

Distinction Between Active and Passive Targeting of Nanoparticles Dictate Their Overall Therapeutic Efficacy

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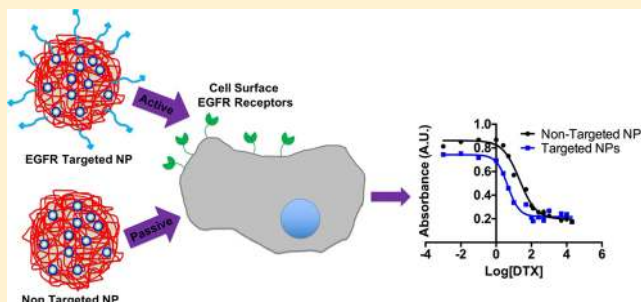
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Supporting Information

ABSTRACT: The role of nanoparticles in cancer medicine is vast with debate still surrounding the distinction between therapeutic efficacy of actively targeted nanoparticles versus passively targeted systems for drug delivery. While it is commonly accepted that methodologies that result in homing a high concentration of drug loaded nanoparticles to the tumor is beneficial, the role of intracellular trafficking of these nanoparticles in dictating the overall therapeutic outcome remains unresolved. Herein we demonstrate that the therapeutic outcome of drug loaded nanoparticles is governed beyond simply enabling nanoparticle internalization in cells.

Using two model polymeric nanoparticles, one decorated with the GE11 peptide for active targeting of the epidermal growth factor receptor (EGFR) and the other without, we demonstrate that EGFR mediated intracellular internalization results in an enhanced therapeutic effect compared to the nontargeted formulation. Our findings demonstrate that the intracellular destination of nanoparticles beyond its ability to internalize is an important parameter that has to be accounted for in the design of targeted drug delivery systems.



INTRODUCTION

Despite significant advances in cancer treatments and diagnosis, cancer still remains as one of the world's most devastating diseases with more than 10 million new cases reported every year.¹ Concomitant with these devastating cancer incidences are the side effects of chemotherapeutics still widely used for the treatment of this disease. Chemotherapeutics are the major class of drugs available in the arsenal against cancer, introduced in the 1950s for the treatment of these diseases.² However, these drugs are highly toxic and can result in severe and debilitating side effects for patients.^{3,4} As a result of this, chemotherapeutics encapsulated in nanoparticle delivery vehicles are a promising field to enhance the effectiveness of these treatments in cancer while avoiding some of these side effects.^{5–8} There is great interest in applying nanoparticle technology to cancer therapies largely resulting from the appealing features that nanoparticles possess. These include therapeutic protection from degradation, improved drug pharmacokinetics, improved intracellular penetration, selective tissue targeting, and the inclusion of imaging modalities.^{9–11} Further to this, the increasing opportunities of “theranostic” nanoparticles which can combine both diagnostic capabilities as well as therapy within

a single entity provide great promise in the fight against cancer.^{12,13}

Nanoparticles due to their unique size are able to exploit the distinct cancer pathology and its molecular biology to in turn result in higher uptake and preferential targeting of therapeutics to the tumor compared to traditional treatments.⁸ Broadly, this is achieved by two methods, “passive” or “active” targeting. Passive targeting is possible due to the unique changes in the cancer vasculature. Due to the rapid growth of tumors, blood vessels and junctions are not formed properly and can be loose and leaky. Owing to the unique size of nanoparticle formulations, they are able to pass through these loose junctions resulting in preferential accumulation at the tumor site over time. This phenomenon is known as the enhanced permeation and retention (EPR) effect.^{10,14} As a result of this and the combinatorial advantages of nanoparticle based delivery vehicles, a number of systems have been approved by the U.S. Food and Drug Administration (FDA) for chemotherapeutic delivery. These formulations include

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69 Doxil (approved 1995), a liposomal delivery system of
70 doxorubicin, Abraxane (approved 2005), an albumin based
71 nanoparticle delivery system of paclitaxel, and in Korea a
72 polymeric micelle delivery vehicle of paclitaxel, Genexol-PM
73 (approved in Korea 2007), have recently been approved.¹⁵ All
74 of these delivery systems are passively targeted formulations
75 and they can suffer from a number of limitations. Lack of
76 control in uptake in these systems can result in off target drug
77 delivery, which may lead to multiple drug resistance, a situation
78 where chemotherapy treatments fail in patients due to the
79 resistance of cancer cells to one or more drugs.^{10,16} Another
80 limitation of the passive strategy is that certain tumors do not
81 strongly exhibit the EPR effect, and the permeability of vessels
82 can be highly heterogeneous throughout a single tumor.¹⁶

