

Display of the human (pro)renin receptor on
Bombyx mori nucleopolyhedrovirus (BmNPV)
particles using Bm cells

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1 **Display of the human (pro)renin receptor on *Bombyx mori***
2 **nucleopolyhedrovirus (BmNPV) particles using Bm cells**

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23 **Abstract**

24

25 The human prorenin receptor (hPRR) was displayed on the surface of *Bombyx mori*
26 nucleopolyhedrovirus (BmNPV) with and without fusion to glycoprotein 64 (GP64) of the
27 BmNPV. hPRR1 is a native hPRR with an additional FLAG peptide sequence inserted
28 between the signal peptide and prorenin-binding domain. hPRR2 has the prorenin-binding
29 domain inserted between amino acid residues ⁸¹Asp and ⁸²Pro of GP64. hPRR4 has the
30 prorenin-binding domain inserted in ⁸¹Asp and ³²⁰Met of partially deleted GP64.
31 Incorporation of hPRR was confirmed in recombinant BmNPV (rBmNPV) but not in
32 cysteine protease-deleted rBmNPV. hPRR1 was observed in ER, but hPRR2 and hPRR4
33 were observed around the endoplasmic reticulum (ER) and in its periphery. rBmNPV-hPRR1
34 and -hPRR2, carrying hPRR1 and hPRR2 respectively, showed binding affinity to human
35 renin, but rBmNPV-hPRR4 did not. The presence of hPRR4 of rBmNPV-hPRR4 was
36 confirmed in western blotting under nonreducing conditions, suggesting that although hPRR4
37 was incorporated in rBmNPV-hPRR4, it behaved as a nonfunctional aggregate. This
38 rBmNPV display system can also be used for analyzing a ligand-receptor interaction.

39

40 **Keywords:** BmNPV; surface display; prorenin receptor; GP64; fusion protein

41

42 Baculoviruses are widely used as expression vectors of recombinant proteins in insect
43 cells and larvae. They permit high-level expression of recombinant proteins that are co- and
44 post-translationally modified. Baculoviruses can also be used to display recombinant proteins
45 on their surfaces (1, 2). A common method for displaying proteins on the surface of
46 baculoviruses involves the use of the transmembrane domain of GP64 present in the
47 baculovirus. The GP64 transmembrane domain can be substituted with that of hemagglutinin
48 (HA) or vesicular stomatitis virus G glycoprotein (VSVG) (3, 4). The N-terminal domain of
49 neuraminidase from influenza virus has also been used as the transmembrane domain (5).
50 To date, *Autographa californica* multiple nucleopolyhedrovirus (AcMNPV) that can infect
51 *Spodoptera frugiperda*, *Trichoplusia ni*, and cabbage loopers, has been the most widely used
52 baculovirus for baculovirus surface display. The display system using *Bombyx mori*
53 nucleopolyhedrovirus (BmNPV) has been employed, but it is in a few (6, 7). BmNPV is a
54 pathogen of the silkworm *B. mori*, and both AcMNPV and BmNPV belong to the family
55 Baculoviridae. BmNPV is also used for recombinant protein production, especially when
56 insect larvae, silkworms, are used as hosts. Silkworms are bigger than and superior to
57 cabbage loopers for recombinant protein production. The baculovirus titer in the hemolymph
58 of BmNPV-infected silkworm larvae is higher than that in the BmNPV-infected Bm5 cell
59 culture supernatant. When insect larvae are used, sterile conditions and culture medium,
60 including serum and growth factors, are not required during the protein production process.
61 Therefore, the insect larvae are advantageous to mass production of virus titer and
62 recombinant protein.

63 In this paper, the human (pro)renin receptor (hPRR) was employed to display on the
64 surface of BmNPV in the culture of *B. mori*-derived Bm5 cells. Three types of constructs

65 were used for display of hPRR. One was the native form of hPRR. Others were fused to
66 GP64 of BmNPV; the prorenin-binding domain of hPRR was either inserted between ⁸¹Asp
67 and ⁸²Pro of GP64 or between ⁸¹Asp and ³²⁰Met of partially deleted GP64. These three
68 constructs were investigated for insect cell localization and binding for prorenin. The
69 BmNPV display system can be used for the large-scale production of recombinant proteins
70 displayed on the surface of the baculovirus. Infection of silkworm larvae with such
71 baculoviruses that display various proteins may then be carried out and used for the
72 functional analysis of receptors, drug delivery system, and vaccines against infectious viruses
73 and protozoa.

74

75 MATERIALS AND METHODS

76

77 **Cell line, cell culture, and baculovirus infection** Bm5 cells derived from *Bombyx mori*
78 were cultured in NIM-Ex medium (Nihon Nosan Co. Ltd., Yokohama, Japan) supplemented
79 with 1% heat-inactivated fetal bovine serum (Sigma-Aldrich, St. Louis, MO, USA) and 1%
80 antibiotic-antimycotic solution (Invitrogen, Carlsbad, CA, USA). Bm5 cells were infected
81 with 10 µl hemolymph from recombinant bacmid-injected silkworm larvae. The multiplicity
82 of infection (M.O.I.) was 0.1–1.

