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**Introduction and Scope of Research:** Our research objective is to develop a novel set of technologies that will target the Jak/Stat signaling cascade in breast cancer (1). These technologies, which involve a new class of affinity reagents and intramolecular delivery tools (2, 3), have the capability of identifying the most important nodes in this signaling pathway and ultimately inhibiting or modifying them to influence effects on breast cancer cell proliferation and death. The interplay between the Jak and Stat components in cytokine signaling has been an area of intense investigation (4, 5). However, although many of the molecular interactions that occur between them and with other signaling partners have been broadly implicated in breast cancer, they are poorly characterized because of a lack of appropriate experimental tools. Consequently, a host of basic questions remain to be answered. Our goal is to develop an experimental framework to sort out the most important interactions in the pathway and establish whether there is a specific Achilles Heel that can be exploited to attack breast cancers in innovative ways. As a long-term goal, we will utilize this information to develop novel synthetic antibody reagents that can be delivered with precision and potency to breast cancer cells.

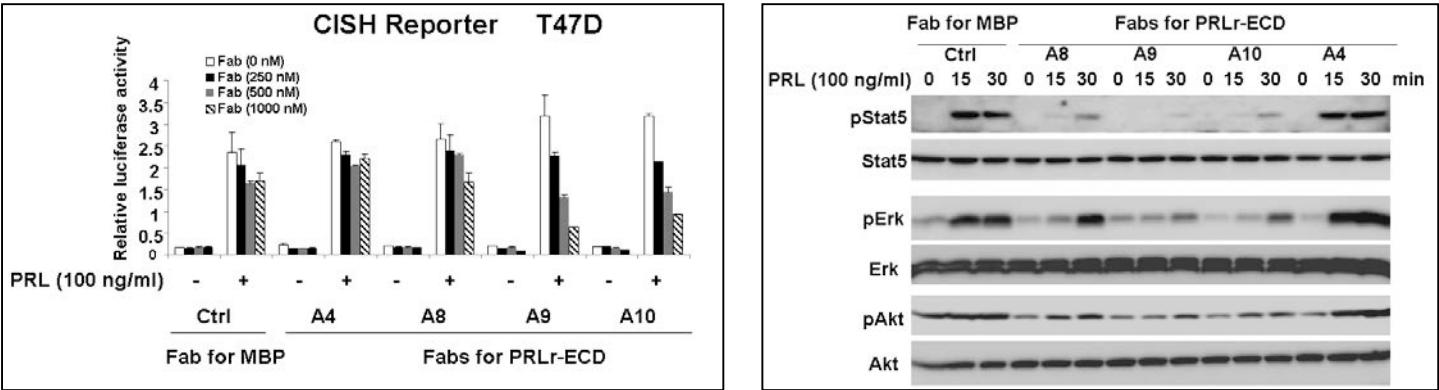
**Research Accomplishments**

**Aim 1- Generate synthetic antigen binders (sABs) to components of the prolactin receptor signaling network.** This was reported in the previous progress report in detail; significant accomplishments are summarized below.

**Aim 2- Test the inhibitory sABs in cell-based assays to evaluate their effectiveness as receptor antagonists.** Our hypothesis is that sABs that block hormone binding will have an inhibitory effect on downstream signaling as measured by Stat activation. To assess this, we have tested four inhibitory sABs in cell based assays to measure inhibition of Stat5 activation (Figure 1).

Cells were incubated with increasing concentrations of each sAB, or a control sAB against bacterial maltose binding protein. The effect of the sABs on prolactin signaling was determined using a dual luciferase luminescence assay under the control a of phospho-Stat promoter (Figure 1A).

The results indicate that 3 sABs (A8, A9 and A10) significantly inhibit prolactin signaling in a concentration dependent manner.



**Figure 1- sAB inhibition of prolactin signaling.** A) Effects on cell signaling based on luciferase expression. B) Measurement of decrease in phospho-Stat5 caused by the four individual inhibitory sABs.

To further examine the effect of the sABs on prolactin receptor signaling, we determined the change in the phosphorylation levels of various factors using western blot. While the control sAB (Fab for MBP) has no effect on prolactin signaling, sABs A8, A9 and A10 strongly inhibit phosphorylation of Stat5 and Erk (Figure 1B). Again, sAB A4 shows no effect on downstream signaling of prolactin as was observed in the luciferase assay.

