

Folic Acid-Mediated Targeting of Cowpea Mosaic Virus Particles to Tumor Cells

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SUMMARY

Cowpea mosaic virus (CPMV) is a well-characterized nanoparticle that has been used for a variety of nanobiotechnology applications. CPMV interacts with several mammalian cell lines and tissues in vivo. To overcome natural CPMV targeting and redirect CPMV particles to cells of interest, we attached a folic acid-PEG conjugate by using the copper-catalyzed azide-alkyne cycloaddition reaction. PEGylation of CPMV completely eliminated background binding of the virus to tumor cells. The PEG-folate moiety allowed CPMV-specific recognition of tumor cells bearing the folate receptor. In addition, by testing CPMV formulations with different amounts of the PEG-FA moiety displayed on the surface, we show that higher-density loading of targeting ligands on CPMV may not be necessary for efficient targeting to tumor cells. These studies help to define the requirements for efficiently targeting nanoparticles and protein cages to tumors.

INTRODUCTION

The ability to target tumors and deliver therapeutics to specific locations in the body is a primary goal in cancer medicine. Tumor-specific targeting has the potential to reduce adverse effects and increase the therapeutic benefit of highly toxic chemotherapies [1]. Tumor-targeting strategies include the use of various types of molecular and supramolecular scaffolds such as liposomes [2], iron oxide nanoparticles [3], silica-gold nanoshells [4], highly branched macromolecules such as dendrimers [5, 6], protein cages [7], and viruses [8–11]. Ligands capable of targeting tumors such as antibodies, peptides, or small molecules are typically attached to the exterior surface. In many cases, drugs or contrast agents can also be encapsulated inside the particles in order to combine targeted cell killing with intravital imaging [12–16].

Recently viruses and protein cages have been studied in the development of tumor-specific targeting strategies [8, 17]. These multivalent protein assemblies have the advantages of simple production, monodisperse size and structure, and surfaces that are compatible with circulation in vivo [10, 18–20]. One such particle is cowpea mosaic virus (CPMV), a plant virus that grows in the common cowpea plant (*Vigna unguiculata*) and forms a 31 nm diameter icosahedral structure [21, 22]. CPMV grows to very high yields in infected plants, and the purification is simple and inexpensive. In addition, CPMV is nonpathogenic for humans, and the products derived from plant virus culture are not contaminated with animal cells or viruses [23, 24]. CPMV particles are highly stable to extremes of temperature, pH, and a variety of organic solvents such as DMSO [21, 25].

Surface modification of CPMV particles for targeting purposes has been performed by genetic and chemical means [26–31]. Chemical modification of CPMV surface lysine residues using fluorescent dye-labeled N-hydroxysuccinimide (NHS) esters or sulfhydryl groups using maleimide and haloacetamide reagents has been extensively characterized [29, 32]. Surface carboxylate groups may also be modified for specific attachment [33]. Recently, CPMV particles were shown to be stable to the conditions of the copper(I)-catalyzed azide-alkyne cycloaddition (CuAAC) reaction, dramatically increasing the variety of ligands that may be efficiently conjugated to the capsid surface [34, 35].

Although it is a plant pathogen and does not replicate in animal cells, CPMV has been shown to interact with a variety of cell types both in vitro and in vivo [17, 36]. Fluorescently labeled CPMV interacts with vascular endothelial cells when injected intravenously, allowing for a clear resolution of vascular structures in a variety of organs, while PEG-coated CPMV does not [36]. Interestingly, CPMV interacts specifically with a 54 kD membrane protein found in mammalian cells, and this interaction is also blocked by PEG coating of the virus [37]. These studies demonstrate that CPMV has significant potential for use in targeting and imaging of both normal vasculature and tumors in vivo but also underscores the need for cell-specific retargeting of the capsid.

One of the best-characterized ligands to be exploited for targeting tumor cells is the vitamin folic acid (FA). FA plays an essential role in human growth and development and at the cellular level in cell division and DNA synthesis [38]. Uptake of FA into cells is mediated by the folate receptor (FR), and binding of FA to FR initiates receptor-mediated endocytosis and internalization of FA [39, 40]. Normally expression of FR on cells is low; however, the demand for FA increases during cellular activation and proliferation. Indeed, FR expression is upregulated on a variety of human tumors, including cancers of the ovary, kidney, uterus, testis, brain, lung, and myelocytic blood cells [12, 41–43]. Upregulation of FR has also been correlated with poor disease outcome [44]. Thus, FA has been studied as a promising targeting ligand for cancer detection, imaging, and treatment [45].

Several nanoparticle formulations have used FA-based targeting strategies. The derivatization of liposomes with FA has been the most intensively investigated thus far [13, 14, 46]. Other nanoparticle formulations employing FA-targeting include FA-coated gadolinium nanoparticles that were shown to increase uptake in tumors, suggesting the possibility of neutron capture therapy [47], poly-amidoamine dendrimers (PAMAM) conjugated to FA [6, 48], and core-shell nanostructures decorated with FA [49]. The FA-targeting strategy has also been applied to gold nanoparticles [50], pegylated micelles [51], and adenoviruses [10] in an attempt to reach FR-expressing cells. PEG spacers have also been used to facilitate binding of the conjugated FA to cell-surface receptors [14, 51–54]. Nevertheless, precise control of attachment of targeting ligands at the nanoscale, as well as establishment of the relationship between ligand density and target cell recognition, has not been achieved.

Several attached large molecules such as peptides, antibody ligands, tumor receptor ligands, and transferrin, have been used to target viral nanoparticles and protein cages [29, 35, 55], but the efficiency of cellular targeting using capsid-arrayed small molecule ligands has been less well studied. We report here the chemical attachment of FA, PEG, and PEG-conjugated FA to CPMV by using the CuAAC reaction in order to direct the capsid specifically to tumor cells. The ability of the PEG molecules to inhibit natural CPMV-cell surface interactions and the ability of FA-modified CPMVs to bind to tumor cells was studied by confocal microscopy and flow cytometry-based binding assays. In addition, CPMV-based formulations that kept the amount of PEG constant and varied the loading of the FA ligand were evaluated for their ability to interact with FR-expressing cells in order to test the hypothesis that polyvalent binding might be advantageous for cell-specific targeting of CPMV.