83 **Construction of hPRRs and bacmids and preparation of hemolymph from silkworm**

84 **larvae** Three hPRRs were constructed as shown in Fig. 1. hPRR1 is the native form and has
85 an additional FLAG peptide sequence behind the signal peptide sequence, as reported
86 previously (8). To fuse hPRR with *gp64* from BmNPV, the *gp64* gene was amplified by PCR
87 with the Bm*gp64*-F and Bm*gp64*-R primers (Table 1). The amplified *gp64* gene product was

88 inserted in the *Kpn I–Hind III* site in pUC18, and the resulting plasmid was named pBmgp64.
89 hPRR2 gene, in which the prorenin-binding domain of hPRR is inserted between the N-
90 terminal domain between ⁸¹Asp and ⁸²Pro of the complete gp64 gene was constructed. The
91 hPRR gene was amplified with the bgl-hPRR-F and bgl-hPRR-R primers (Table 1), and the
92 amplified hPRR gene was inserted into the *BamH I* site in pBmgp64. hPRR4 gene, in which
93 the prorenin-binding domain of hPRR is inserted between ⁸¹Asp and ³²⁰Met of partially
94 deleted gp64, was constructed. In hPRR4, the prorenin-binding domain of hPRR was put
95 between N-terminal (amino acid 1-81) and C-terminal (amino acid 320-511) domains of
96 GP64. This is for investigation of the deletion effect of internal domain of GP64 (amino acid
97 82-319) in hPRR4. The hPRR PCR product was inserted into the *BamH I–Sph I* site in
98 pBmgp64. The hPRR1, hPRR2, and hPRR4 genes were amplified with the Bmgp64-F and
99 Bmgp64-R primers (Table 1), and the three amplified hPRR genes were inserted into
100 pENTR/D/TOPO by the TOPO cloning method. Recombinant bacmids were constructed
101 using *Escherichia coli* BmDH10Bac (9) and BmDH10Bac (BmNPV-CP⁻) that lacks its own
102 cysteine protease (10). Hemolymph containing recombinant BmNPV (rBmNPV) was
103 prepared using constructed recombinant bacmids according to a previous study (9, 10).

104 **Baculovirus isolation** Bm5 cells were infected with each hemolymph containing
105 rBmNPV and cultivated for 1–2 days. The culture media were centrifuged at 8000 × *g* for 10
106 min, and the culture supernatants were recovered and overlaid on a 25% sucrose solution
107 (25% sucrose in 5 mM NaCl and 10 mM EDTA). These were further centrifuged at 114000 ×
108 *g* for 1 h. The supernatants were removed, and the pellets were washed with phosphate-
109 buffered saline (PBS, pH 6.2). The pellets were resuspended in small volumes of PBS and
110 used for further experiments.

111 **Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and**
112 **western blotting** Proteins were separated by SDS-PAGE using 12% polyacrylamide or 5–
113 15% gradient polyacrylamide gels that were subsequently subjected to western blotting. For
114 electrophoresis under nonreducing conditions, the samples were mixed with sample buffer
115 without β -mercaptoethanol and boiled. After SDS-PAGE, the proteins were blotted onto a
116 polyvinylidene fluoride (PVDF) membrane using the Mini Trans-Blot Electrophoretic
117 Transfer Cell (Bio-Rad, Hercules, CA, USA). After blocking in 5% skim milk in Tris-
118 buffered saline containing 0.1% Tween 20 (TBST), the membrane was incubated for 1 h in
119 either 1: 10000 diluted mouse anti-FLAG M2 antibody (Sigma-Aldrich, St. Louis, MO,
120 USA) or 1: 4000 diluted rabbit anti-Bmcp64 polyclonal antibody (BioGate Co., Ltd., Gifu,
121 Japan). The membrane was washed with TBST and then incubated for 1 h in 1: 20000 diluted
122 anti-mouse or anti-rabbit IgG antibody labeled with horseradish peroxidase (GE Healthcare
123 UK Ltd., Buckinghamshire, England). Detection was carried out using the ECL Plus Western
124 blotting reagent (GE Healthcare UK Ltd.). Specific bands were detected on a Fluor-S MAX
125 MultiImager (Bio-Rad).

126 **Immunofluorescence microscopy** Bm5 cells were infected with hemolymph and
127 cultivated for 2 days. The culture medium was sampled, and infected cells were attached to
128 polylysine-coated slide glass. These cells were fixed with 8% formalin solution (Wako Pure
129 Chem. Ind. Ltd., Osaka, Japan) for 20 min and washed with PBS (pH 7.0). Any remaining
130 aldehyde was quenched with 50 mM NH_4Cl in PBS. The fixed cells were washed with PBS
131 and blocked overnight with 8% (w/v) bovine serum albumin (BSA) in PBS. After blocking
132 with BSA, the cells were incubated for 2–3 h with 1: 1000 dilution of mouse anti-FLAG M2
133 antibody in PBS and 2% BSA. The cells were washed with PBS and incubated for 1 h with 1:

134 100 dilution of fluorescein isothiocyanate (FITC)-conjugated goat anti-mouse antibody
135 (Jackson ImmunoResearch Laboratories Inc., West Grove, PA, USA) in PBS and 2% BSA.
136 The endoplasmic reticulum (ER) was stained with 1 μ M ER-TrackerTM Red (Molecular
137 Probes, Carlsbad, CA, USA) at 4°C for 30 min. After staining, the cells were examined by
138 confocal laser scanning microscopy (TCS-LS, Leica Microsystems, Heidelberg, Germany).

