



23 results of the VBN and silica sol-gel interface indicate a general applicability for VBN-based  
24 bioseparations and biosensing applications.

25

## 26 **Keywords**

27 Virus-based nanomaterial; molecular pharming; nanobiotechnology; tobamovirus; plant-made  
28 pharmaceuticals; silica sol-gel; bioseparations; biosensing; antibody purification; monoclonal  
29 antibody.

30

## 31 **Abbreviations**

32 Cy5-TMV, Cyanine 5 conjugated TMV; hIgG, human immunoglobulin G; PEG, polyethylene  
33 glycol; PLGA, poly(lactic-co-glycolic acid); TMOS, tetramethyl orthosilicate; TMV, tobacco  
34 mosaic virus; TVCV, turnip vein clearing virus.

## 35 **1. Introduction**

36 Virus-based nanomaterials (VBNs) are being studied for various medical applications as versatile  
37 nanomachines that can be manufactured at an industrial scale with high fidelity and low costs<sup>1,2</sup>.  
38 Reports in literature have primarily focused on the design of novel VBNS. With sufficient novelty  
39 and value of VBNS having been demonstrated in several application areas, there is a need to  
40 consider more advanced VBN-based systems to leverage the potential of existing VBNS and move  
41 this technological platform towards the clinic and market.

42 To date, several studies have explored VBNS as structural and/or functional elements within larger  
43 system arrangements. Recent examples include: hot-melt extrusion of trivalent vaccine candidates  
44 mixed into slow-release PLGA implants<sup>3</sup>; surface conjugation of osteogenic VBN nanofibers to  
45 the surface of 3D-printed bioceramic bone scaffolding<sup>4</sup>; electrostatic layer-by-layer assembly of  
46 free-standing VBN biofilms<sup>5</sup>; surface adsorption of immunosorbent VBNS onto gold sensor chips<sup>6</sup>;  
47 magnetic particle conjugation to immunosorbent VBNS for protein purification<sup>7</sup>.

48 Silica sol-gel chemistries represent an alluring set of matrices for bioencapsulation and more  
49 advanced VBN-based systems. Extensive literature supports favorable silica sol-gel chemistry  
50 characteristics of high structural uniformity, stability, pore size tunability, optical properties, and  
51 biodegradability for various biomedical applications<sup>8,9</sup>. A range of live cells<sup>10-12</sup> and enzymes<sup>13-15</sup>  
52 have been studied for bioencapsulation in silica sol-gel matrices to stabilize viability and activity  
53 as well as to improve ease of use for the intended application. There have not yet been any such  
54 studies using VBNS, with the exception of a study of viral encapsulation focused on extended  
55 release of viral vectors for gene therapy<sup>16</sup>.

56 In this study we present entrapment and utility of plant virus-based immunosorbent nanoparticles  
57 (VINs) in silica sol-gel matrices by pore confinement, representing a novel system configuration  
58 for VBNs in general. VINs display antibody-binding proteins on their external coat protein surface,  
59 and they have been used as simple and bioregenerable reagents for biosensing and therapeutic  
60 antibody purification<sup>6,17,18</sup>. This entrapment of VINs represents the first use of plant virus-based  
61 VBNs in a silica sol-gel matrix and the first application of VBN technology for utility in an intact  
62 silica sol-gel matrix. We demonstrate that the silica sol-gel matrix can immobilize these large  
63 biomolecules over a long-duration (~30 days) in an environment that preserves their  
64 immunosorbent capture and elution functionalities and is sufficiently diffusive for antibodies to  
65 freely enter and leave the matrix. We also show that the silica sol-gel encapsulated VINs can be  
66 used to purify antibodies from a complex mixture, in this case, crude *Nicotiana benthamiana* plant  
67 extract (representing the production of therapeutic antibodies in plants, formerly known as  
68 molecular pharming<sup>19</sup>), to overcome reusability and bioprocessing challenges encountered when  
69 using VINs in solution to purify antibodies.

70

## 71 **2. Materials & Methods**

### 72 **2.1 Virion production**

73 Turnip vein clearing virus (TVCV) presenting a coat protein display of the D and E domains of  
74 *Staphylococcus aureus* Protein A was used as the plant virus-based immunosorbent nanoparticles.  
75 The VINs were produced according to a previously reported study by agroinfiltration of *N.*  
76 *benthamiana* plants using expression vector pICH25892 and processed with a polyethylene glycol

77 (PEG)-based purification scheme<sup>7</sup>. Final yield was approximately 300 mg VIN per kg *N.*  
78 *benthamiana* leaf tissue, as determined by Bradford total soluble protein assay.