RESULTS

Previous studies have shown that CPMV recognizes and is taken up by a variety of mammalian cell types [36, 37]. To first ask whether CPMV conjugated directly to FA could redirect this natural targeting, an NHS ester of FA was

conjugated to surface lysines on CPMV. A parallel reaction performed under identical conditions with fluorescein-NHS resulted in the attachment of 100 ± 10 dyes per particle; we assume that the analogous folate conjugation gave a similar result [56, 57]. The natural uptake of CPMV is significant [36], and CPMV-FA particles showed no measurable increase in binding to FR-expressing KB cells in comparison to unmodified CPMV particles as measured by flow cytometry binding analysis and cell uptake studies (data not shown). Thus, the natural cellular interactions of CPMV needed to be modified in order to achieve FR-specific targeting.

We previously showed that PEG (3,400 Da) inhibits the natural targeting and uptake of CPMV [36]. In addition, the importance of a PEG spacer for efficient cellular recognition of FA-conjugated nanoparticles has also been previously noted [2]. To accommodate tailored loading of both PEG and FA on CPMV particles, a relatively short (500 Da) PEG chain was introduced as shown in Figure 1. A monodisperse azide-PEG-amine was coupled to the NHS ester of folic acid to give the FA-azide **2a**. The corresponding polyvalent alkyne nanoparticle was obtained by reacting CPMV with a large excess of the alkynyl NHS ester reagent **1**. This procedure has been shown to acylate 150–200 of the exposed lysine side chain amines of CPMV [56] and was chosen to provide an excess of available surface alkyne groups for subsequent attachment. The use of the CuAAC ligation reaction eliminates the need for protecting group manipulations and allows for reproducible attachments to be achieved with a minimum amount of material.

FA derivatives are known to form aggregates at high concentrations [58], and there are some indications that too many FA molecules displayed on a surface can inhibit binding to the target receptor, perhaps because of such aggregation [59]. To address this phenomenon, we limited the density of PEG-FAs on CPMV by using a relatively low concentration of azide derivatives in the CuAAC bioconjugation step. An estimate of the loading was obtained by performing a parallel reaction by using N_3 -PEG-fluorescein (**2b**) with CPMV-alkyne under identical conditions, as we have previously described [57]. This transformation provided a value of 60 ± 6 molecules per virus for the attachment of PEG-Fluorescein or PEG-FA, where a 1 mg/ml CPMV solution translates to a folic acid concentration of 10 μ M.

The integrity of the conjugated virus preparations was verified by analytical size exclusion chromatography (Superose 6), which also showed by a slightly smaller elution volume that the CPMV-(PEG-FA)₆₀ conjugate appears to be slightly larger than unmodified CPMV (Figure 2A). It is possible that the early elution of CPMV-(PEG-FA)₆₀ could also be caused by altered interaction of PEG-FA moiety with the column material in comparison to the unmodified particle (designated CPMV-WT). The sharpness of the peaks for CPMV-(PEG-FA)₆₀ in comparison to CPMV-WT suggests that the modification of the particles within the population is relatively uniform on a per-particle basis. This uniformity was supported by transmission electron