139 **Surface plasmon resonance (SPR) experiments** The surface plasmon resonance
140 experiments were performed in an SPR optical biosensor (IASys plus; IASYS Affinity
141 Sensors Ltd., Saxon Hill, Cambridge, UK) using a carboxymethyl (CM) sensor cuvette.
142 Renin (2.5 μ g; Cayman, Ann Arbor, MI, USA) was immobilized on the surface of the CM
143 sensor cuvette, and unreacted CM groups on the sensor cuvette, which did not have any
144 immobilized protein, were blocked with ethanolamine to control nonspecific binding. PBS
145 (pH 6.2) was used as the running buffer.

146

147 **Results**

148

149 **Expression of hPRRs in Bm5 cells using rBmNPV and rBmNPV-CP⁻** Production of
150 the GFP_{uv}- β 1,3-*N*-acetylglucosaminyltransferase2 fusion protein (GGT2) was significantly
151 improved when the rBmNPV-CP⁻ bacmid was used because GGT2 degradation by the
152 cysteine protease was reduced (10). Therefore, to express hPRRs on rBmNPV surfaces,
153 rBmNPV-CP⁻ was used. However, there are no reports on display on the surface of BmNPV-
154 CP⁻. Three bacmids rBmNPV-CP⁻-hPRR1, -hPRR2, and -hPRR4 were constructed, and
155 hemolymph from each bacmid-injected silkworm larvae was prepared. Bm5 cells were
156 infected with each rBmNPV-CP⁻, and the cell cultures were sampled at a post-infection time

157 (P.I.T.) of 2 days. During this period, the viability of Bm5 cells was 95%–97%. The Bm5
158 cells were separated from the cultures, and the intracellular expression of each hPRR in Bm5
159 cells was confirmed by western blotting (Data not shown). The estimated molecular weights
160 of the hPRRs in the Bm5 cells are as follows: hPRR1, 38.4 kDa; hPRR2, 95.8 kDa; and
161 hPRR4, 64.7 kDa. The culture supernatant was overlaid on a 25% sucrose solution, and each
162 rBmNPV-CP⁺ was recovered by centrifugation. The expression of each hPRR in each
163 rBmNPV-CP⁺ was confirmed by western blotting. GP64 was observed in all samples, but the
164 hPRR was not detected (Fig. 2A).

165 Three types of rBmNPVs (-hPRR1, -hPRR2, and -hPRR4) were prepared by silkworm
166 expression system, and used for infection to Bm5 cells. Each hPRR was expressed
167 intracellularly (Data not shown). The presence of each hPRR in each rBmNPV was examined
168 (Fig. 2B) and compared against that in rBmNPV-CP⁺. In particular, hPRR1 seemed to be
169 incorporated in rBmNPV-hPRR1 with its own native single transmembrane domain (Fig. 2B,
170 left, lane 1). Recently, it has been reported that several transmembrane proteins could be
171 displayed on the surface of baculoviruses through their native transmembrane domains (11–
172 13). hPRR2 and 4 were incorporated in rBmNPV using C-terminal domain (amino acid 320–
173 511) of GP64 (Fig. 2B, left, lanes 2 and 3). This domain, including transmembrane domain, is
174 enough for incorporation of hPRR into rBmNPV. GP64s in rBmNPV-hPRR1, -hPRR2, and -
175 hPRR4 were detected by the polyclonal anti-gp64 antibody, which is against the 400–416
176 amino acid sequence of GP64 of rBmNPV (Fig. 2B, right). hPRR2, in which the prorenin-
177 binding domain is inserted into complete GP64, was examined together with GP64 (Fig. 2B,
178 right lane 2). However, the hPRR4 couldn't be detected with the same anti-GP64 antibody

179 because its molecular weight is similar to that of GP64 (Fig. 2B, right, lane 3). These results
180 show that hPRR, hPRR2 and hPRR4 was incorporated in rBmNPV.

181

182 **Analysis of hPRRs under nonreducing conditions** GP64 is present intracellularly in
183 the form of two types of homotrimers (trimers I and II) under nonreducing conditions (14).
184 Trimers I and II of GP64 with molecular weights above and below 220 kDa, respectively,
185 were observed in each rBmNPV. This was confirmed by western blot using the anti-Bm $gp64$
186 polyclonal antibody under nonreducing conditions (Fig. 3A). The hPRR1 (38.4 kDa) was
187 present as a monomer under both reducing and nonreducing conditions (Fig. 3B). The hPRR2
188 (288 kDa) was also observed above GP64 trimers I and II (around 220 kDa) when the anti-
189 Bm $gp64$ (Fig. 3A) and anti-FLAG M2 antibodies were used under nonreducing conditions
190 (Fig. 3B). Under reducing conditions it was observed between 80 and 100 kDa (95.8 kDa)
191 (Fig. 3A). This suggests that hPRR2 is also a trimer. The hPRR4 was not clearly observed by
192 the anti-Bm $gp64$ antibody under either reducing or nonreducing conditions because its
193 molecular weight is similar to that of GP64 (Fig. 3A). The hPRR4 (64.7 kDa) was detected
194 between 60 and 80 kDa under reducing conditions by using the anti-FLAG M2 antibody (Fig.
195 3B). Under nonreducing conditions, the hPRR4 was stacked in a well. This suggests that
196 hPRR4 formed aggregates. A protein band at 220 kDa was detected under nonreducing
197 conditions, which were also observed in rBmNPV-CP⁻-GGT2. This means that this band was
198 nonspecific (data not shown).