79 Wild-type tobacco mosaic virus (wt-TMV) was produced via mechanical inoculation of ~5-week  
80 old *N. benthamiana* plants by lightly sprinkling three leaves per plant with Celite<sup>®</sup> 545 (Millipore  
81 Sigma, Burlington, MA, USA) as an abrasive aid and gently rubbing 100  $\mu$ L of 0.01 mg/mL wt-  
82 TMV in 0.01 M potassium phosphate buffer pH 7.0 per each of the three leaves. The plant leaves  
83 were washed with water 20 minutes after inoculation. Leaf tissue was collected after infection  
84 symptoms presented ~1 week post-inoculation and frozen at -80 °C for storage.

85 Extraction of wt-TMV from frozen *N. benthamiana* leaf tissue was performed with a 5:1 (w/v)  
86 extraction ratio with 0.1 M potassium phosphate pH 7.0 with 0.1% (v/v) beta-mercaptoethanol  
87 using a chilled mortar and pestle. The plant extract was filtered through three-layered cheese cloth,  
88 mixed with equal parts chloroform and n-butanol up to 1:1 (v/v) ratio, centrifuged at 8,000 x g and  
89 4 °C for 10 minutes, and the upper aqueous phase layer was collected. PEG-based precipitation  
90 was performed by addition of 4% (w/v) PEG 8,000 and 1% (w/v) NaCl, incubation of the mixture  
91 for 30-60 minutes at 4 °C, and centrifugation at 8,000 x g and 4 °C for 15 minutes. The pellet was  
92 resuspended in 50 mM Tris-HCl pH 7.0 with a glass rod and let to sit at 4 °C for 30-60 minutes.  
93 The resuspended solution was centrifuged again at 8,000 x g and 4 °C for 10 minutes. The resulting  
94 supernatant was then ultracentrifuged at 50,000 RPM and 4 °C for 90 minutes using a 70.1 Ti rotor  
95 (Beckman Coulter, Brea, CA, USA) with 1 mL 15% sucrose cushion per tube. The pellet was  
96 resuspended again in 50 mM Tris-HCl pH 7.0 with a glass rod and let to sit at 4 °C overnight. The  
97 resuspended solution was added on top of a 10 – 40% (w/v) sucrose gradient and ultracentrifuged  
98 at 30,000 RPM and 4 °C for 90 minutes using a SW40 swinging bucket rotor. A 50% sucrose  
99 solution was used as a plug to fractionate and collect the wt-TMV containing solution.

100 Purification of wt-TMV was also performed according to the VIN purification protocol. Final yield  
101 was ~ 800 mg wt-TMV per kg *N. benthamiana* leaf tissue, as determined by UV absorbance  $A_{260\text{nm}}$   
102 spectroscopy measurements.

103 The fluorescent reporter particle used in this study, a chemical conjugation of the Cyanine 5  
104 fluorophore to the exterior of the wt-TMV coat protein (Cy5-TMV), was generated using purified  
105 wt-TMV and a two-step reaction composed of a diazonium coupling and click reaction step  
106 according to a method presented by Bruckman and Steinmetz<sup>20</sup>. Spin filters were used to separate  
107 the Cy5-TMV from leftover reaction reagents. Fluorophore dye loading is calculated using  
108 extinction coefficients at 260 nm of  $\epsilon_{\text{Cy5}} = 250,000 \text{ mL/cm/mg}^{20}$  and  $\epsilon_{\text{TMV}} = 3 \text{ mL/cm/mg}^{21}$ .

109

### 110 **2.3 Silica sol-gel synthesis**

111 The silica sol-gel synthesis was performed according to the method for entrapment of liposomes  
112 in silica gel detailed by Zeno and team<sup>22</sup>. In brief, 3.8 mL of tetramethyl orthosilicate (TMOS) was  
113 added dropwise to 2.75 mL of 0.002 M HCl in a beaker chilled by an ice bath. This mixture was  
114 tip sonicated for 15 minutes, added to a round-bottom flask, and rotary evaporated at 340 mbar  
115 reduced pressure and 50 °C for 2-3 minutes before being passed through a 0.22  $\mu\text{m}$  filter to yield  
116 the final silica sol. The sol was combined with 3 parts PBS by volume and 1 part of a PBS solution  
117 containing either VIN, wt-TMV, or Cy5-TMV at ~0.300 mg/mL concentration (according to total  
118 soluble protein measurement for VIN or UV-vis measurement for TMV).