83 Active targeting of nanoparticles is where a targeting moiety
84 (e.g., ligand, antibody, peptide) is introduced onto the
85 nanoparticle system to target specific changes in cancer cell
86 biology, which are highly upregulated in comparison to the
87 healthy surrounding cells and tissues.^{10,17} In this process,
88 nanoparticles will recognize and bind to target cells through
89 ligand–receptor interactions, and the bound nanoparticles are
90 internalized before the drug is released inside the cell, resulting
91 in less off target drug release compared to passively targeted
92 systems. One method to improve the retention of nano-
93 particles and in turn their therapeutic payload in cancer tumors
94 is active targeting to cell membrane receptors that are
95 overexpressed in cancer cell lines, such as the epidermal
96 growth factor receptor (EGFR).^{18,19} EGFR up regulation has
97 been implicated in the aggressiveness of several cancers
98 including breast (overexpression evident in approximately
99 50% of triple negative breast cancer tumors),²⁰ renal, ovarian,
100 colon, and nonsmall cell lung cancer.²¹ As a result of this
101 targeting, the EGF receptor is an exciting prospect for actively
102 targeted chemotherapeutic nanoparticle formulations in the
103 treatment of a range of cancers.

104 It is now widely accepted that irrespective of the targeting
105 strategy adopted in the design for drug delivery, i.e., active or
106 passive, it is pivotal for the nanoparticles to become
107 therapeutically relevant, and their intracellular uptake is a
108 crucial factor. Furthermore, it is widely established that the
109 cellular internalization is highly dependent on the size, shape,
110 and surface chemistry of the nanoparticles. However, what
111 remains poorly understood is how the intracellular trafficking
112 and intracellular destination of nanoparticles affect the overall
113 therapeutic outcome. In this study, we show that intracellular
114 destination of an actively targeted polymeric nanoparticle and a
115 passively targeted nanoparticle with comparable internalization
116 abilities differ vastly in their overall therapeutic outcome. We
117 demonstrate this using docetaxel (DTX) loaded polymeric
118 nanoparticles with and without GE11 peptide for targeting the
119 EGFR, which is highly overexpressed in a number of cancers.²¹

120 ■ RESULTS AND DISCUSSION

121 The role of nanoparticle targeting is of great interest in the
122 application of nanoparticles for the treatment of disease and
123 injuries, especially their role in cancer medicine. Despite this
124 interest, there are limited studies available which look to
125 compare actively targeted systems, such as the ERP effect, with
126 passively targeted systems. This study demonstrates the
127 superior effectiveness in relevant *in vitro* cancer models of an
128 EGFR targeted polymeric nanoparticle delivery vehicle of
129 docetaxel (DTX) to a nontargeted nanoparticle comparison.
130 The polymeric nanoparticles were prepared by an oil-in-water

emulsion process where poly(glycidyl methacrylate) (PGMA)
131 was first synthesized and then modified with 6-maleimidohex-
132 anoic acid and rhodamine B through epoxide ring opening
133 reactions. The attachment of the rhodamine B fluorescent
134 probe allowed for *in vitro* fluorescent tracking of the
135 nanoparticles, while the 6-maleimidohexanoic acid modifica-
136 tion was used to anchor the GE11 EGFR targeting peptide to
137 the surface of the nanoparticle. The GE11 peptide, with
138 sequence YHWYGYTPQNVI as published previously,^{22,23} was
139 synthesized with a custom tail of four glycine units and a
140 cysteine residue (YHWYGYTPQNVI GGCGC) to allow for
141 consistent and precise attachment to the nanoparticle surface
142 through thiol-maleimide chemistry. The four-glycine linkers
143 providing flexibility in the GE11 targeting moiety away from
144 the nanoparticle surface an important consideration for
145 targeting peptides.^{24,25} For DTX-loaded nanoparticles, the
146 chemotherapeutic was dissolved in the “oil” phase during the
147 emulsion process for encapsulation.
148

The nanoparticles are approximately 190 nm in diameter
149 (PGMA-DTX-GE11, 195.3 nm, PDI 0.104; PGMA-DTX, 150
189.3 nm, PDI 0.102) with the GE11 targeted nanoparticles
151 exhibiting a higher zeta potential (PGMA-DTX-GE11, $24.2 \pm$
152 5.8 mV; PGMA-DTX NPs, -12.6 ± 8.4 mV) (Figure 1) 153