199

200 **Intracellular distribution of hPRRs** Expressed GP64 enters the cellular secretory
201 pathway where it is glycosylated, oligomerized, and transported to the plasma membrane (14).

202 During the budding and assembly of budded baculoviruses, GP64 is localized in discrete
203 areas on the plasma membrane, and these sites appear as budding sites (15, 16). Therefore,
204 whether each hPRR is able to localize in the plasma membrane in a manner similar to that of
205 GP64 or not was investigated. The hPRR (hPRR1 in this report) has been reported to localize
206 in the ER in HeLa-S3 cells (17) and in the intracellular vesicular complexes and plasma
207 membrane in cardiomyocytes (18). In this study, hPRR1 was observed in ER, at the same site
208 stained by ER tracker. However, hPRR2 and hPRR4 were observed in the periphery of ER
209 (Fig. 4). It indicates that expressed hPRR1, hPRR2 and hPRR4 entered into the secretory
210 pathway in Bm5 cells and the transmembrane domain in each variant is essential for its
211 distribution in Bm5 cells.

212

213 **Renin-binding by hPRR-displaying rBmNPVs** To confirm the expression of each
214 hPRR on the surface of each rBmNPV, SPR experiments were performed. Human renin was
215 immobilized in the wells of the CM sensor cuvette, and 5 µg of each rBmNPV was added to
216 the wells. Control rBmNPV (rBmNPV-CP'-GGT2) was found to bind to immobilized renin
217 to a slight extent (Fig. 5). In the cases of hPRR1 and hPRR2, the response increased with the
218 incubation time, indicating that rBmNPV-hPRR1 and -hPRR2 bound to the immobilized
219 renin, indicating that hPRR1 and hPRR2 were localized on the surfaces of the rBmNPVs
220 respectively. The response of rBmNPV-hPRR4 was at the same level as that of the control,
221 indicating that the binding of rBmNPV-hPRR4 to renin was nonspecific. This suggests that
222 although hPRR4 might be incorporated in rBmNPV-hPRR4 through the transmembrane
223 domain of GP64, it may not be functional.

224

Discussion

225

226

227 Surface display of recombinant proteins and functional peptides in baculovirus systems
228 has been reported earlier. The AcMNPV baculovirus has been mainly used in these systems,
229 but BmNPV has been used in very few cases (6, 7). In general, display of recombinant
230 proteins has been achieved by fusion with GP64 present on the baculovirus surface. However,
231 recombinant proteins with their own native transmembrane domains without fusion to GP64
232 was also expressed and displayed on the surface of baculoviruses (11–13, 19, 20). In this
233 study, we tried to express hPRR on the surface of BmNPV by fusing it with GP64 of
234 BmNPV or in its native form.

235 hPRR1 and 2 were expressed on the surface of rBmNPV. The hPRR1 without fusing to
236 GP64 was displayed on the surface of rBmNPV but not on that of AcMNPV (21). This
237 suggests that the mechanism of protein display on the surface of BmNPV was differed from
238 AcMNPV. All three hPRRs were not incorporated in rBmNPV-CP⁻, which lacks the cysteine
239 protease v-cath. This protease has a cathepsin L-like sequence, and its substrate specificity is
240 similar to that of cathepsin B (22). When insect cells are infected with a baculovirus, this
241 protease accumulates as a propeptide, and the death of host cells activates this propeptide
242 (23). Moreover, in baculovirus-infected larvae, this cysteine protease plays an important role
243 in the liquefaction of infected larvae. The relationship between this cysteine protease and the
244 incorporation of recombinant proteins has not yet been investigated.

245 The hPRR2, which was constructed by inserting the prorenin-binding domain of hPRR
246 into the site between ⁸¹Asp and ⁸²Pro in GP64, could be displayed on the surface of rBmNPV
247 in an active form. Deletion of the N-terminal half of AcMNPV GP64 did not prevent its

248 oligomerization. However, deletion of the predicted alpha-helical region (amino acids 298 to
249 346) and hydrophobic region II (amino acids 330 to 338) prevented the oligomerization of
250 *Orgyia pseudotsugata* multiple nucleopolyhedrovirus (OpMNPV) GP64 (14). hPRR2,
251 composing of the prorenin-binding domain of hPRR, inserted into full-length GP64, was
252 displayed on the surface of BmNPV, and its rBmNPV-hPRR2 was able to bind to human
253 renin. However, hPRR4, which was constructed by inserting the prorenin-binding domain of
254 hPRR into the site between ⁸¹Asp and ³²⁰Met of GP64, was expressed, but rBmNPV-hPRR4
255 did not bind to renin (Fig. 5). In previous report, purified hPRR4 was not able to bind to
256 human renin (24). The hPRR4 was detected under nonreducing conditions in the upper part
257 of the SDS-PAGE gel in this study (Fig. 3), suggesting that it is present in rBmNPV particles
258 as non-functional aggregates. GP64 from AcMNPV or BmNPV has 15 cysteine residues and
259 is expressed as a trimer. Moreover, the oligomerization domain, which is in the form of a
260 leucine zipper motif, is located between amino acids 298 and 339 in GP64 of AcMNPV (25).
261 This oligomerization domain is also present in GP64 of BmNPV. Twenty-five amino acids
262 from this domain in GP64 were deleted in hPRR4, which may result in the aggregation of
263 hPRR4 (25).