119 The prepared solution was aliquoted as 40  $\mu\text{L}$  droplets (or 2  $\mu\text{L}$  droplets for the fluorescent  
120 microscope imaging) onto parafilm at room temperature for gel bead formation. Post-gelation (~1  
121 hr) beads were submerged in equilibration buffer (PBS pH 7.0) at 4 °C prior to use.

122

## 123 **2.4 Binding and elution of human immunoglobulin G**

124 The binding and elution of human immunoglobulin G (hIgG) with gel beads was performed  
125 following a series of processing steps for batch operation: equilibration, sample loading, impurity  
126 wash, and elution. Equilibration consists of a 150  $\mu$ L PBS buffer bath to submerge each 40  $\mu$ L gel  
127 bead, which was individually contained in a 2 mL tube, that was nutated at 4 °C for a minimum  
128 duration of 24 hours and exchanged with fresh PBS buffer a minimum of four times throughout  
129 the duration. Previous reports in literature have provided 24 hours of equilibration prior to silica  
130 sol-gel use<sup>23</sup>, presumably to ensure stabilization of internal pore charges.

131 The hIgG samples were prepared as 80  $\mu$ L of 0.25 mg/mL hIgG in either PBS or 0.22  $\mu$ m filtered  
132 *N. benthamiana* extract. The clarified *N. benthamiana* extract was prepared by 3:1 (v/w) chilled  
133 mortar and pestle extraction of 6-week-old *N. benthamiana* leaf tissue frozen at -80 °C, filtration  
134 through 4-layered cheesecloth, centrifugation at 8,000 x g and 4 °C for 15 minutes, and an  
135 additional filtration through a dead-end 0.22  $\mu$ m syringe filter. For sample loading, all equilibration  
136 buffer was removed and the 80  $\mu$ L of sample was added to submerge the silica sol-gel bead, which  
137 was kept at 4 °C nutating for 24 hours.

138 Impurity wash consisted of a minimum of four buffer exchanges into fresh PBS buffer over a  
139 period of no less than 24 hours at 4 °C while nutating.

140 Elution consisted of PBS buffer removal, addition of 80  $\mu$ L 0.1 M glycine buffer pH 2.5,  
141 incubation for 4 hours at 4 °C while nutating, recovery of the elution solution, and pH  
142 neutralization with 0.1 volumes of 0.1 M Tris-HCl pH 9.0.

