinz-Taheny, K., Osborne, T.S., Novak, R.J., Lewis, B.A., Firth, M.L., 2003.
- 565 West Nile virus encephalitis and myocarditis in wolf and dog. Emerg Infect Dis 9(10),
- 566 1303-1306.

567	Ma, H., Dang, Y., Wu, Y., Jia, G., Anaya, E., Zhang, J., Abraham, S., Choi, J.G., Shi, G., Qi, L.,
568	Manjunath, N., Wu, H., 2015. A CRISPR-Based Screen Identifies Genes Essential for West-
569	Nile-Virus-Induced Cell Death. Cell Rep 12(4), 673-683.
570	Makino, Y., Suzuki, T., Hasebe, R., Kimura, T., Maeda, A., Takahashi, H., Sawa, H., 2014.
571	Establishment of tracking system for West Nile virus entry and evidence of microtubule
572	involvement in particle transport. J Virol Methods 195, 250-257.
573	Maric, M., Maculins, T., De Piccoli, G., Labib, K., 2014. Cdc48 and a ubiquitin ligase drive
574	disassembly of the CMG helicase at the end of DNA replication. Science 346(6208),
575	1253596.
576	Martina, B.E., Koraka, P., van den Doel, P., Rimmelzwaan, G.F., Haagmans, B.L., Osterhaus, A.D.,
577	2008. DC-SIGN enhances infection of cells with glycosylated West Nile virus in vitro and
578	virus replication in human dendritic cells induces production of IFN-alpha and TNF-alpha.
579	Virus Res 135(1), 64-71.
580	Meyer, H., Bug, M., Bremer, S., 2012. Emerging functions of the VCP/p97 AAA-ATPase in the
581	ubiquitin system. Nat Cell Biol 14(2), 117-123.
582	Meyer, H., Weihl, C.C., 2014. The VCP/p97 system at a glance: connecting cellular function to
583	disease pathogenesis. J Cell Sci 127(Pt 18), 3877-3883.
584	Morizono, K., Chen, I.S., 2014. Role of phosphatidylserine receptors in enveloped virus infection. J
585	Virol 88(8), 4275-4290.
586	Panda, D., Rose, P.P., Hanna, S.L., Gold, B., Hopkins, K.C., Lyde, R.B., Marks, M.S., Cherry, S.,
587	2013. Genome-wide RNAi screen identifies SEC61A and VCP as conserved regulators of
588	Sindbis virus entry. Cell Rep 5(6), 1737-1748.
589	Paz, S., 2015. Climate change impacts on West Nile virus transmission in a global context. Philos
590	Trans R Soc Lond B Biol Sci 370(1665).
591	Perera-Lecoin, M., Meertens, L., Carnec, X., Amara, A., 2014. Flavivirus entry receptors: an
592	update. Viruses 6(1), 69-88.