264 Incorporation of hPRRs in rBmNPVs and immunofluorescent microscopic observation
265 suggest that hPRR1 was expressed in ER, but hPRR2 and hPRR4 were in the periphery of ER
266 (Fig. 4). This indicates that the C-terminal domains of hPRR and BmNPV GP64, which
267 include the transmembrane and cytoplasmic domains, direct proteins to go through the
268 secretory pathway in Bm5 cells and be expressed in the membrane fraction in the secretory
269 pathway. Previously, it has been reported that hPRR1 localizes in the ER in HeLa-S3 and Sf-
270 9 cells (17) and in the intracellular vesicular complexes and plasma membrane in

271 cardiomyocytes (18). In this study, most hPRR1 was observed mainly at ER. During the
272 budding and assembly of budding baculoviruses, nucleocapsids are thought to migrate
273 through the cytoplasm and bud through the plasma membrane at sites where GP64 is
274 concentrated (16). It suggested hPRR1 might be also expressed in the plasma membrane and
275 then hPRR1 may be displayed on the surface of rBmNPV. However, proteins expressed in
276 ER were also displayed on the surface of baculoviruses (20). It is possible that hPRR1
277 expressed in ER was displayed on the surface of rBmNPV, not expressed in the plasma
278 membrane.

279 hPRR1, with its native single transmembrane domain, and hPRR2 with the
280 transmembrane domain of GP64, were successfully displayed on the surface of rBmNPV and
281 showed biological function. The BmNPV-based surface display system may be useful for
282 analyzing receptors and the baculovirus drug delivery system, which require very high titers
283 of baculoviruses.

284

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288

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377

Figure legends

378

379 FIG. 1. Constructions of hPRRs. hPRR1 is native hPRR with an additional FLAG peptide
380 sequence inserted between the signal peptide and prorenin-binding domain. hPRR2 was
381 constructed by inserting the prorenin-binding domain of hPRR between ⁸¹Asp and ⁸²Pro of
382 GP64 from BmNPV. hPRR4 was constructed by inserting the prorenin-binding domain of
383 hPRR into the site between ⁸¹Asp and ³²⁰Met of partially deleted GP64.

384

385 FIG. 2. Expression of individual hPRRs in rBmNPV-CP⁻ (A) and rBmNPV (B). Infected
386 Bm5 cells and culture supernatant were separated from cell cultures. Each rBmNPV was
387 isolated from its culture supernatant by ultracentrifugation. An aliquot of each sample was
388 analyzed by CBB staining and western blotting with the mouse anti-FLAG M2 antibody or
389 rabbit anti-BmGP64 antibody. Lanes 1, 2, and 3 in (A) represent rBmNPV-CP⁻-hPRR1, -
390 hPRR2, and -hPRR4, respectively, while lanes 1, 2, and 3 in (B), rBmNPV-hPRR1, -hPRR2,
391 and -hPRR4, respectively.

392

393 FIG. 3. Analysis of oligomerization under both reducing and nonreducing conditions. Each
394 sample was electrophoresed on SDS-PAGE gels under both reducing and nonreducing
395 conditions and then analyzed by western blotting using the rabbit anti-Bm_{gp}64 antibody (A)
396 and the mouse anti-FLAG M2 antibody (B). Lanes 1, 2, and 3 represent rBmNPV-hPRR1, -
397 hPRR2, and -hPRR4, respectively.

398

399 FIG. 4. Immunofluorescence localization of each hPRR in Bm5 cells infected with rBmNPVs.

400 Bm5 cells were infected with each rBmNPV (rBmNPV-hPRR1, -hPRR2, and -hPRR4) and
401 cultivated for 2 days prior to staining with the mouse anti-FLAG M2 antibody, and followed
402 by staining with the FITC-conjugated goat anti-mouse IgG antibody and ER-Tracker™ Red.
403 Bar indicates 10 μm. Each hPRR was shown to be green fluorescence by staining with the
404 mouse anti-FLAG M2 antibody. ER was stained as red fluorescence.