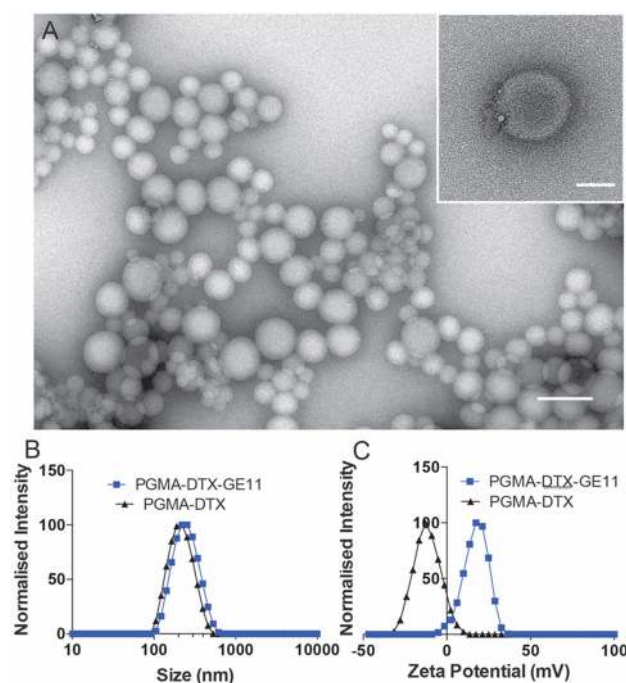


Figure 1. Physical characterization of the targeted DTX loaded nanoparticles. (A) Transmission electron microscopy images of the GE11 targeted DTX loaded nanoparticles with high magnification inset (scale 200 nm, inset 50 nm), (B) dynamic light scattering assessment of nanoparticle size, and (C) zeta potential for the targeted and nontargeted nanoparticle formulations.

resulting from the attachment of the slightly cationic peptide.
154 Drug loading was assessed by reverse phase high-pressure
155 liquid chromatography (HPLC) with the GE11 targeted
156 nanoparticles found to contain $24.0 \pm 1.3\%$ w/w DTX,
157 whereas the nontargeted nanoparticles contained a load of
158 $21.8 \pm 0.8\%$ w/w (see the Supporting Information Figure S1 for the
159 standard curve). The slight difference between particle loading
160 is a result of batch-to-batch variation in the nanoparticle
161

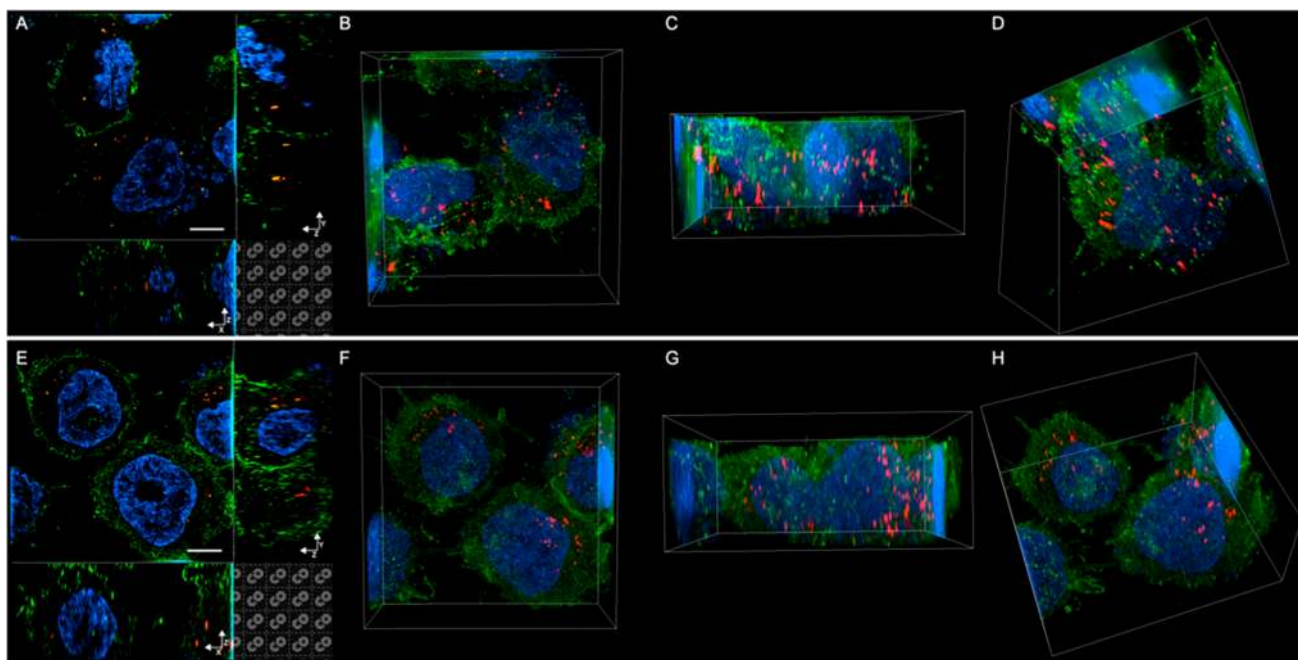


Figure 2. High-resolution confocal microscopy and structured illumination microscopy (SIM) assessment of nanoparticle internalization. (A–D) GE11-PGMA targeted nanoparticles and (E–H) PGMA nontargeted nanoparticles after 24 h incubation with MDA-MB-231 cells. (A) High-resolution confocal microscopy image taken from a central slice of the z-stack showing GE11-PGMA targeted internalization with the x - y , z - y , and z - x profiles displayed. 3D reconstructions of the stack obtained from SIM images (B) top, (C) side, and (D) isometric views. (E) High-resolution confocal microscopy image taken from a central slice of the z-stack showing PGMA nontargeted internalization with the x - y , z - y , and z - x profiles displayed. 3D reconstructions of the stack obtained from SIM images (F) top, (G) side, and (H) isometric views. Hoescht labeling of the nuclei (blue), anti-EGFR (green), and rhodamine labeled nanoparticles (red). Scale bars are 5 μ m.