- Plevka, P., Battisti, A.J., Sheng, J., Rossmann, M.G., 2014. Mechanism for maturation-related
 reorganization of flavivirus glycoproteins. J Struct Biol 185(1), 27-31.
- 595 Pye, V.E., Dreveny, I., Briggs, L.C., Sands, C., Beuron, F., Zhang, X., Freemont, P.S., 2006. Going
 596 through the motions: the ATPase cycle of p97. J Struct Biol 156(1), 12-28.
- Ramanathan, H.N., Ye, Y., 2012. The p97 ATPase associates with EEA1 to regulate the size of
 early endosomes. Cell Res 22(2), 346-359.
- Read, R.W., Rodriguez, D.B., Summers, B.A., 2005. West Nile virus encephalitis in a dog. Vet
 Pathol 42(2), 219-222.
- 601 Ritz, D., Vuk, M., Kirchner, P., Bug, M., Schütz, S., Hayer, A., Bremer, S., Lusk, C., Baloh, R.H.,
- 602 Lee, H., Glatter, T., Gstaiger, M., Aebersold, R., Weihl, C.C., Meyer, H., 2011.
- Endolysosomal sorting of ubiquitylated caveolin-1 is regulated by VCP and UBXD1 and
 impaired by VCP disease mutations. Nat Cell Biol 13(9), 1116-1123.
- Roby, J.A., Setoh, Y.X., Hall, R.A., Khromykh, A.A., 2015. Post-translational regulation and
 modifications of flavivirus structural proteins. J Gen Virol 96(Pt 7), 1551-1569.
- Samuel, M.A., Diamond, M.S., 2006. Pathogenesis of West Nile Virus infection: a balance between
 virulence, innate and adaptive immunity, and viral evasion. J Virol 80(19), 9349-9360.
- 609 Scholle, F., Girard, Y.A., Zhao, Q., Higgs, S., Mason, P.W., 2004. trans-Packaged West Nile virus-
- 610 like particles: infectious properties in vitro and in infected mosquito vectors. J Virol 78(21),
 611 11605-11614.
- 612 Seguin, S.J., Morelli, F.F., Vinet, J., Amore, D., De Biasi, S., Poletti, A., Rubinsztein, D.C., Carra,
- 613 S., 2014. Inhibition of autophagy, lysosome and VCP function impairs stress granule
 614 assembly. Cell Death Differ 21(12), 1838-1851.
- Shimojima, M., Takenouchi, A., Shimoda, H., Kimura, N., Maeda, K., 2014. Distinct usage of three
 C-type lectins by Japanese encephalitis virus: DC-SIGN, DC-SIGNR, and LSECtin. Arch
 Virol 159(8), 2023-2031.

- Shirato, K., Kimura, T., Mizutani, T., Kariwa, H., Takashima, I., 2004a. Different chemokine
 expression in lethal and non-lethal murine West Nile virus infection. J Med Virol 74(3),
 507-513.
- 621 Shirato, K., Miyoshi, H., Goto, A., Ako, Y., Ueki, T., Kariwa, H., Takashima, I., 2004b. Viral
- envelope protein glycosylation is a molecular determinant of the neuroinvasiveness of the
 New York strain of West Nile virus. J Gen Virol 85(Pt 12), 3637-3645.
- Smit, J.M., Moesker, B., Rodenhuis-Zybert, I., Wilschut, J., 2011. Flavivirus cell entry and
 membrane fusion. Viruses 3(2), 160-171.
- Stolz, A., Hilt, W., Buchberger, A., Wolf, D.H., 2011. Cdc48: a power machine in protein
 degradation. Trends Biochem Sci 36(10), 515-523.
- Suthar, M.S., Diamond, M.S., Gale, M., 2013. West Nile virus infection and immunity. Nat Rev
 Microbiol 11(2), 115-128.
- Takada, A., Ebihara, H., Feldmann, H., Geisbert, T.W., Kawaoka, Y., 2007. Epitopes required for
 antibody-dependent enhancement of Ebola virus infection. J Infect Dis 196 Suppl 2, S347356.
- 633 Takahashi, H., Ohtaki, N., Maeda-Sato, M., Tanaka, M., Tanaka, K., Sawa, H., Ishikawa, T.,
- 634 Takamizawa, A., Takasaki, T., Hasegawa, H., Sata, T., Hall, W.W., Kurata, T., Kojima, A.,
- 635 2009. Effects of the number of amino acid residues in the signal segment upstream or
- downstream of the NS2B-3 cleavage site on production and secretion of prM/M-E virus-like
- 637 particles of West Nile virus. Microbes Infect 11(13), 1019-1028.
- Troupin, A., Colpitts, T.M., 2016. Overview of West Nile Virus Transmission and Epidemiology.
 Methods Mol Biol 1435, 15-18.
- 640 Verma, R., Oania, R., Fang, R., Smith, G.T., Deshaies, R.J., 2011. Cdc48/p97 mediates UV641 dependent turnover of RNA Pol II. Mol Cell 41(1), 82-92.
- Wang, Q., Li, L., Ye, Y., 2008. Inhibition of p97-dependent protein degradation by Eeyarestatin I. J
 Biol Chem 283(12), 7445-7454.