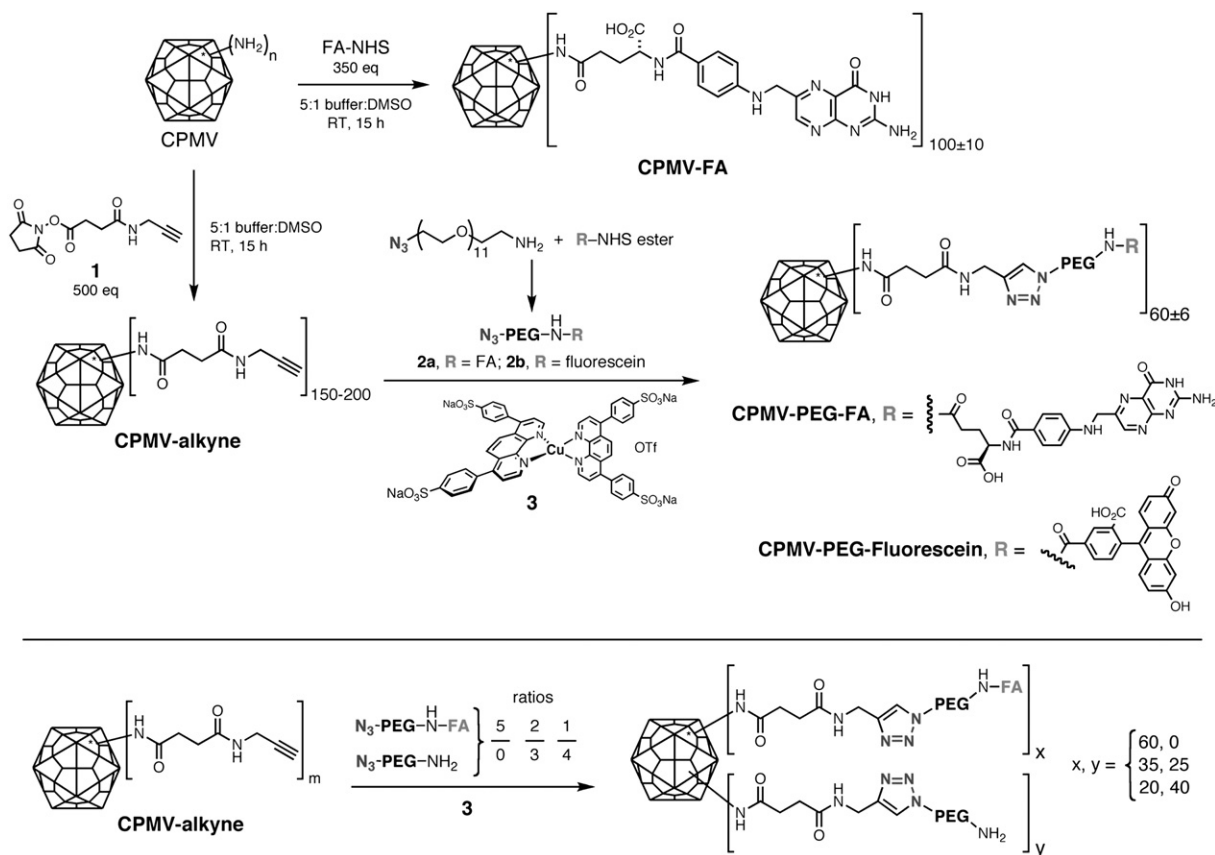


Figure 1. Synthesis of the PEG-Folic Acid Conjugates

Scheme for synthesis of FA-conjugated CPMV.

microscopy (TEM) studies (Figures 2E and 2F) of negatively stained samples, which confirmed that the CPMV-(PEG-FA)₆₀ particles were intact and showed no evidence of aggregation. Visual analysis of the samples showed that 100% of the CPMV-(PEG-FA)₆₀ particles exhibited a halo-like structure on the surface of the particle that appeared to be uniform within the population and which was absent in unmodified CPMV (Figures 2E and 2F).

Successful PEG-FA conjugation was further confirmed by gel electrophoresis and western immunoblotting. Figure 2B shows the large (L) and small (S) capsid proteins of wild-type and PEG-FA-derivatized CPMV; unlike larger PEG chains used in previous studies [60], the small PEG-FA moiety used here shifted the positions of the fully denatured S or L bands only slightly toward higher molecular weight. Western immunoblots appear in Figures 2C and 2D, by using higher concentrations of capsid protein. The use of anti-CPMV antibody showed the L and S subunits as well as minor amounts of higher molecular weight species that arise from incomplete denaturation of the durable CPMV capsid (Figure 2C). After attachment of PEG-FA, the higher molecular weight bands increased in intensity, presumably because the FA-derivatized protein subunits were more prone toward aggregation or resistant to denaturation (Figure 2C). Anti-FA antibody showed all of

the bands from CPMV-(PEG-FA)₆₀ to bear FA (Figure 2D). Of the five surface lysines found on CPMV, two dominate the acylation reactivity of the particle—one each on the S (lysine 38) and L (lysine 99) subunits [56, 61]. The average of 60 PEG-FA chains per particle may therefore be assumed to be distributed largely among the 120 sites provided by these two residues on the capsid exterior.

The ability of the relatively short (500 Da) PEG chains to inhibit the natural targeting of CPMV was assessed by flow cytometry. Two tumor cell lines, KB and HeLa, were used for these studies. KB, a human nasopharyngeal carcinoma cell line, has been widely used in tumor-targeting studies due to its high expression of FR and affinity for FA [42, 45, 48, 62–64]. HeLa cells, a well-established epitheloid cervical carcinoma, have been reported to express less FR than KB cells. The cellular uptake of CPMV particles in KB or HeLa cells after 2 hr incubation at 37°C was analyzed by fluorescence microscopy (Figure 3). As previously described, unmodified CPMV showed significant uptake in HeLa and KB cell lines (Figures 3A and 3D), while the decoration of CPMV with PEG inhibited this phenomenon (Figures 3B and 3E). Both types of cells also showed increased uptake of FA-conjugated viruses compared to unmodified CPMV (Figures 3C and 3F). Interestingly, uptake of CPMV-(PEG-FA)₆₀ in HeLa cells appeared as an

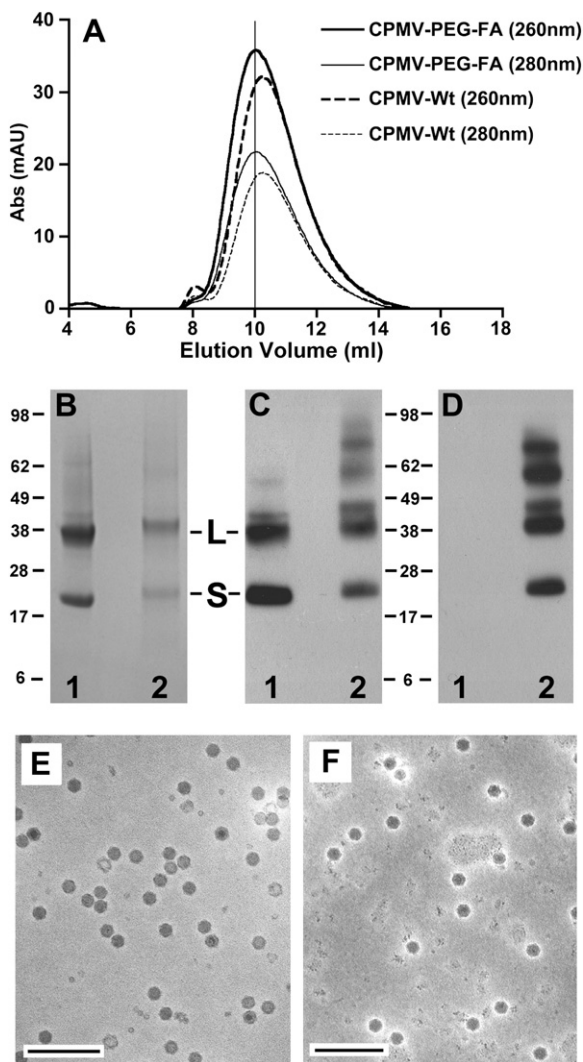


Figure 2. Virus Integrity Assessed by Size Exclusion Chromatography, Western Blot, and TEM