405

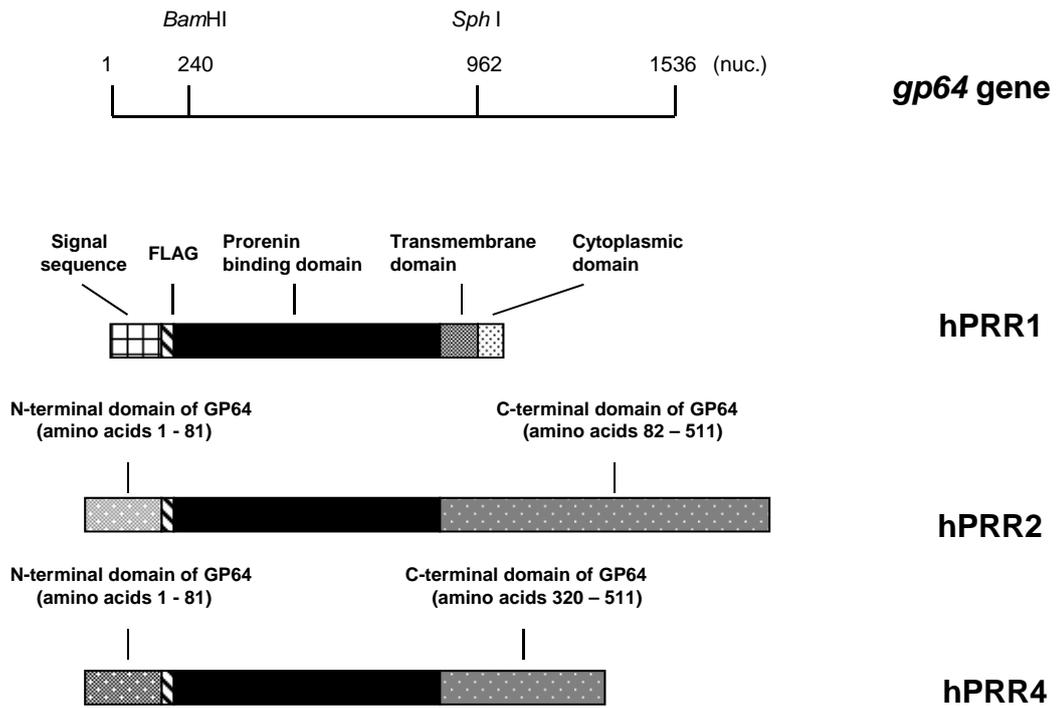
406 FIG. 5. SPR analysis of the binding of rBmNPV-hPRR1, -hPRR2, and -hPRR4 to human
407 renin. Each rBmNPV was isolated by ultracentrifugation, and 5 μg was subjected to SPR
408 analysis. Human renin (2.5 μg) was immobilized on the surface of the CM sensor cuvette,
409 and unreacted CM groups were blocked with ethanolamine. The sensor cuvette that had no
410 immobilized human renin was also blocked with ethanolamine and used as a control for
411 nonspecific binding. PBS (pH 6.2) was used as the running buffer. The arrows indicate the
412 point at which the samples were added to the sensor cuvettes. rBmNPV-CP⁻-GGT2 was used
413 as the negative control.

414 Table 1. Primers

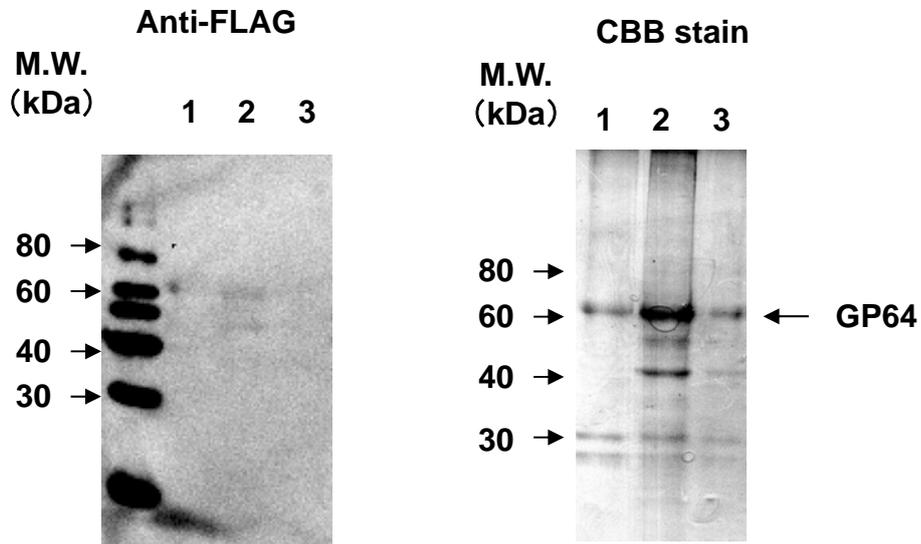
Name	5' → 3'
Bmcp64-F	caccggtaccatggtaggcgctattgtttatacg
Bmcp64-R	cccaagctttaatattgtctactattacggtttc
bgl-hPRR-F	gaagatctccgactacaaggacgacgacgacaag
bgl-hPRR-R	gaagatctagatattcaaaattatacttatatgc
sph-hPRR-R	acatgcatgcatatattcaaaattatacttatatgc