644	Wang, Q., Shinkre, B.A., Lee, J.G., Weniger, M.A., Liu, Y., Chen, W., Wiestner, A., Trenkle,
645	W.C., Ye, Y., 2010. The ERAD inhibitor Eeyarestatin I is a bifunctional compound with a
646	membrane-binding domain and a p97/VCP inhibitory group. PLoS One 5(11), e15479.
647	Welsch, S., Miller, S., Romero-Brey, I., Merz, A., Bleck, C.K., Walther, P., Fuller, S.D., Antony,
648	C., Krijnse-Locker, J., Bartenschlager, R., 2009. Composition and three-dimensional
649	architecture of the dengue virus replication and assembly sites. Cell Host Microbe 5(4), 365-
650	375.
651	Wilcox, A.J., Laney, J.D., 2009. A ubiquitin-selective AAA-ATPase mediates transcriptional
652	switching by remodelling a repressor-promoter DNA complex. Nat Cell Biol 11(12), 1481-
653	1486.
654	Wolf, D.H., Stolz, A., 2012. The Cdc48 machine in endoplasmic reticulum associated protein
655	degradation. Biochim Biophys Acta 1823(1), 117-124.
656	Wong, H.H., Kumar, P., Tay, F.P., Moreau, D., Liu, D.X., Bard, F., 2015. Genome-Wide Screen
657	Reveals Valosin-Containing Protein Requirement for Coronavirus Exit from Endosomes. J
658	Virol 89(21), 11116-11128.
659	Xia, D., Tang, W.K., Ye, Y., 2016. Structure and function of the AAA+ ATPase p97/Cdc48p. Gene
660	583(1), 64-77.
661	Yamanaka, K., Sasagawa, Y., Ogura, T., 2012. Recent advances in p97/VCP/Cdc48 cellular
662	functions. Biochim Biophys Acta 1823(1), 130-137.
663	Yi, Z., Fang, C., Zou, J., Xu, J., Song, W., Du, X., Pan, T., Lu, H., Yuan, Z., 2016. Affinity
664	Purification of the Hepatitis C Virus Replicase Identifies Valosin-Containing Protein, a
665	Member of the ATPases Associated with Diverse Cellular Activities Family, as an Active
666	Virus Replication Modulator. J Virol 90(21), 9953-9966.
667	Yu, I.M., Zhang, W., Holdaway, H.A., Li, L., Kostyuchenko, V.A., Chipman, P.R., Kuhn, R.J.,
668	Rossmann, M.G., Chen, J., 2008. Structure of the immature dengue virus at low pH primes
669	proteolytic maturation. Science 319(5871), 1834-1837.

670	Zaitsev, B.N., Benedetti, F., Mikhaylov, A.G., Korneev, D.V., Sekatskii, S.K., Karakouz, T.,
671	Belavin, P.A., Netesova, N.A., Protopopova, E.V., Konovalova, S.N., Dietler, G., Loktev,
672	V.B., 2014. Force-induced globule-coil transition in laminin binding protein and its role for
673	viral-cell membrane fusion. J Mol Recognit 27(12), 727-738.
674	Zhong, X., Pittman, R.N., 2006. Ataxin-3 binds VCP/p97 and regulates retrotranslocation of ERAD
675	substrates. Hum Mol Genet 15(16), 2409-2420.
676	Zidane, N., Ould-Abeih, M.B., Petit-Topin, I., Bedouelle, H., 2013. The folded and disordered
677	domains of human ribosomal protein SA have both idiosyncratic and shared functions as
678	membrane receptors. Biosci Rep 33(1), 113-124.
679	Centers for Disease Control and Prevention (CDC), 2016.West Nile virus disease cases and deaths
680	reported to CDC by year and clinical presentation, 1999-2015. (accessed 18.08.16).
681	
682	Acknowledgements
683	This work was supported by the Program for Leading Graduate Schools "Fostering Global Leaders
684	in Veterinary Science for Contributing to One Health", and grants (16H06429, 16H06431) from the
685	Ministry of Education, Culture, Sports, Science and Technology (MEXT), Japan. We thank Dr.
686	Takashima for donating the WNV NY99 6-LP strain and Dr. Takada for providing pseudotyped
687	VSV.
688	
689	Author contributions statement
690	P. Wallaya, S.K., M.S., Y.O and H.S. conceived the experiments, P. Wallaya and S.K. conducted
691	the experiments and analysed the data. P. Wallaya, S.K., M.S., Y.O., and H.S. contributed
692	reagents/materials/analysis tools. P. Wallaya, S.K., M.C., W.H., Y.O. and H.S. wrote the paper. All
693	authors reviewed the manuscript.
694	