143

## 144 **2.5 Protein analysis**

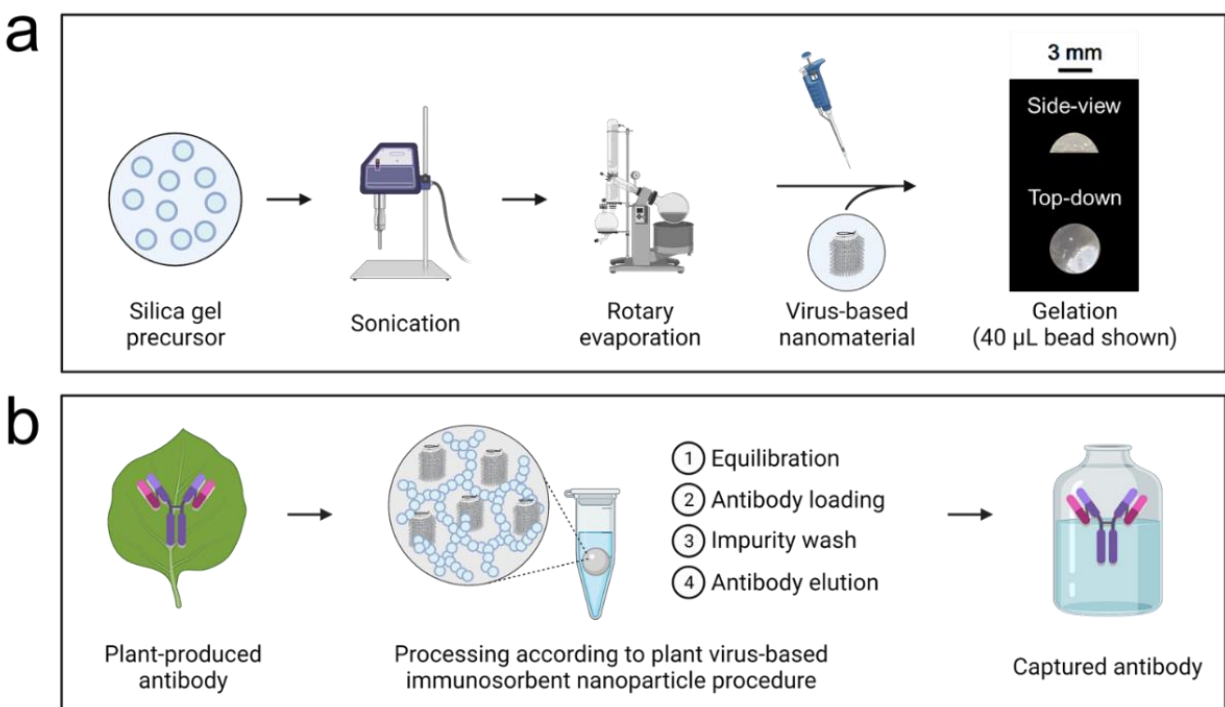
145 Protein analyses using methods of the Bradford assay, SDS-PAGE, western blot, and transmission  
146 electron microscopy (TEM) were performed according to previously reported methods<sup>7</sup>. UV-vis  
147 measurements were made using a Quartz SUPRASIL<sup>®</sup> quartz cuvette (Hellma Analytics,  
148 Plainview, NY, USA) and a SpectraMax<sup>®</sup> M4 spectrophotometer (Molecular Devices, San Jose,  
149 CA, USA). Fluorescence microscopy was performed with an Eclipse 80i microscope (Nikon,  
150 Tokyo, Japan) equipped with a 4x objective. Immediately before imaging, Cy5-TMV loaded beads  
151 were transferred from a PBS bath to a slide and excess PBS was removed with a pipette. All images  
152 were collected using the same exposure time.

153

## 154 **3. Results**

155 An illustration of the silica sol-gel functionalization with entrapped plant virus-based  
156 immunosorbent nanoparticles and the example use case presented in this study is depicted in  
157 Figure 1.





158

159 **Figure 1.** Illustrated schematics of (a) synthesis of sol-gel functionalized with entrapped virus-based  
160 based nanomaterials (including example photographs of gels synthesized in this study), and (b) the  
161 example use case presented in this work of gel-entrapped plant virus-based immunosorbent  
162 nanoparticles for affinity purification of plant-made antibodies.