(A) Size exclusion FPLC analysis of CPMV-PEG-FA and WT-CPMV. (B) Coomassie stain of WT-CPMV (1) and CPMV-(PEG-FA)₆₀ (2), showing small (S) and large (L) capsid subunit proteins. (C–D) Western blots of WT-CPMV (1) and CPMV-(PEG-FA)₆₀ at higher concentration (2). Left panel, detection using rabbit anti-CPMV antibody. Right panel, detection using mouse anti-folic acid antibody. L, large subunit; and S, small subunit. (E–F) Typical TEM image of WT-CPMV (E) and CPMV-(PEG-FA)₆₀ (F), showing intact particles. The bar represents 200 nm.

evenly dispersed fluorescence in the cytoplasm, while CPMV uptake in KB cells showed a more punctate distribution of the virus throughout the cells. CPMV is naturally targeted to lysosomes [36], although the endocytic pathway used to reach this compartment is not well defined. The uptake of FR in either cell type is not well defined and has been shown to involve both clathrin-mediated and caveolin-mediated endocytic pathways. Differences between uptake patterns in each cell type could be due to differences in receptor expression level (e.g., saturation of the clathrin-mediated pathway in HeLa cells spilling into the

caveolae pathway or vice versa), or due to differences in the localization of endocytic vesicles or the trafficking pattern of vesicle recycling between the two cell types [63, 65, 66]. Nevertheless, our data indicate that the endocytic pathway used for uptake and trafficking of CPMV-PEG-FA is modified in comparison to CPMV-WT.

Differences in cell-surface binding were measured by flow cytometry, where virus binding and subsequent steps were carried out at 4°C in order to minimize endocytosis. CPMV-(PEG-FA)₆₀ bound to KB cells much better than to HeLa cells (94% and 22%, respectively, relative to cells alone) (Figures 4A and 4B, solid histograms), and both cell types interacted hardly at all with CPMV-PEG lacking the terminal FA moiety (1.6% and 0.7% to KB and HeLa cells, respectively) (Figures 4A and 4B). These results are consistent with the difference in the expression of FR receptors on the two cell types (approximately 2.8×10^5 receptors per cell for KB cells versus 4-fold fewer receptors per cell for HeLa [67, 68]). Further details are provided in the Supplemental Data and Figure S1 available with this article online.

To further show that the interaction between CPMV-PEG-FA and tumor cells was due to specific interactions between the particles and FR on cells, the ability of free folic acid to compete for binding of CPMV-(PEG-FA)₆₀ to KB cells was assessed by flow cytometry. In the experiment shown in Figure 5A, free folic acid at greater than 10 μM substantially inhibited CPMV-(PEG-FA)₆₀ binding, and less than 1 μM folic acid had no effect. CPMV-(PEG-FA)₆₀ was used at a concentration of 17 nM in particles, or 1 μM total FA. Nonlinear regression analysis (Prism, GraphPad, Inc., San Diego, CA) of a representative experiment revealed an IC₅₀ value of 2.75 μM. The average IC₅₀ of four separate experiments was 4.9 μM (Figure 5A). If each folate unit on CPMV acted as a free folate molecule and each virus-cell contact was made by a single FA-FR interaction, the IC₅₀ value would, by definition, be equal to the virus-displayed FA concentration of 1 μM. Thus, FA displayed on the capsid surface binds with similar affinity on a per-folate basis than free folic acid. On a per-particle basis, the avidity of CPMV-(PEG-FA)₆₀ for KB cells is much stronger (by approximately 300 times) than free FA.

To further explore whether controlling FA loading at the nanoscale could influence interaction with FR, three different versions of CPMV-PEG-FA were formulated that had an equal loading of PEG but varied in the number of FA units at the end of the PEG chains. Separate portions of CPMV-alkyne were reacted with azide-PEG-FA and two different combinations of azide-PEG-FA and azide-PEG-amine, giving rise to the attachment of 60, 35, and 20 PEG-FA units per particle (Figure 1, bottom). These are designated CPMV-(PEG-FA)₆₀, CPMV-(PEG-FA)₃₅, and CPMV-(PEG-FA)₂₀, respectively. Comparative western blots using identical amounts of virus particles confirmed the relative loading values by Spot Density Analysis (Alpha Innotech, San Leandro, CA) (Figure S2A). While little difference in the cell uptake of each of these particles was observed at saturating conditions, titration of binding

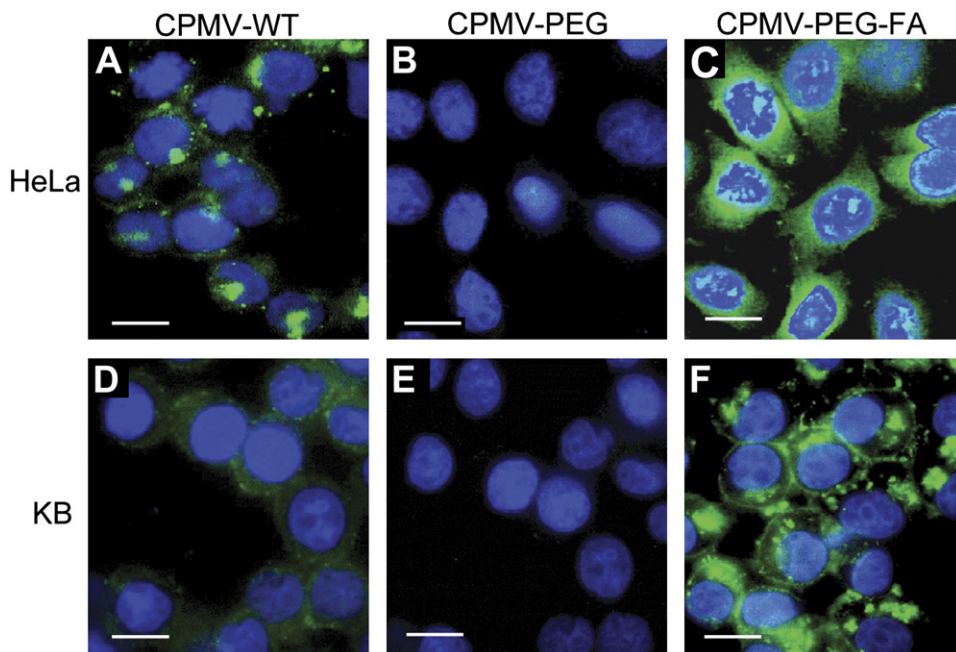


Figure 3. Fluorescent Microscopy of Monolayers of HeLa and KB Cells

Cells were incubated with the indicated virus particles (37°C, 2 hr), followed by permeabilization and treatment with anti-CPMV antibody and then with a green fluorescent secondary antibody conjugate. Nuclei are stained with DAPI (blue). Top, HeLa cells; bottom, KB cells. Scale bar, 10 μ m.