162 emulsion preparation process. Quantification of the maleimide
 163 functionalization and the subsequent attachment of GE11
 164 targeting peptide was achieved using a Fluorometric Thiol
 165 Quantification Assay Kit. Attachment of GE11 peptide was
 166 found to be 22.4 nmol mg⁻¹ of the targeted nanoparticle
 167 formulation (see the Supporting Information Figure S2). From
 168 physical characterization it was evident that the nanoparticles
 169 were similar in size, shape, and drug loading with the major
 170 difference being the attachment of the GE11 targeting peptide.
 171 Drug release was assessed by HPLC, and it was evident that
 172 DTX was released at a faster rate from the PGMA nontargeted
 173 nanoparticle (71.2% DTX released) when compared to the
 174 GE11 targeted nanoparticle (57.1% released) over a period of
 175 6 days (Supporting Information Figure S3). At 24 h, the
 176 nontargeted formulation had released 7.2% more DTX than
 177 the GE11 targeted formulation, an important result with
 178 downstream *in vitro* experiments based on 24 h incubation
 179 with the therapeutic nanoparticles for comparison.

180 The uptake of both the GE11 targeted and the nontargeted
 181 nanoparticle without DTX loading was assessed by confocal
 182 microscopy in the breast cancer cell line MDA-MB-231. MDA-
 183 MB-231 cells were chosen as DTX has been used extensively in
 184 the treatment of breast cancers, and the overexpression of
 185 EGFR with these cells has been demonstrated previously.^{26,27}
 186 It was anticipated the targeted nanoparticle would have
 187 enhanced cellular association and in turn increased uptake
 188 compared to the nontargeted formulation; however, this was
 189 not evident by confocal microscopy (see the Supporting
 190 Information Figure S4). In fact, the nontargeted nanoparticles
 191 had significant cellular association most likely due to
 192 nonspecific binding and cellular membrane association events.
 193 To further probe nanoparticle internalization, high-resolution

structured illumination microscopy (SIM) was used to confirm
 194 uptake and make comparisons between the targeted and
 195 nontargeted nanoparticles (Figure 2 and Supporting Informa-
 196 tion S5 video links of SIM 3D reconstructions of the PGMA-
 197 GE11 NP and PGMA NP internalization). Cellular internal-
 198 ization was clearly evident for both nanoparticles by SIM
 199 microscopy, confirming both the targeted and nontargeted
 200 nanoparticles were successful at internalizing within a 24 h
 201 time period.
 202

Further assessment of the nanoparticle fate once internalized
 203 was conducted by investigating the colocalization of nano-
 204 particles with both endosomes and lysosomes by immunohisto-
 205 chemistry (see the Supporting Information Figure S6A,B).
 206 Quantitative analysis of these confocal images by comparing
 207 the Pearson's correlation coefficients (a measure to compare
 208 the colocalization) of both the GE11 targeted and the
 209 nontargeted PGMA nanoparticles with endosomes or
 210 lysosomes found significant differences in the intracellular
 211 localization of the nanoparticles following 24 h incubation (see
 212 the Supporting Information Figure S6C). Manders et al. were
 213 the first to use the Pearson correlation coefficient for
 214 fluorescent microscopy colocalization studies.²⁸ Briefly, the
 215 Pearson's correlation coefficient measures the pixel by pixel
 216 covariance in the signal levels of two images, with a perfect
 217 correlation of 1 and a perfectly inverse correlation of -1.²⁹ For
 218 this analysis, only the signal from the respective nanoparticle
 219 images with the endosome or lysosome immunohistochemistry
 220 images were compared to avoid background effects from
 221 alternate fluorescent labels present in the stacked image. This
 222 analysis showed the EGFR targeted nanoparticle associated
 223 significantly less with endosomes and lysosomes when
 224 compared to the nontargeted PGMA nanoparticle. Further to
 225

226 this, both nanoparticles associated more with lysosomes
 227 compared to endosomes after 24 h of incubation (see the
 228 Supporting Information Figure S6). Hence despite similar
 229 levels of cellular internalization of the nanoparticles within
 230 MDA-MB-231 cells, it is apparent that the internal trafficking
 231 of these nanoparticles is affected by the EGFR targeting, which
 232 may also contribute to differences in the therapeutic delivery
 233 and efficacy of these nanoparticle drug delivery vehicles.
 234 To assess the therapeutic efficacy of these DTX loaded
 235 nanoparticles, the half maximal inhibitory concentration
 236 (IC₅₀) was determined in both MDA-MB-231 and the
 237 MCF-7 breast cancer cell lines, respectively (Figure 3). In