163

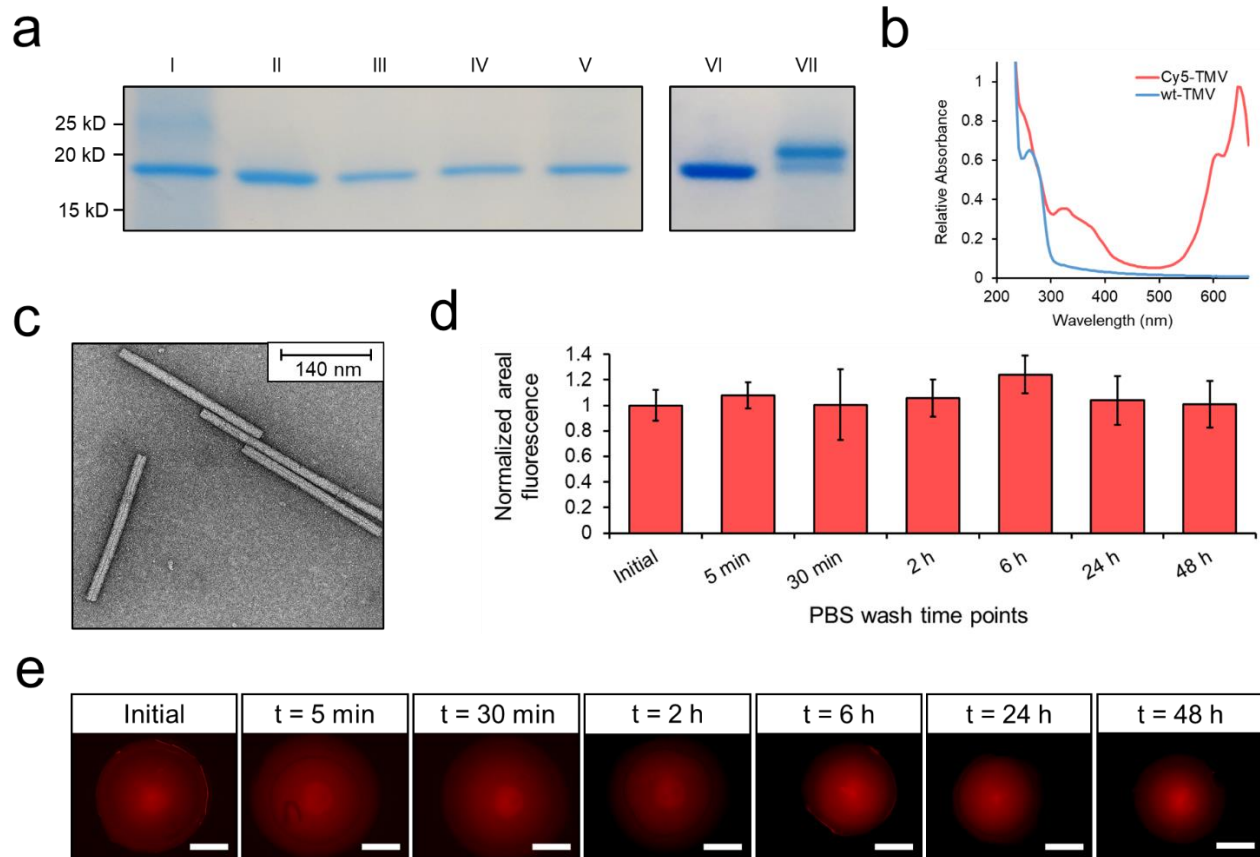
164 Firstly, a fluorescent reporter VBN was developed to investigate silica sol gel entrapment. As  
165 shown in Figure 2a, we produced wt-TMV *in planta* and purified the wt-TMV to ultra-high levels  
166 of purity. We then generated Cy5-TMV as a reporter system using previously reported methods to  
167 conjugate Cyanine 5 fluorophore to the exterior surface of wt-TMV coat<sup>20</sup>. UV-Vis absorbance  
168 spectroscopy 200 – 700 nm spectrums of wt-TMV and Cy5-TMV are comparable to the spectrum  
169 results of the previously reported study (Figure 2b). Notably differences of the Cy5-TMV from  
170 the wt-TMV include the introduction of an absorbance peak at around 324 nm, which is consistent

171 with the introduction of a diazonium bond (needed for the alkyne addition reaction), and another  
172 peak at around 646 nm, which is consistent with the sulfo-Cy5 azide absorbance. We estimate a  
173 dye loading of ~32%, which is to say that on average ~680 of an estimated 2,130 coat proteins per  
174 assembled virion were conjugated with a Cy5 fluorophore on the exterior surface. Negative stain  
175 TEM images were used to confirm structural integrity of the Cy5-TMV (Figure 2c).

176 Plant virion entrapment due to crosslinking physical immobilization during silica sol-gel bead  
177 synthesis was then assessed using the Cy5-TMV reporter system. Normalized areal fluorescence  
178 measurements are shown over the course of a 48-hour study in Figure 2d. Representative  
179 fluorescence microscopy images used in the quantitative measurements are shown in Figure 2e.  
180 Fluorescence results indicate that the initial Cy5-TMV concentration in the silica sol-gel matrix  
181 was maintained throughout the period of examination. There were appreciable decreases in gel  
182 bead volume due to minor breakages and shrinkage over the course of multiple microscope images  
183 (Supplementary Information, Figure S1). We attribute this observed behavior to the small bead  
184 size (2  $\mu$ L), forceps manipulation, and multiple exposures to a dry environment used for  
185 fluorescent microscope imaging. This observation of areal shrinkage was not noted in the beads  
186 used for functional testing (40  $\mu$ L).

187 There was no evidence of VIN or wt-TMV lost in the PBS wash buffer during standard  
188 equilibration to suggest incomplete entrapment. However, we performed an experiment using  
189 lower wash buffer volumes (1 vol buffer: 1 vol gel) to improve the limit of detection and found  
190 that < 10% of the initial VIN or wt-TMV mass added to the gel was recovered in the wash buffer  
191 during equilibration, peaking at an initial brief wash timepoint and decaying rapidly from there  
192 (Supplementary Information, Figure S2). This suggests that a small amount of the VIN or wt-TMV

193 was not adequately entrapped during sol-gel synthesis and that it can be readily cleared from  
194 solution.