415

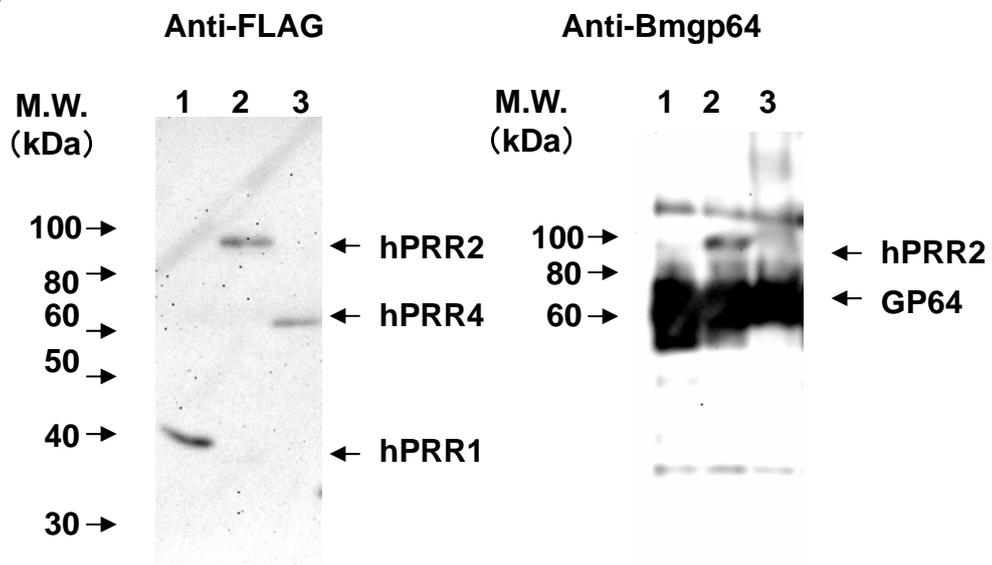
Fig. 1, Kato et al.



(A)



(B)



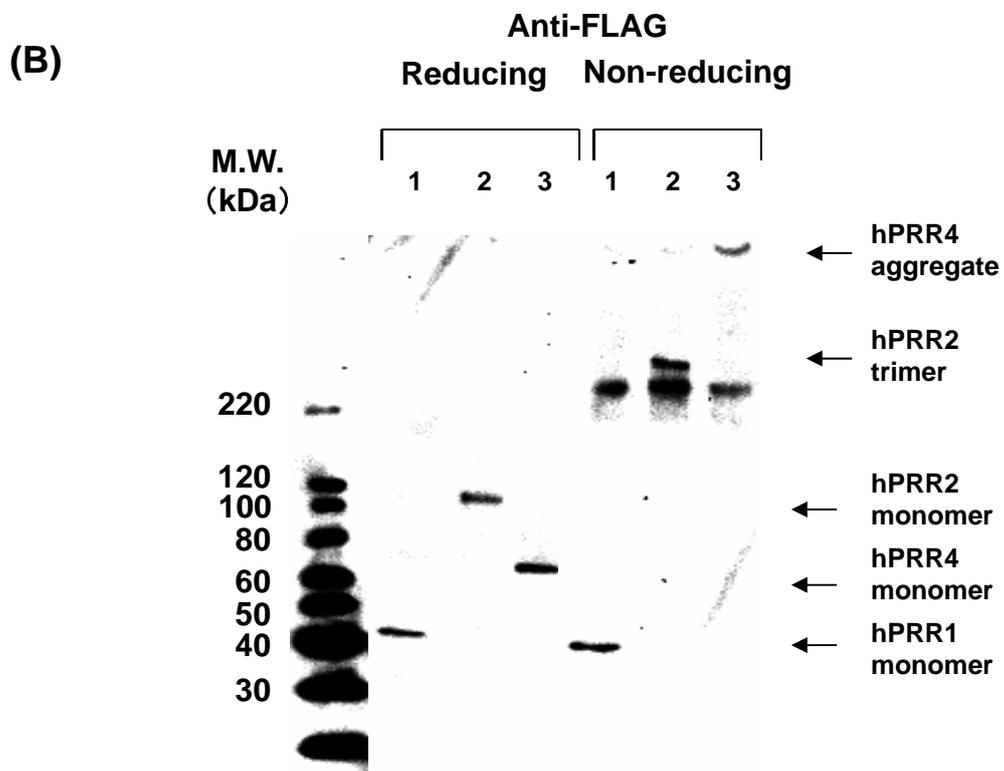
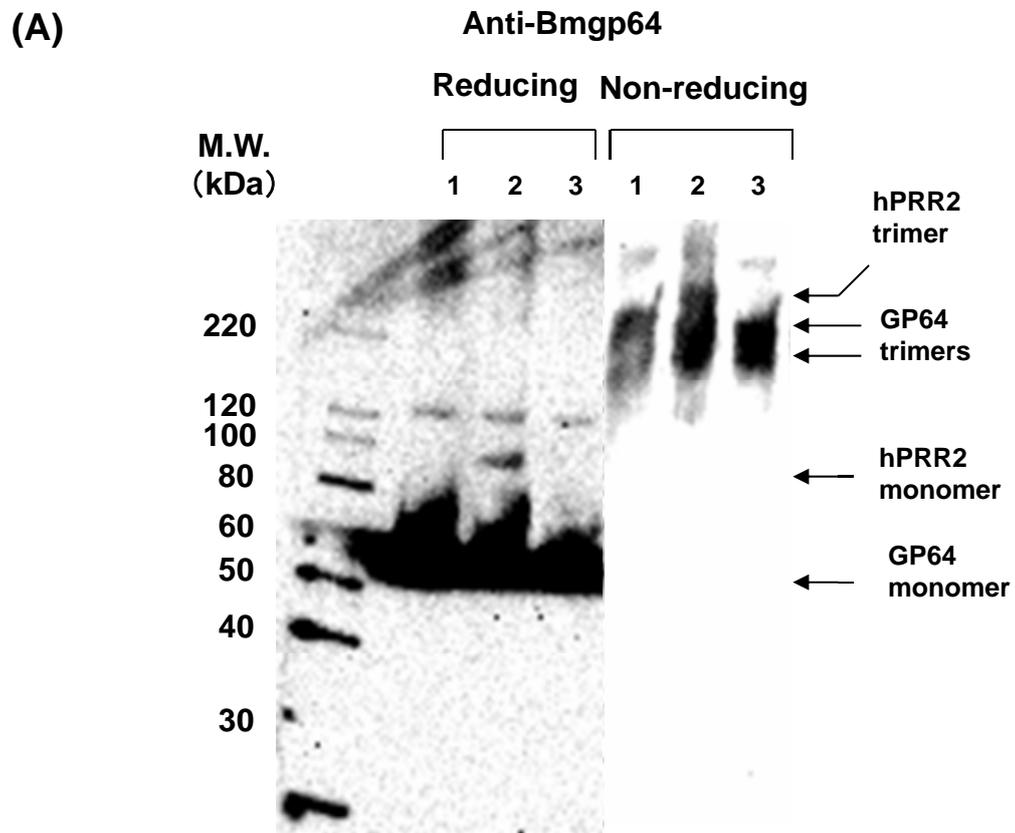