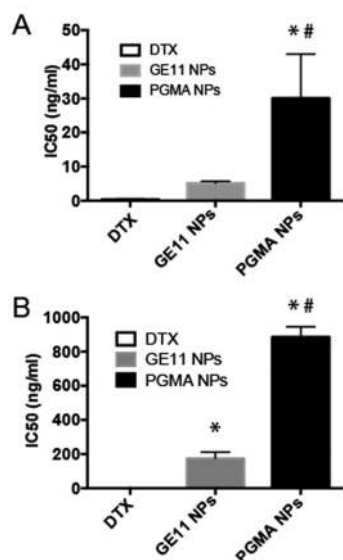


Figure 3. IC₅₀ analysis of docetaxel (DTX), the GE11 targeted DTX nanoparticle formulation (GE11 NPs), and the nontargeted PGMA DTX nanoparticles (PGMA NPs) in both MDA-MB-231 and MCF-7 cells. (A) MDA-MB-231 and (B) MCF-7 breast cancer cell lines. Data displayed as mean ± SD and statistically assessed with a one-way ANOVA followed by a Bonferroni multiple-comparisons test. Significance (**p* < 0.05) for comparison to free DTX and (#*p* < 0.05) for significance between the targeted GE11 NPs and the nontargeted PGMA NPs.

238 assessing the IC₅₀ of these formulations, it was evident that
 239 the GE11-PGMA targeted nanoparticles had a significantly
 240 lower IC₅₀ value compared to the nontargeted formulations in
 241 both MDA-MB-231 (5.1 ng/mL for GE11-PGMA NPs vs 30.1
 242 ng/mL for PGMA NPs, Figure 3A) and MCF-7 (173.5 ng/mL
 243 for GE11-PGMA NPs vs 886.2 ng/mL for PGMA NPs, Figure
 244 3B) cell lines. This is despite the targeted nanoparticle
 245 releasing 7% less DTX than the nontargeted nanoparticle in a
 246 24 h period. These data presented together suggest that the
 247 active targeting of nanoparticles to cancerous cells results in
 248 active uptake and in turn direct delivery of the chemo-
 249 therapeutic at the site of cell activation.

250 To investigate this further and to test whether the GE11-
 251 PGMA targeted nanoparticle was indeed acting through an
 252 EGFR mediated pathway, an experiment was performed to first
 253 block the EGFR receptor in culture with a blocking antibody
 254 before incubation with the targeted and nontargeted DTX
 255 containing nanoparticles at their half inhibitory (IC₅₀)
 256 concentrations, respectively (Figure 4). The presence of the
 257 EGFR blocking antibody by itself in culture media did not
 258 significantly alter cell viability (Figure 4). Pretreatment of cells

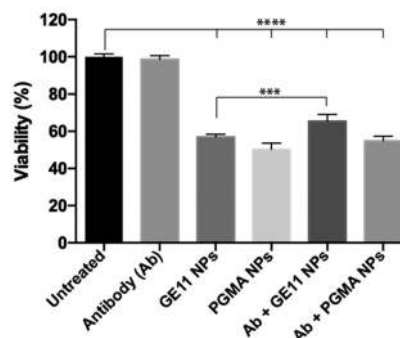


Figure 4. Blocking of the EGFR receptor significantly affects the efficacy of the GE11 targeted nanoparticle (GE11 NPs). MDA-MB-231 cells were plated and preincubated with an EGFR blocking antibody for 1 h before treatment with the IC₅₀ concentration of each nanoparticle formulation and compared. Data displayed as mean ± SD and statistically assessed with a one-way ANOVA followed by a Bonferroni multiple-comparisons test. Significance (***) *p* < 0.001, *****p* < 0.0001).

with this blocking antibody prior to nanoparticle treatment 259
 resulted in significant rescue of cell viability compared to 260
 GE11-PGMA targeted nanoparticle treatment only, but this 261
 rescue was not observed for the nontargeted PGMA 262
 nanoparticle (Figure 4). These data taken together support 263
 an active uptake of the GE11 targeted nanoparticles mediated 264
 by the EGFR due to the significant rescue observed. 265
 Furthermore, the nontargeted nanoparticles are internalized 266
 by cells via passive processes not dependent on the EGFR. 267
 Previous studies investigating similar nanoparticles found that 268
 inhibition of clathrin mediated endocytosis (chlorpromazine), 269
 caveolin mediated endocytosis (nystatin/progesterone), and 270
 micropinocytosis (*N,N*-dimethylamiloride) had no significant 271
 effect on PGMA based polymeric nanoparticles of similar size, 272
 and hence it was assessed that these specific cellular uptake 273
 pathways were not required for polymeric nanoparticle cellular 274
 internalization.³⁰ The activation of the EGFR resulting in 275
 cellular internalization via clathrin dependent uptake is also 276
 supported extensively in the literature.^{31,32} It is evident that 277
 cellular uptake of a therapeutic payload into a target cell is not 278
 alone sufficient for efficacy, but the pathway of internalization 279
 plays an integral role in the therapeutic outcome of 280
 nanoparticle delivery agents. 281