revealed a clear difference in binding that correlated with the loading of FA per particle (Figure S2B). Moreover, in a competition assay using free FA to inhibit binding, in each case, approximately the same concentration of free FA was required to disrupt binding of the particles to KB cells as was presented on the virus surface (Figure 5B). In other words, in no case was a substantial per-folate polyvalent effect observed, but in all cases, the folate-decorated particles bound substantially better than free folic acid molecules alone.

DISCUSSION

In this work, we demonstrate the specific binding and endocytosis of a FA-derivatized CPMV particle by cells bearing FR. Direct, controlled chemical conjugation of a monodisperse PEG-FA moiety to the CPMV surface by copper (I)-catalyzed azide-alkyne cycloaddition (CuAAC) was efficient and resulted in the attachment of a maximum

of 60 ± 6 molecules of FA per particle. Wild-type CPMV exhibited significant binding, which was eliminated by the conjugation of the relatively short PEG chains to the virus surface. The addition of the FA moiety then allowed the particles to be recognized specifically by the tumor cell lines. Modulating the extent of FA loading using the CuAAC reaction demonstrated excellent control of particle modification at the nanoscale and correlated with tumor cell recognition. These studies suggest that CPMV nanoparticles can be effectively redirected by surface conjugation to ligands of interest, allowing specific uptake into tumors while avoiding nonspecific uptake into normal cells.

At least two factors may contribute to the inability of HeLa or KB cells to specifically recognize FA directly conjugated to the CPMV capsid (as opposed to FA being displayed on the end of PEG chains). The simplest explanation is that endogenous and potentially multivalent CPMV-cell surface interactions are of higher affinity than

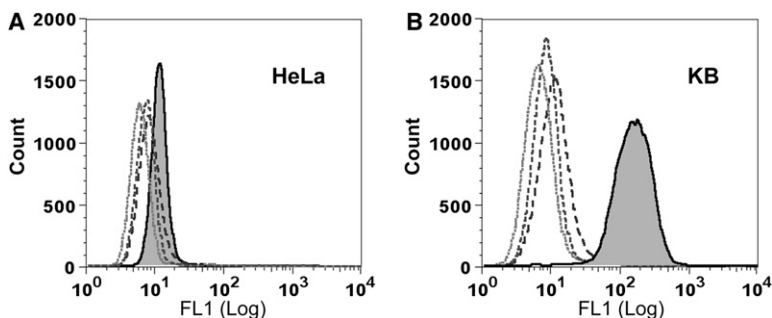


Figure 4. Measurement of Virus Binding to Tumor Cells Using Flow Cytometry

(A) Binding of CPMV to HeLa tumor cells. (B) Binding of CPMV to KB tumor cells. The peaks from left to right are cells only (dotted histogram), CPMV-PEG (short dashed lines), WT-CPMV (long dashed lines), and CPMV-(PEG-FA)₆₀ (filled histograms), respectively.

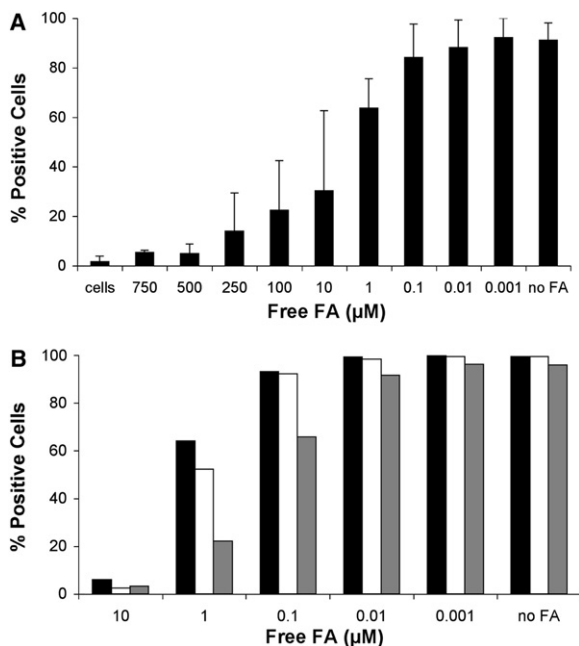


Figure 5. Competition of CPMV-PEG-FA Using Free FA

(A) CPMV-PEG-FA bearing 60 FA/particle (1 μM FA) was incubated with KB cells in the presence of increasing amounts of FA as indicated. Error bars represent the mean ± SD of triplicate experiments.

(B) Competition of CPMV-(PEG-FA)₂₀ (gray), CPMV-(PEG-FA)₃₅ (white bars), or CPMV-(PEG-FA)₆₀ was performed with increasing amounts of FA as indicated. In each case, the concentration of virus particles was 17 nM, giving rise to the following concentrations of displayed FA: CPMV-(PEG-FA)₂₀ = 0.33 μM FA; CPMV-(PEG-FA)₃₅ = 0.6 μM FA; CPMV-(PEG-FA)₆₀ = 1 μM FA.

the FA-FR interaction. Second, the conjugated FA units may simply be rendered inaccessible to FR by virtue of being attached to the rigid capsid structure by a short tether; previous reports have noted steric impediments to binding in other settings [14, 52, 53]. Alternatively, if endocytosis is initiated by multivalent binding, rather than a single-point contact (as has long been known for certain carbohydrate receptors [69, 70]), there may be a minimum distance that adjacent FAs need to achieve before such a process is initiated. The PEG linker can address both potential deficiencies, and studies are in progress to attempt to distinguish between these possibilities. It should be noted that the PEG linker used here is significantly shorter than has been previously employed for FA nanoparticles [52, 53, 64].