695 Additional information

696 **Competing financial interests:** The authors declare no competing financial interests.

697

698 Figure Legends

699 **Fig. 1.**

WNV infection is inhibited in the presence of VCP inhibitors. (A) WNV infection in the presence
of either EerI (left panels) or MDBN (right panels). HeLa cells were inoculated with WNV

702 (MOI=1) and then treated with EerI or MDBN at 1 h.p.i. Cells were harvested at 24 h.p.i. and

stained with anti-JEV antibody (Kimura et al., 1994; Kobayashi et al., 2012) that has cross

reactivity with WNV antigen (green). Cell nuclei were counterstained with DAPI (blue). (B)

Positivity of WNV-infected cells from (A). Mean \pm SD from triplicate experiments is shown; * p <

706 0.05, ** p < 0.01 (one-way ANOVA). (C) The culture supernatants from (A) were collected at 24

h.p.i. and the viral titers of the harvested supernatants were examined by plaque assay. Mean \pm SD

from three independent experiments is shown; * p < 0.05, ** p < 0.01 (one-way ANOVA).

709

710 Fig. 2.

711 WNV infection is inhibited in VCP knockdown cells. (A) HeLa cells were treated with either 712 siRNA against VCP [siVCP (1), (2) and (3)] or control siRNA (siCont). The siRNA-treated cells 713 were inoculated with WNV (MOI=1) at 48 h.p.i. The inoculated cells were harvested at 24 h.p.i. 714 The expression of endogenous VCP protein and WNV envelope protein after treatment with the 715 indicated siRNA were examined by immunoblotting with mouse anti-VCP antibody and mouse 716 anti-WNV/Kunjin envelope protein. The expression of actin was examined after reprobing as an 717 endogenous control. (B) WNV-infected cells from (A), after 24 h incubation with WNV, the cells 718 were harvested and examined by immunofluorescence assay. WNV-infected cells were stained with 719 anti-JEV antibody (green) and cell nuclei were counterstained with DAPI (blue). (C) Positivity of WNV-infected cells from (B). Mean \pm SD from three independent experiments is shown; ** p <720 721 0.01 (one-way ANOVA). (D) The culture supernatants from (A) were collected at 24 h.p.i and the

viral titers of the harvested supernatants were determined using plaque assay. Mean \pm SD from three independent experiments is shown; ** p < 0.01 (one-way ANOVA).

724

725 **Fig. 3**.