282 CONCLUSIONS

283 Despite significant efforts, the role of active targeting moieties
 284 in nanomedicine versus passively targeted systems still remains
 unclear. In this study, we present data on two model polymeric
 drug delivery agents: one actively targeted through decoration
 of the nanoparticle surface with the EGFR targeting GE11
 peptide; and one without, enabling passive internalization by
 cells. This study demonstrates the importance that active
 targeting of the nanoparticle system has on dictating the
 intracellular trafficking and furthermore the direct role that this
 intracellular trafficking has on the efficacy of therapeutic
 nanoparticles in cancer treatment. These results suggest that
 careful consideration of actively targeted nanoparticles cannot
 only influence the cellular uptake of the therapeutic cargo but
 also have downstream ramifications on the cellular recognition
 and intracellular processing of the cargo once inside the cell.
 For nanoparticle drug delivery systems to reach the potential
 they clearly possess in the treatment of cancer, thorough 299

300 studies on the efficiency of targeting need to occur to ensure
301 optimal efficacy from these systems can be achieved.

302 ■ MATERIALS AND METHODS

303 **Materials.** Glycidyl methacrylate (GMA, 97%, Sigma), 2,2'-
304 azobis(2-methylpropionitrile) (AIBN, 0.2 M in toluene, Sigma),
305 methyl ethyl ketone (MEK, 2-butanone; Fischer Chemical, LR grade),
306 Pluronic F-108 (Mn = 14 600 g mol⁻¹, Sigma), 6-maleimidohexanoic
307 acid (Sigma), rhodamine B (Kodak), sodium acetate (Sigma), Tris(2-
308 carboxyethyl)phosphine hydrochloride (Sigma), and docetaxel (LC
309 Laboratories, 97%) were all used as received. The GE11 peptide was
310 synthesized with a GGGC linker sequence at the carboxyl terminus
311 with the following sequence (YHWYGYTPQNVIGGGGC, China
312 Peptides) and it was used as received. Tissue culture and
313 immunohistochemistry reagents such as minimum essential medium
314 (Invitrogen), fetal bovine serum (Invitrogen), Trypsin/EDTA
315 (Sigma), GlutaMAX (Invitrogen), paraformaldehyde (Sigma),
316 mouse anti-EGFR Monoclonal antibody (H11, Life Technologies,
317 catalog no. MA5-13070), Hoechst (Sigma), goat antimouse Alexa
318 Fluor 488 (Life Technologies, catalog no. A11001), EEA1 antibody
319 (Abcam, ab2900), LAMP2 antibody (Santa Cruz, sc-18822 (H4B4)),
320 goat antirabbit AF488 (ThermoFisher, A-11034), goat antimouse
321 AF647 (ThermoFisher, A-21235), and fluorobrite (Invitrogen) were
322 used as received. Coverglass used for SIM imaging were 170 ± 5 μm
323 from Marienfeld, Germany. All other reagents were of analytical
324 grade.

325 **Poly(glycidyl methacrylate) (PGMA) Synthesis.** PGMA (Mn =
326 186 200 g/mol, PDI 1.644) was synthesized by the free radical
327 polymerization of glycidyl methacrylate in methyl ethyl ketone (1:1
328 ratio) at 80 °C for 18 h. The obtained polymer was purified by
329 multiple precipitations from the MEK solution using methanol.
330 Azobis(isobutyronitrile) (AIBN) was used as the initiator with
331 product and molecular weight confirmed by NMR and GPC,
332 respectively.

333 **Synthesis of PGMA Nanoparticles.** PGMA (200 mg) was
334 reacted with 6-maleimidohexanoic acid (MHA, 300 mg) in MEK (30
335 mL, 70 °C, 5 h). The solvent was removed by a rotovap to
336 approximately 2 mL, and the MHA modified PGMA (PGMA-MHA)
337 precipitated in diethyl ether. The MHA-PGMA was reacted with
338 rhodamine B (RhB, 20 mg) in MEK (30 mL, 70 °C, 5 h). The
339 PGMA-RhB-MHA was collected by precipitating in diethyl ether and
340 dried under a flow of N₂ to remove residual ether.

341 PGMA nanoparticles were prepared using a nonspontaneous
342 emulsification process as described previously.^{30,33} The modified
343 polymer (PGMA-RhB-MHA for GE11 attachment post nanoparticle
344 formulation or PGMA-RhB for nontargeted nanoparticles without
345 GE11 modification, approximately 100 mg) was dissolved in a mixture
346 of chloroform (1.5 mL) and MEK (4.5 mL) along with docetaxel (20
347 mg). This organic phase was added dropwise, with rapid stirring, to an
348 aqueous phase of Pluronic F-108 (12.5 mg mL⁻¹, 30 mL), and the
349 emulsion was homogenized with a probe-type ultrasonicator on low
350 power for 1 min. The organics were evaporated, and the sample
351 centrifuged at 3000g (45 min) to remove large aggregates and excess
352 polymer. The supernatants were collected, and the nanoparticles
353 centrifuged at 20 000g (20 min) to collect the nanoparticles.
354 Nanoparticles were washed a further 3 times with aqueous pluronic
355 F-108 (12.5 mg mL⁻¹, 30 mL), before storage at 4 °C of the stock and
356 aliquots collected and freeze-dried for concentration determination.