Our studies coupling 20–60 PEG-FA moieties to CPMV resulted in targeting that compares favorably to other FA-conjugated nanoparticle formulations. For example, conjugation of FA to PAMAM dendrimers resulted in loading of about 2.5 molecules of FA residues per dendrimer, due to an undesired reaction occurring between the carboxylic acid groups of FA and the amine groups of the dendrimer. In this case, no binding enhancement was observed: a concentration of 30 nM of free FA reduced by 50% the amount of association to KB cells reached by 30 nM of the dendrimer-FA conjugate [5, 48]. Coupling

of FA to 10 nm Au nanoparticles has also been reported [50], but only 20% of the KB cells exposed to the nanoparticles actively took up the FA-conjugate. This uptake could be suppressed by 2 μM free FA.

PEGylation of CPMV, even with the small 500 Da material here, resulted in effective suppression of the particle's natural cellular interactions. The strategy has been used previously in many other studies by our groups and others. For example, multifunctional PEGylated micelles conjugated to FA and carrying the antitumor drug adriamycin showed significant increase in cytotoxicity compared to nontargeted micelles. However, longer incubation times also induced significant cytotoxicity when nontargeted micelles were used, suggesting that PEGylation was not sufficient in blocking nonspecific binding of the micelles to tumor cells [51]. Attempts have also been made to retarget adenovirus particles to KB cells [10]. The adenovirus surface was modified with FA-PEG conjugate (24% occupancy), resulting in a ~40% increase in transfection efficiency over nontargeted pegylated virus. Nevertheless, the pegylated adenovirus was still able to transduce cells via its natural coxsackie-adenovirus receptor (CAR), suggesting that the degree of PEGylation was not sufficient to effectively retarget the virus.

One drawback in most chemical nanoparticle formulations is the lack of control over the spatial distribution on the outer shell of the ligand of interest. A sophisticated rational design of ligand orientation and stoichiometry would allow for the creation of a polyvalent scaffold that enhances the targeting properties of the nanoparticle and offers the possibility of cell targeting with lower concentrations of targeting molecules than would be possible for the monomeric form [71]. The increased binding of CPMV-PEG-FA over free FA (factor of approximately 2 per FA, or 120 per particle) is not sufficient to clearly indicate the existence of polyvalent (simultaneous multipoint) interactions [72] with FR, compared to other known examples using carbohydrates and cell surface lectins [73–77] or liposome-displayed anthrax peptides and receptors [78]. In part, this may be due to the nature of the ligand-receptor pair, since more substantial avidity effects were observed by using CPMV displaying carbohydrate moieties [28, 79] (E. Kaltgrad and M.G. Finn, personal communication). Interestingly, a substantial polyvalent effect has recently been quantified by SPR measurements of binding affinity for FA displayed on dendrimers [78, 80]. While the more flexible nature of the dendrimer scaffold may better match cellular receptor patterns, as has been previously hypothesized [78], the target is also different (protein immobilized on SPR chips versus cell-surface receptors), and competition with free monovalent folate was not reported in the dendrimer case [80]. For CPMV-PEG-FA, the roughly linear trend of free FA inhibition with variation of PEG-FA loading further suggests that monovalent FA-FR interactions are dominant but also that local concentration of the FA ligand plays a role in binding of cells to the polyvalent scaffold. It seems likely that the effective off-rate of the particle is slowed by facile reassociation with FR receptors brought about by the high local concentration of

FA on the particle surface. Such effects may be all that is necessary to provide enough increased avidity and endocytosis to allow for effective tumor targeting [81].

Prior attempts to target tumors with displayed FA have yielded mixed results depending on the tumor site [47, 82, 83]. Endothelial cell targeting has also been widely studied in order to reach the tumor vasculature [15, 84]. Thus, a combination of targeting moieties could enhance the specificity of nanoparticles to tumors in vivo. Toward these ends, the construction and in vivo testing of more sophisticated particles, taking advantage of the precise control of amino acid side-chain positioning and reactivity to install PEGs and tethered targeting moieties, is in progress in our laboratories.

SIGNIFICANCE

The ability to specifically target therapeutics and imaging modalities to tumors is an important goal in biomedicine. Viruses and protein cage nanoparticles have proven utility in this area, however many of these materials have natural cellular interactions. CPMV in particular provides tremendous imaging resolution of normal and tumor vasculature; however, the particles have significant affinity for cell-surface proteins. Thus, any natural in vivo interaction of the nanoparticle must be masked and the particle retargeted to a tumor ligand of interest. In this article, we use copper (I) catalyzed azide-alkyne cycloaddition chemistry to precisely attach a defined number of folic acid tumor ligands, while redirecting the natural specificity of the particles by coating the remaining surface of the particles with low molecular-weight polyethylene glycol. The resulting particles are effectively retargeted specifically to the folic-acid receptor. The density of ligands may be controlled at the nanoscale, and competitive binding versus free folic acid was shown to track linearly with the density of folate attached to the nanoparticle scaffold. The high local concentration of folic acid on the capsid surface enables these particles to be targeted efficiently to cell-surface FR receptors. To our knowledge, this is the first time that small-molecule ligands have been successfully used to provide targeting specificity for virus-based nanoparticles.

EXPERIMENTAL PROCEDURES

All reagents, unless otherwise specified, were purchased from commercial suppliers and used without further purification. Bifunctional N_3 -PEG-NH₂ [$N_3(CH_2CH_2O)_{11}CH_2CH_2NH_2$, MW 570, PDI = 1.0] was obtained from Polypure, Inc. Copper(I) triflate was prepared according to literature procedures [85].