Fig. 4, Kato et al.

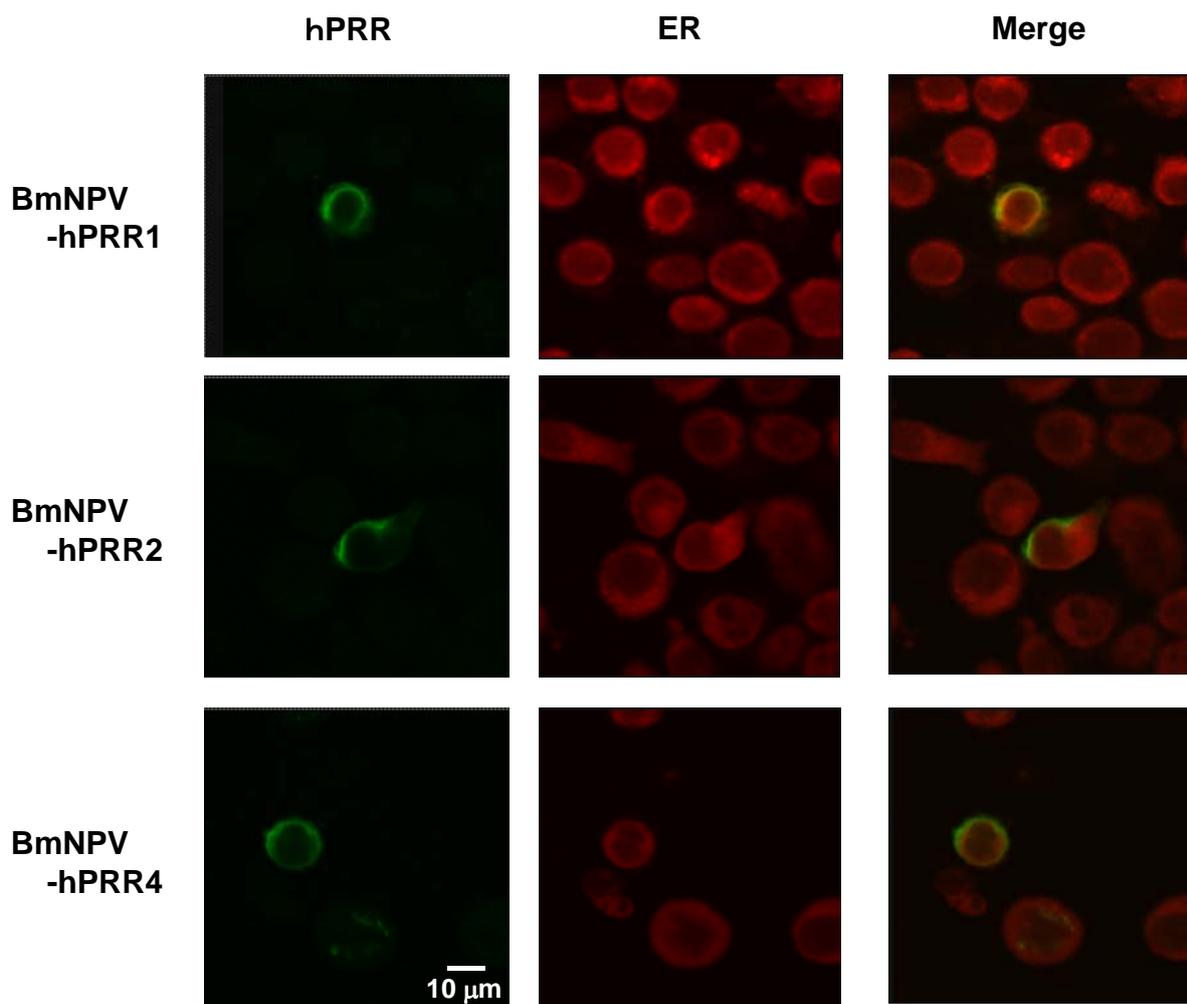


Fig. 5, Kato et al.