357 **GE11 Peptide Attachment.** The GE 11 peptide was dissolved in
358 a degassed sodium acetate buffer (50 mM, pH 4.5) prepared in
359 Pluronic F108 (12.5 mg mL⁻¹) to a final GE11 concentration of 5 mg
360 mL⁻¹. 6-Maleimidohexanoic acid coated nanoparticles (4 mg) were
361 centrifuged (16 000g, 0.5 h) and suspended in the acetate buffer (50
362 mM, 1 mL). This process was repeated a second time (solution A).
363 Tris(2-carboxyethyl)phosphine hydrochloride (TCEP, 40 mg) was
364 dissolved in the acetate buffer (50 mM, 920 μL), and an aliquot of the
365 GE11 peptide stock was added (80 μL) and left to incubate (0.5 h) at
366 room temperature (RT, solution B). Solution A was combined with
367 solution B and left to stir slowly (24 h, RT, under argon). Following

peptide attachment, the nanoparticles were washed by repeated
368 centrifugation (2× acetate buffer, final wash Pluronic F108 solution, 369
16 000g, 0.5 h) before suspending in 1× phosphate buffered saline
370 (PBS) for ongoing *in vitro* studies. 371

372 **Nanoparticle Characterization.** Dilute samples of nanoparticles
373 in MQ water were assessed for hydrodynamic size and zeta potential
374 by dynamic light scattering on a Zetasizer Nano ZS (Malvern
375 Instruments). Nanoparticles were drop-casted on carbon coated
376 transmission electron microscopy (TEM) grids and imaged with an
377 accelerating voltage of 120 kV on a JEOL 2100 TEM for physical
378 characterization. Lyophilized aliquots of the nanoparticles were used
379 for drug loading determination by reverse phase high-pressure liquid
380 chromatography (RP-HPLC). The lyophilized aliquots were weighed
381 before being suspended in methanol (1 mL). The suspension was
382 sonicated and vortexed to dissolve the encapsulated docetaxel. The
383 nanoparticles were removed via centrifugation (16 000g, 0.5 h), and
384 the supernatant containing the dissolved free drug was analyzed using
385 RP-HPLC. The measurements were performed on a Waters 2695
386 separation module with a Waters 2489 UV/vis detector (determi-
387 nation λ = 232 nm) using reverse phase isocratic elution (methanol-
388 water, 70:30; flow rate at 1.0 mL min⁻¹) through a C18 column (150
389 mm × 4.60 mm, 5 μm, 25 ± 5 °C). The measurements were
390 compared against a standard curve performed in identical run
391 conditions (see the Supporting Information Figure S1 for standard
392 curve). 6-Maleimidohexanoic acid and the final peptide concentration
393 attached to the nanoparticles was quantified using a fluorometric thiol
394 quantification assay kit as per the manufacturer's instructions (abcam,
395 ab112158, see the Supporting Information Figure S2 for the standard
396 curve).

397 **Drug Release Experiment.** Docetaxel release was assessed by
398 RP-HPLC following a previously published protocol by Singh et al.
399 using the above-mentioned running conditions for drug loading
400 characterization.³⁴

401 **Cell Culture.** MDA-MB-231 or MCF-7 cells were cultured in
402 Minimum Essential Medium α with 10% fetal bovine serum (FBS),
403 1× GlutaMAX, and 0.15% sodium bicarbonate in a humidified
404 incubator at 37 °C and 5% CO₂. Cells were cultured without
405 antibiotics or antimycotics. Cells were washed with PBS, trypsinised,
406 collected by centrifuge (1 000g, 5 min), and subcultured before
407 reaching 80% confluence. Passage numbers were kept low for all
408 experiments.

409 **IC50 Analysis.** Cells were plated in 96 well plates at 5000 cells/
410 well in 50 μL of media. Plates were incubated for 24 h and checked
411 visually for consistent cell density before treatments were applied.
412 Nanoparticle concentrations were calculated by diluting according to
413 total DTX mass from HPLC loading analysis. DTX and nanoparticle
414 stock solutions were prepared in a minimum of 7 concentrations
415 between 10 and 0.0001 mg/mL in DMSO. The stocks were sterilized
416 with ultraviolet light, then diluted 1 in 250 parts media. Treatments in
417 media were applied in triplicate at 50 μL per well then thoroughly
418 agitated. Treatments containing DMSO were diluted by 1/2 by
419 addition to media in wells, for a total of 0.2% v/v DMSO in 100 μL of
420 media per well. The outer wells of each plate were filled with PBS to
421 maintain even evaporation in treated cells. Each plate also contained
422 three wells of cells with 100 μL of media and three with 100 μL of
423 media + 0.2% v/v DMSO as controls. Treatments were allowed to
424 incubate with cells for a total of 24 h at which point cell culture media
425 containing each of the treatments were removed by inverting the
426 plate, gently shaking, and dabbing onto a sterile paper towel. Cells
427 were then immediately washed twice with PBS (100 μL per well),
428 agitated for 1 min, then the PBS was removed. Warmed media was
429 immediately applied (100 μL per well). Plates were then incubated for
430 48 h, accumulating to a total 72 h of *in vitro* study period.