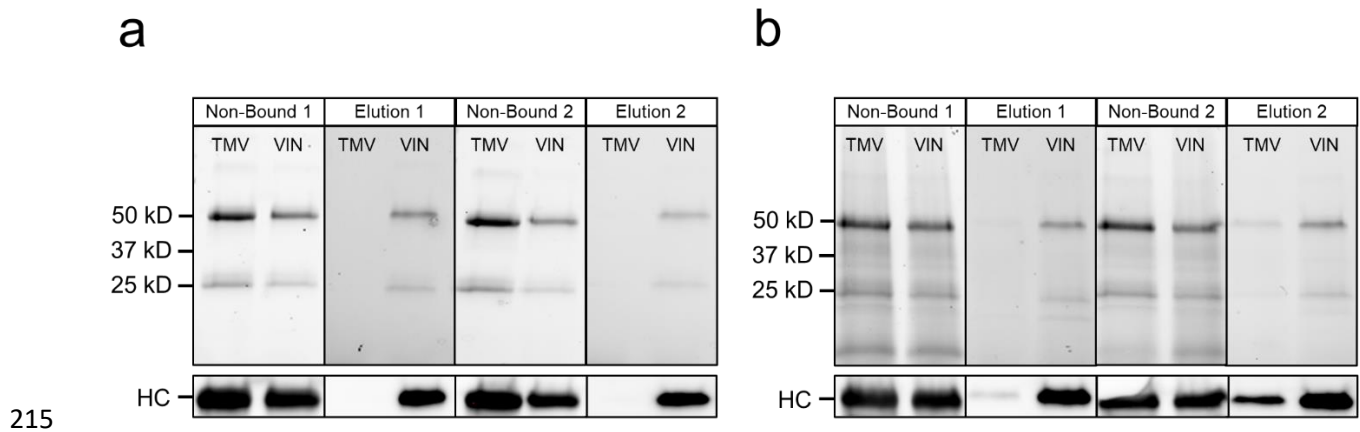
195

196 **Figure 2.** Purification and immobilization of fluorescently labelled plant virions (a) SDS-PAGE  
197 results of an ultracentrifuge-based purification of wt-TMV from *Nicotiana benthamiana* leaf  
198 tissue, which is then chemically conjugated with Cy5 dye in a two-step reaction to generate Cy5-  
199 TMV. Lane definitions: I – filtered plant extract, II – PEG-precipitated pellet resuspension, III –  
200 microfuge pellet, IV – ultracentrifuge pellet resuspension, V – ultracentrifuge wash, VI – initial  
201 wt-TMV pre-conjugation, VII – Cy5-TMV post-conjugation. (b) UV-Vis absorbance spectra of  
202 wt-TMV and Cy5-TMV. (c) Negative stain transmission electron microscope image of the Cy5-  
203 TMV. (d) Normalized areal fluorescence per 2  $\mu$ L volume silica bead containing Cy5-TMV over  
204 48 hours. Beads were exchanged into fresh PBS buffer after each measurement. Error bars

205 represent one standard deviation with biological triplicate. (e) Fluorescent microscope images of  
206 the silica sol-gel beads containing Cy5-TMV over 48 hours of submerged gel wash. Scale bars  
207 represent 500  $\mu\text{m}$ .

208

209 The bind-and-elute immunosorbent functionality of the VIN-entrapped silica sol-gel beads was  
210 then examined in this study. Assessment of this functionality using samples of hIgG in PBS and  
211 hIgG in clarified *N. benthamiana* plant extract is shown in each condition for two consecutive  
212 cycles of bind-and-elute in Figure 3. The results show that the VIN-entrapped beads yielded a high  
213 recovery of hIgG in the elution step for both the PBS and plant extract conditions, whereas the  
214 TMV-entrapped beads did not.



216 **Figure 3.** Use of silica sol-gel functionalized with entrapped plant virus-based immunosorbent  
217 nanoparticles (VIN) to purify monoclonal antibodies. Reducing condition SDS-PAGE (upper) and  
218 western blot (lower) results of the non-bound liquid sample after loading and low pH elution of  
219 human immunoglobulin G for a first and second use cycle shown for a sample loading consisting  
220 of human immunoglobulin G spiked into a solution of (a) clean PBS and (b) sterile-filtered

221 *Nicotiana benthamiana* extract. Lanes are loaded with fixed volume. HC, heavy chain of the  
222 human immunoglobulin G.