Propagation of CPMV in Plants

The primary leaves of Kentucky cowpea plants (*Vigna unguiculata*) were mechanically inoculated as 10 day old seedlings, bearing two primary leaves and with secondary leaves just beginning to show. Virus stocks were initiated from pCP1 and pCP2 plasmid cDNAs encoding full-length copies of the two RNA moieties of CPMV, RNA-1, and RNA-2, respectively. Carborundum was first dusted onto the leaves to aid in the wounding process. At approximately 3 weeks post inocu-

lation, the symptomatic leaves were harvested, weighed, and frozen at -70°C until virus purification. The virus was purified from the infected leaves by a method described elsewhere [86].

Synthesis of N_3 -PEG-FA

FA-NHS ester was prepared from FA according to literature procedures [2]. FA-NHS (100 mg, 0.19 mmol) in anhydrous DMSO (10 ml) was treated with N_3 -PEG-NH₂ (50 mg, 0.09 mmol), and the mixture was stirred while warming gently (ca. 35°C) for 20 hr. DMSO was removed under high vacuum at RT. The residue was treated with CH_2Cl_2 , and the solids were scraped gently from the walls of the flask. The mixture was sonicated to form a slurry, which was transferred to a fritted disk, and washed thoroughly with diethyl ether, CH_2Cl_2 , and THF. The residual dark orange solid was dried and extracted with water until the aqueous extract was colorless. The aqueous fractions were combined and evaporated under vacuum to yield N_3 -PEG-FA (compound 2) as viscous orange oil (60 mg, 0.06 mmol, 66%). ^1H NMR (200 MHz, D₂O): δ 8.59 (s, 1H), 7.52 (d, 2H), 6.64 (d, 2H), 4.48 (s, 1H), 4.15 (m, 2H), 3.58–3.35 (m, 40), 2.35–2.20 (m, 12H). An analogous procedure was employed for the synthesis of N_3 -PEG-fluorescein.

Preparation of CPMV-Alkyne

Compound 1 (50 mg, 0.20 mmol) was dissolved in DMSO (1 ml) and added to a solution of wild-type CPMV (5 ml, 7 mg/ml, 75 μM in protein subunits, 0.1 M phosphate buffer [pH 7.0]). The mixture was gently agitated at RT for 15 hr and purified by sucrose gradient fractionation (10%–40% sucrose in 0.1 M [pH 7.0] phosphate buffer, Beckman SW-28 Ti rotor, 28000 rpm, 3 hr). The intact virus was collected as a pale white band under intense illumination on a gradient fraction collector and subjected to ultracentrifugation (Beckman 50.2 Ti rotor, 42000 rpm, 3 hr) to form a colorless pellet. The solution was decanted and the colorless pellet was dissolved under N_2 with sufficient buffer (Tris-Cl, 0.1 M [pH 8.0]) to obtain a concentration of 7.3 mg/ml.

Conjugations to CPMV-Alkyne

The following reaction protocol was carried out in an inert atmosphere (N_2) glove box with O_2 level kept below 6 ppm until the final gel filtration step. For each of the three N_3 -PEG-FA loading levels, a 2 ml Eppendorf centrifuge tube was charged with CPMV-alkyne (7.3 mg/ml, 110 μl) and buffer (Tris-Cl, 0.1 M [pH 8.0], 260, 250, 220 μl , respectively). Degassed aqueous solutions of N_3 -PEG-FA (25 mM; 10, 20, 50 μl , respectively) and N_3 -PEG-NH₂ (25 mM; 40, 30, 0 μl , respectively) were added simultaneously to the virus solution and mixed by gentle agitation. Thus, the total azide and PEG concentrations were kept constant for each of the three reactions, and the amount of FA versus amine caps on the PEG chains were varied. A solution of copper(I) triflate (100 mM, CH_3CN , 50 μl) was combined with a solution of sulfonated bathophenanthroline (100 mM in 100 mM Tris-Cl [pH 8.0], 100 μl) (We subsequently learned that Tris-Cl slightly inhibits the CuAAC reaction relative to HEPES buffer, and so the latter is now recommended) to form a catalyst of the formulation 3. An aliquot of the catalyst mixture (16 μl) was added to the tube containing the virus. The reaction mixture was immediately placed on a rotisserie for continuous agitation, and kept under N_2 at RT for 15 hr. The product (designated CPMV-PEG-FA) was purified by three passages through size exclusion gel-filtration columns (BioRad, p-100), which removed all residual catalyst and excess N_3 -PEG-FA. The integrity of the virus was verified by analytical size exclusion chromatography (Superose 6) and transmission electron microscopy (TEM, samples stained with 0.2% uranyl acetate; images acquired on a 100 Kv Philips Technai electron microscope). Conjugations of N_3 -PEG-NH₂ and N_3 -PEG-Fluorescein to CPMV-alkyne were performed in the same manner by using the following reagents: CPMV-alkyne (7.3 mg/ml, 110 μl), buffer (Tris-Cl, 0.1 M [pH 8.0], 250 μl), N_3 -PEG-NH₂, or N_3 -PEG-Fluorescein (25 mM, 20 μl). Virus concentrations were determined by UV-vis spectroscopy: at 0.1 mg/ml, CPMV gives a standard absorbance of 0.8 at 260 nm. The average molecular weight of the CPMV virion is 5.6×10^6 .

Synthesis of CPMV-FA

FA-NHS (31 mg, 58 μ mol) was dissolved in DMSO (1 ml) and added to a solution of wild-type CPMV (5 ml, 3 mg/ml, 32 μ M in protein subunits, 0.1 M phosphate buffer [pH 7.0]). The mixture was gently agitated at RT for 15 hr and purified by sucrose gradient fractionation (10%–40% sucrose in 0.1 M [pH 7.0] phosphate buffer, Beckman SW-28 Ti rotor, 28000 rpm, 3 hr). The intact virus was collected as a pale white band under intense illumination on a gradient fraction collector and subjected to ultracentrifugation (Beckman 50.2 Ti rotor, 42000 rpm, 3 hr) to form a colorless pellet. The solution was decanted, and the colorless pellet was dissolved with sufficient buffer (Tris-Cl, 0.1 M [pH 8.0]) to obtain a concentration of 5.0 mg/ml. CPMV-FA was characterized in the manner described above for CPMV-alkyne + azide conjugates.