431 Assays were performed using warmed CellTiter 96 Aqueous One
432 Solution Cell Proliferation Assay ((3-(4,5-dimethylthiazol-2-yl)-5-(3-
433 carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium), "MTS")
434 solution at 20 μL per well. Plates were read 3 h after the reagent was
435 applied with a PerkinElmer EnSpire Multimode plate reader. 435
436 Absorbance was measured at 490 nm in triplicate and averaged. 436
437 IC50 curves were constructed from triplicate assay data in GraphPad 437

438 Prism. Data sets were normalized, then analyzed by 4-parameter
439 nonlinear regression with an inhibition model.

440 **EGFR Blocking Experiment.** MDA-MB-231 cells were plated in
441 96 well plates at 5000 cells/well in 50 μ L of media. After 24 h, cells
442 were treated with a blocking antibody anti-EGFR (H11, Life
443 Technologies, catalog no. MA5-13070) at 1 μ g/mL or PBS (control)
444 diluted in media for 1 h followed by a 24 h incubation with either
445 targeted GE11-PGMA-DTX-NPs or untargeted PGMA-DTX-NPs at
446 their determined IC50 concentrations with MDA-MB-231 cells. At a
447 time 24 h after the incubation, media was changed for fresh media and
448 the plate remained in the incubator for a further 48 h. After this time,
449 cell viability was assessed using CellTiter-Glo 2.0 following the
450 manufacturer's instructions.

451 **Confocal Imaging.** MDA-MB-231 cells were cultured as above
452 on glass coverslips in 6-well plates and treated for 24 h with
453 nanoparticle formulations (10 μ g/mL), before thorough washing with
454 PBS and fixation with 4% paraformaldehyde. Samples were
455 immunohistochemically labeled with Mouse anti-EGFR Monoclonal
456 antibody (1:1000, 3% BSA in PBS, 4 $^{\circ}$ C, overnight), washed with PBS
457 (twice, 5 min), and labeled with goat antimouse Alexa Fluor 488
458 (1:500 in Fluorobrite, 37 $^{\circ}$ C, 30 min). Secondary was then removed,
459 washed with PBS (twice, 5 min), and stained with Hoechst (1:1000,
460 PBS, RT, 20 min) before washing with PBS (twice, 5 min).
461 Colocalization studies were treated as per the above samples with the
462 following antibodies (endosomes, 1 $^{\circ}$ rabbit-EEA1, 2 $^{\circ}$ antirabbit
463 AF488 and lysosomes, 1 $^{\circ}$ mouse-LAMP1, 2 $^{\circ}$ antimouse AF647.
464 Samples were mounted on glass microscope slides and imaged on a
465 Nikon Ti-E inverted confocal microscope with a Nikon A1Si spectral
466 detector system and a Nikon 60 \times oil immersion, 1.49NA objective.
467 For super resolution imaging, images were captured on the Nikon N-
468 SIM system with a Nikon SR 100 \times , 1.49NA objective. Cells were
469 treated as above but were seeded on a coverglass of accurate thickness
470 (170 \pm 5 μ m from Marienfeld, Germany) to aid in reconstruction
471 efficiency.

472 **Statistical Analysis.** All results are displayed as means \pm SD. Data
473 was analyzed using the GraphPad Prism version 6.0 data management
474 software to conduct ANOVA on groups of data. Statistically
475 significant differences between each treatment were determined
476 using Bonferroni post hoc tests (* p < 0.05, ** p < 0.01, *** p <
477 0.001, **** p < 0.0001).

478 ■ ASSOCIATED CONTENT

479 ● Supporting Information

480 The Supporting Information is available free of charge on the
481 ACS Publications website at DOI: 10.1021/acs.lang-
482 muir.8b02946.

483 Complete synthetic procedures and protocols, support-
484 ing experiments including RP-HPLC standard curve and
485 drug loading for docetaxel, thiol assay quantification,
486 supporting confocal imaging analysis, SIM video links,
487 and Pearson colocalization analysis of nanoparticles with
488 endosomes or lysosomes (PDF)

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496 Notes

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511 ■ ABBREVIATIONS

BSA, bovine serum albumin; DTX, docetaxel; EGFR, 512
epidermal growth factor receptor; EPR effect, enhanced 513
permeation and retention effect; FDA, United States Food 514
and Drug Administration; GE11, epidermal growth factor 515
targeting peptide; GE11-PGMA-DTX-NPs, polymeric nano- 516
particles loaded with docetaxel and containing the GE11 517
targeting peptide; IC50, half maximal inhibitory concentration; 518
PBS, phosphate buffered saline; PGMA, poly(glycidyl meth- 519
acrylate); PGMA-DTX-NPs, nontargeted polymeric nano- 520
particles loaded with docetaxel; RP-HPLC, reverse phase 521
high-pressure liquid chromatography; SIM, structured illumi- 522
nation microscopy 523

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