223

224 The small amount of hIgG in the elution for the wt-TMV negative control using *N. benthamiana*  
225 extract (Figure 3b) indicates that the bind-and-elute functionality cannot be solely ascribed to  
226 specific Protein A-Fc interactions. Additional negative controls of silica sol-gel synthesized with  
227 bovine serum albumin and with no proteinaceous solution also corroborate these results (data not  
228 shown). The hIgG recovered in the elution for the plant extract-containing sample condition  
229 indicates some degree of non-specific binding interactions which may be present. Minor yellowish  
230 coloration of the gel bead, which closely resembled the color of the loaded sample, after impurity  
231 washing provides visual support for this observation (data not shown).

232 The second use cycle of hIgG recovery in elution indicates that the VIN functionality is preserved  
233 over multiple uses. A three-week lag period between the gel synthesis and completion of the  
234 second use cycle also provides additional support for the entrapment and stability of VIN in the  
235 silica sol-gel environment. An increase in hIgG recovery for the second use cycle elution of the  
236 wt-TMV bead and plant extract-containing sample condition is observed, which may indicate an  
237 undesired behavior for performance such as incomplete hIgG elution from the first use cycle or an  
238 additive effect of the non-specific binding mechanism between the first and second use cycle such  
239 as what may be observed should a subset of the plant extract constituents accumulatively bind to  
240 the silica sol-gel matrices.

241 The hIgG captured from the plant extract sample was recovered at ~60% purity in both use cycles,  
242 based on gel densitometry measurement. The SDS-PAGE results qualitatively support that the

243 hIgG purity was significantly increased from the capture and elution procedure. A reduction in  
244 recovery for the second use cycle was observed for hIgG in PBS (~50% first elution) and in plant  
245 extract (~80% first elution).

246 A substantial fraction of the hIgG sample loaded was not bound in either the VIN or wt-TMV  
247 conditions. We attribute this to the excess loading concentration, which was used in this proof-of-  
248 concept study to determine maximal elution given the employed gel synthesis and operational  
249 configuration. Each gel bead was synthesized containing a total of ~3  $\mu\text{g}$  VIN, which, based on  
250 previous study of VIN in free solution recovering ~1.5 mg hIgG/mg VIN in excess hIgG  
251 conditions, can be estimated to correspond to a binding capacity of ~4.5  $\mu\text{g}$  hIgG. Each gel was  
252 incubated in a liquid bath containing ~20  $\mu\text{g}$  hIgG. Based on Bradford total soluble protein assay  
253 results, we recovered ~1.9  $\mu\text{g}$  hIgG from PBS for the first cycle elution from the immobilized VIN  
254 sol-gel bead, corresponding to ~10% hIgG recovery of the initial sample load. This binding  
255 capability represents ~42% of that of VIN in free solution.

256

#### 257 **4. Discussion**

258 In this study, we report encouraging proof-of-concept results for novel VBN entrapment and  
259 functionality within the pores of silica sol-gel matrices with broad applicability in protein  
260 purification and biosensing. VIN-containing silica sol-gel beads were used to capture hIgG from  
261 PBS or plant extract and elute the hIgG in a low pH environment for two consecutive use cycles  
262 spanning approximately 30 days post-synthesis. The high surface area to volume ratio of the rod-  
263 like VBNs used in this study proved to be an amenable geometry for pore entrapment with  
264 considerable active binding site availability without the need for significant optimization.

265 Future work to investigate the significance of VBN morphology on performance would be valuable  
266 for understanding the possible VBN silica sol-gel design space. Relevant characteristics for  
267 investigation include the geometry of the virion, either rod-like or icosahedral, the rigidity of the  
268 geometry, given that rod-like virions can be classified as stiff (e.g., tobacco mosaic virus) or  
269 flexible (e.g., potato virus X), and the size of the virion, which can be readily extended in rod-like  
270 virions through genome augmentation.