726 The role of VCP in distinct steps of WNV life cycle was investigated. (A) WNV-VLP infection 727 after the indicated siRNA treatment, HeLa cells were treated with siRNA against VCP, siVCP (1) 728 or control siRNA (siCont). The indicated siRNA treated-cells were inoculated with WNV-VLPs (16 729 HAU) at 48 h post transfection and incubated for 72 h. Relative quantification of WNV-RNA 730 normalized to human β -actin from (A) examined by qRT-PCR. Mean \pm SD from three independent 731 experiments is shown; ** p < 0.01 (one-way ANOVA). (B) Relative quantification of WNV-RNA 732 expression levels normalized to human β -actin of siRNA-treated cells inoculated with WNV at the early time point of infection. siRNA-treated HeLa cells were inoculated with WNV after 48 h post 733 734 siRNA transfection. The inoculated cells were incubated on ice for 1 h, followed by washing 5 735 times with PBS and then transferred to 37 °C. After 1 h incubation, the cells were harvested by 736 trypsin and prepared for qRT-PCR. Mean \pm SD from two independent experiments in triplicate is shown; * p < 0.05 (one-way ANOVA). (C) HeLa cells were treated with siRNA against VCP, 737 738 siVCP (1) or control siRNA (siCont). After 24 h post transfection, the cells were transfected with 739 plasmid set for WNV-VLP production and incubated for 72 h. The culture supernatants were 740 harvested and inoculated on Vero cell monolayers in 10-fold serial dilutions. The viral titers of the 741 harvested supernatants were determined as IFU/ml. Mean \pm SD from three independent experiments 742 is shown; ** p < 0.01 (one-way ANOVA). (D) HeLa cells were treated with siRNA, either siVCP 743 (1) or siCont, and then transfected with plasmid containing WNV DNA replicon, pCMV-WNrep-744 DsRed at 24 h post siRNA treatment and incubated. The transfected cells were harvested at 72 h 745 post transfection of pCMV-WNrep-DsRed. Relative quantification of WNV-RNA normalized to 746 expression of human β -actin was examined by qRT-PCR. Mean \pm SD from three independent experiments is shown; ** p < 0.01 (one-way ANOVA). 747

Figure 1



Figure 2



B

С

Positivity of WNV-infected

0

siCont

(1)







(2)

siVCP

(3)



Figure 3







D



Supplementary Fig. 1.

Cell viability of HeLa cells after treatment with either EerI or MDBN. The HeLa cells were treated with the indicated concentrations of either EerI (left) or MDBN (right). Cell viability was examined by a MTT assay at 24 h post treatment.

A



Supplementary Fig. 2.

WNV infection in SK-N-SH cells is attenuated in the presence of EerI. (A) WNV infection in human neuroblastoma, SK-N-SH cells. SK-N-SH cells were inoculated with WNV (MOI=1). After 1 h.p.i., the inoculated-cells were treated with EerI at the indicated concentration. DMSO treated-cells were used as control. The cells were fixed and prepared for IFA at 24 h.p.i and stained with anti-JEV antibody which has cross reactivity with WNV antigen (green). Cell nuclei were stained with DAPI (blue). (B) Positivity of WNV-infected cells from (A). Mean \pm SD from triplicate experiments is shown; * p < 0.05. ** p < 0.01 (one-way ANOVA). (C) Cell viability of SK-N-SH at 24 h after exposure to EerI with absorbance values obtained after MTT assay.



Supplementary Fig. 3

Representative figure showing IFA results of WNV infection in siVCP (1) compared to siCont treated-HeLa cells at 12, 24 and 48 h.p.i. The HeLa cells were treated with either siVCP or siCont and then incubated at 37 °C for 48 h. Thereafter, the cells were infected with WNV (MOI=1) and incubated at 37 °C for 12, 24 and 48 h.p.i., respectively. After incubation at the indicated time points, the cells were then fixed and prepared for IFA followed by staining with anti-JEV antibody that has cross reactivity with WNV antigen (green). Cell nuclei were stained with DAPI (blue).



Supplementary Fig. 4.

VCP knockdown does not affect infection of pseudotyped VSV. (A) HeLa cells were treated with siRNA against VCP (siVCP) or control siRNA (siCont) for 48 h. Thereafter, the siRNA-treated cells were inoculated with pseudotyped VSV. Pseudotyped VSV positive cells were determined by expression of green fluorescence protein (GFP), which was carried by the pseudotyped VSV replicon, at 8 h.p.i. nuclei were stained with DAPI (blue). (B) The positivity of GFP from (A). Mean \pm SD from triplicate experiments is shown. The significance was analyzed using a one-way ANOVA.

Highlights

- Inhibition of VCP by chemical inhibitors decreased WNV infection in a dosedependent manner.
- Knockdown of endogenous VCP level using siRNA suppressed WNV infection.
- Depletion of VCP levels suppressed WNV infection at the early stages of WNV replication cycle.
- Depletion of VCP levels lowered nascent WNV genomic RNA.
- VCP participates in early stages and viral genomic RNA replication.