Western Blotting

To detect CPMV, protein bands were visualized by Western blotting with anti-CPMV IgG (purified from rabbit polyclonal antisera on a protein G column; Amersham Pharmacia, Uppsala, Sweden) and diluted 1:1000 in blocking buffer (PBS, 5% normal goat serum, 0.05% Tween20). ImmunoPure goat anti-rabbit IgG, peroxidase conjugated, diluted 1:20,000 (Pierce, Rockford, IL) was used as secondary antibody. Folic acid conjugated to CPMV was detected by western blotting with mouse anti-Folic acid monoclonal antibody (Sigma-Aldrich, St. Louis, MO) diluted 1:1000 in blocking buffer (PBS, 5% normal goat serum, 0.05% Tween20). Goat anti-mouse peroxidase conjugated, diluted 1:20,000 (Pierce, Rockford, IL) was used as secondary antibody. In both cases, detection of peroxidase was carried out by using SuperSignal West Pico Chemiluminescent Substrate (Pierce, Rockford, IL).

Cell Culture and Binding Studies

HeLa cells and KB cells, the latter a human nasopharyngeal epidermal carcinoma [62], were grown continuously as a monolayer by using FA-free RPMI1640 medium (GIBCO, Invitrogen, Carlsbad CA) containing 10% heat-inactivated fetal bovine serum (FBS), penicillin (50 units/ml), streptomycin (50 μ g/ml), and 2 mM L-glutamine at 37°C in a 5% CO₂/95% air humidified atmosphere. The concentration of folic acid was 5–6 nM in FA-free medium containing serum, close to the natural physiologic concentration (5–30 nM) [52, 87]. On the day before each experiment, the medium was replaced with FA-free RPMI 1640 containing all the supplements mentioned above, except 10% FBS.

Measurement of Virus Binding to HeLa and KB Cells Using Flow Cytometry

HeLa and KB cells, grown overnight in FA-depleted medium, were treated with trypsin, counted, and distributed in 100 μ l portions into a 96-well V-bottom shaped plate at a concentration of 5 \times 10⁵ cells/ml. Ten micrograms of different virus preparations were added to each well and the cells were incubated on ice at 4°C for 1 hr. The cells were then washed four times with ice-cold PBS buffer containing 1 mM EDTA and 25 mM HEPES (pH 7.5), centrifuged at 1,600 rpm for 6 min at 4°C. Rabbit anti-CPMV primary antibody was then added to the cells in a 100 μ l volume, and the cells were incubated on ice at 4°C for 30 min and washed as above. The same procedure was then repeated with goat anti-rabbit IgG AlexaFluor 488 conjugated antibody (Invitrogen, Carlsbad, CA), with incubation conducted in the dark. Finally, the cells were fixed with 2% formaldehyde in PBS buffer containing 1 mM EDTA and 25 mM HEPES (pH 7.5). The samples were then analyzed with a FACS Calibur instrument (BD Biosciences, Franklin Lakes, NJ). Approximately 40,000 events were collected for each sample, and the data were analyzed with FlowJo software (Tree Star, Inc, Ashland, OR). Binding and competition data were further analyzed by nonlinear regression analysis (Prism, GraphPad, Inc.). Independent control experiments established that the PEG-conjugated CPMV was recognized with approximately the same efficiency as native CPMV by the anti-CPMV antibody.

Cellular Uptake of the FA-Conjugated Virus in HeLa and KB Cells

Cells were seeded in a 12-well plate containing 12 mm sterile glass coverslips at 1 \times 10⁵ cells/well and grown for 48 hr as previously described. On the day of the experiment, cells were washed once with FA-depleted medium, and 10 μ g of different virus preparations were added to each well, and the cells were incubated at 37°C in a 5% CO₂/95% air humidified atmosphere for 2 hr. Cells were then washed four times with FA-depleted medium to remove unbound virus, on a rocker at room temperature for five minutes. Cells were then fixed with 4% paraformaldehyde in PBS for 20 min. After four washes with PBS buffer, cells were permeabilized with 0.1% Triton X-100 in PBS for 15 min. Nonspecific binding was blocked by incubating the cells in 5% goat serum in PBS for 1 hr. Rabbit anti-CPMV antibody was added to the cells in 1% goat serum, 0.1% Triton X-100 in PBS, and cells were incubated at room temperature for 45 min with gentle agitation. Unbound antibody was then removed by washing four times with PBS. Goat anti-rabbit IgG AlexaFluor 488 conjugated antibody (Invitrogen) was added in 1% goat serum in PBS, and cells were gently agitated for a further 35 min. During the last 5 min of secondary antibody incubation, cell nuclei were stained by adding 100 μ l of 4',6-diamidino-2-phenylindole (DAPI 1:1000 dilution in water). Cells were then washed four times with PBS, and coverslips covered with cells were mounted on slides with Vecta Shield mounting medium (Vector Laboratories). Cells were imaged by using a Nikon Eclipse TS100 microscope, with a 100 \times oil objective.

Competition Assay

KB cells, grown overnight in FA-depleted medium, were treated with trypsin, counted, and distributed in 100 μ l portions into a 96-well V-bottom shaped plate at a concentration of 5 \times 10⁵ cells/ml. Ten micrograms of CPMV-PEG-FA was added to the cells together with different concentrations of free FA, ranging from 750 μ M to 1 nM. Cells were then treated as described above, and the samples were analyzed using flow cytometry. The competition assay was also carried out with CPMV-PEG-FA that had been synthesized so that different amounts of PEG-FA conjugate were displayed on the surface of the virus.

Supplemental Data

Supplemental Data include further characterization of CPMV-PEG-FA binding, including comparisons of CPMV particles with different loadings of FA, and are available at <http://www.chembiol.com/cgi/content/full/14/10/1152/DC1/>.

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