271 Non-specific binding interactions of the hIgG to the silica sol-gel were observed in this study, as  
272 observed for the wt-TMV and plant extract sample condition. We hypothesize that constituents of  
273 the plant extract bind to the silica sol-gel and in turn the extract-gel complex increases the non-  
274 specific binding interactions with hIgG. Additionally, we hypothesize that the low pH elution  
275 conditions (pH 2.5) and the isoelectric points of the silica sol-gel matrix (pH 2.0 for silica) and  
276 hIgG (> pH 6.0) may generate a non-specific binding environment that could result in incomplete  
277 hIgG elution into the bulk liquid. This behavior was not observed in the experimental execution  
278 of this study but should be considered in the future. The isoelectric points of wt-TMV coat protein,  
279 pH  $\sim 4^{24}$ , and VIN coat protein fusion, pH  $\sim 3.7$  as estimated using ExPASy  
280 ([web.expasy.org/compute\\_pi/](http://web.expasy.org/compute_pi/)), are both net negatively charged at the neutral gelation condition  
281 and not expected to non-specifically bind with the silica sol-gel matrix. We do suggest that future  
282 works more rigorously resolve concerns of non-specific binding interactions through optimization  
283 of silica sol-gel composition, perhaps considering doping (3-aminopropyl) triethoxysilane into the  
284 formulation, as has been successfully employed in previous studies<sup>25</sup>.

285 The reduction of effective binding capacity for silica sol-gel entrapped VIN was within the range  
286 of previously reported reductions in enzyme activity; silica sol-gel entrapped horseradish  
287 peroxidase and glucose-6-phosphate dehydrogenase enzymes were reported to exhibit specific

288 activity of 73% and 36% of the specific activities of the free enzymes, respectively<sup>23</sup>. Further  
289 investigation is required to understand the relative contributions of protein activity modulation,  
290 such as from VIN exposure to gelation conditions or the internal pore environment pH, and loss  
291 of accessibility of the active site, such as from diffusional limitations in the sol-gel matrix or  
292 partitioning of VIN into hIgG-inaccessible silica sol-gel matrix pores.

293 The findings of this study were consistent with the results of Kangasniemi and team for extended  
294 release of adenovirus from silica sol-gel in finding that the VBNs in silica sol-gel were stable for  
295 extended durations (weeks to months) and could be evenly distributed in the sol-gel post-  
296 entrapment, as noted by the nearly linear VBN release profile during *in vitro* silica dissolution in  
297 their study<sup>16</sup>. The fundamental difference in observations was that Kangasniemi and team showed  
298 VBN activity to be retained for VBN that were released into free solution via silica dissolution,  
299 whereas this study reports VBN activity while entrapped within the silica sol-gel matrices. Our  
300 work also presents the first entrapment of plant virus-based VBN in silica sol-gel, which can be  
301 used in a wide variety of applications not amenable to mammalian virus-based VBN due to  
302 inherent advantages of safety (e.g., non-infectious to humans) and inexpensive and simple  
303 production.

304 The VIN silica sol-gel system presented in this study could serve as the foundation for a robust  
305 and reusable platform for biosensing. For example, the VIN system could be readily configured  
306 for immunosensing, in which the desired detection antibody would be first introduced to and fixed  
307 (via non-covalent bonding) in the VIN-entrapped gel matrix, allowing the subsequent addition of  
308 the desired analyte solution for sensing. The detection antibody could be removed via low pH  
309 elution and the same VIN-entrapped gel could be reused with a different detection antibody, as  
310 desired. This hypothesized example system may provide means for easy reuse and flexibility of



311 sensing targets as well as an enhanced sensitivity over antibody-only systems due to the increased  
312 sensing surface area of the structural scaffolding of the VIN. This hypothesis of the potential for  
313 increased sensitivity is strongly supported by our recently reported results that VIN coupled to  
314 magnetic particles could achieve ~25x higher binding capacity of antibodies as compared to  
315 current industry standards for affinity protein capture with magnetic particles on a per unit mass  
316 basis<sup>7</sup>.

317 Future works to further exploit the advantages of the highly tractable silica sol-gel chemistries for  
318 more sophisticated geometries and fine-tuned pore architecture would be valuable. For example,  
319 a previous study showed synthesis of bimodal pore distribution silica sol-gel in monolithic  
320 nanoflow columns in which the smaller pores are used to entrap the ligand to functionalize the  
321 column and the larger pores promote bulk flow and diffusion of the target molecule to the ligand  
322 active site<sup>26</sup>. This system architecture could be valuable for developing scale-down VBN-based  
323 bioseparation or biosensing technologies.

324

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334

335 The illustrations were created in part using Biorender.com.

336

## 337 **6. Disclosure Statement**

338 The authors have no competing interests to report.

339

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