Additionally, we hypothesize that inhibiting the proline isomerase activity of cytoplasmic CypA will effectively turnoff downstream signaling by eliminating the switch that ultimately leads to Jak activation. To test this, we have used the sAB pipeline to generate a family of high affinity binders that are specifically targeted to block the active site of CypA and block binding of cyclosporine. Four inhibitory sABs have been evaluated using surface plasmon resonance to establish their binding affinities to CypA (Table 1). The affinities range from 2 nM to 12 nM.

sABs	#3	#4	#5	#8
Kd (nM)	3	2	7	12

Table 1. SPR analysis of binding affinity of selected FABs. Purified sABs were injected over Sensor Chip NTA where 6His-tag CypA was immobilized on the surface. The equilibrium constant Kd was used to measure the binding affinity of two components.

A future goal is to use our receptor mediated delivery technology to introduce an inhibitory sAB into live cells to assess the consequence of selectively blocking CypA on prolactin signaling.

## Key Research Accomplishments

- Used phage display to select sABs that act as inhibitors to CypA and CypB (previously reported).
- Generated and characterized a sAB against the PRLr-ECD and showed it was a potent inhibitor against PRL binding to the ECD (previously reported).
- Determined the X- ray crystal structure of the sAB-PRLr-ECD crystal structure at 2.8Å resolution (new).
- Established that sAB binding interferes with PRL binding through an allosteric mechanism (new).
- Assess the effects of CypA inhibition on prolactin signaling (new).

## Reportable Outcomes

### Publications-

Rizk SS, Luchniak A, Brawley CM, Rock RS, Kossiakoff AA. (2009) "An engineered substance P variant for receptor-mediated delivery of synthetic antibodies into tumor cells." Proc Natl Acad Sci USA. 106:11011-5.

Duguid E, Zhang J, Rizk SS, Symborska A, Kouido J, Clevenger CV and Kossiakoff AA. “ The structure and function of a synthetic antibody-complexed to the extracellular domain of the prolactin receptor: signaling inhibition through an allosteric mechanism” (in preparation).

## **Conclusion**

The reagents that are being generated are of exceptional importance to both basic and translational science in that they will provide a means of delivering inhibitory probes intracellularly into viable cells. The generation of sABs as outlined above has been successful and is following the timeline identified in the SOW.

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**Appendicies –** Attached paper.

**Supporting Data –** None.

# An engineered substance P variant for receptor-mediated delivery of synthetic antibodies into tumor cells

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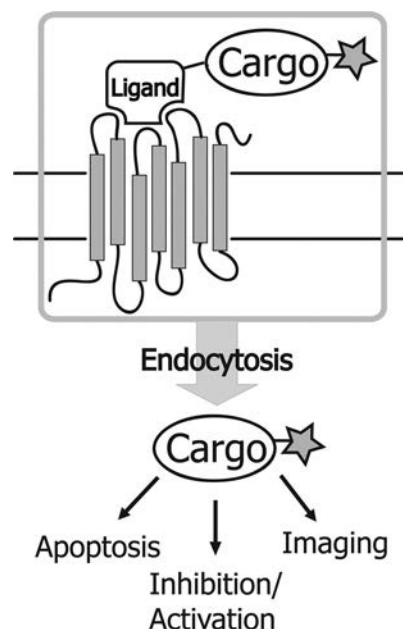
We have developed and tested a robust delivery method for the transport of proteins to the cytoplasm of mammalian cells without compromising the integrity of the cell membrane. This receptor-mediated delivery (RMD) technology utilizes a variant of substance P (SP), a neuropeptide that is rapidly internalized upon interaction with the neurokinin-1 receptor (NK1R). Cargos in the form of synthetic antibody fragments (sABs) were conjugated to the engineered SP variant (SPv) and efficiently internalized by NK1R-expressing cells. The sABs used here were generated to bind specific conformational forms of actin. The internalized proteins appear to escape the endosome and retain their binding activity within the cells as demonstrated by co-localization with the actin cytoskeleton. Further, since the NK1R is over-expressed in many cancers, SPv-mediated delivery provides a highly specific method for therapeutic utilization of affinity reagents targeting intracellular processes in diseased tissue.

actin | drug delivery | synthetic antibody fragment

**D**irected delivery of bioactive reagents into cells is one of the most intensely pursued objectives in biomedical research, yet few major breakthroughs have materialized that can be broadly applied in both laboratory research and therapeutic applications. Transfection reagents (1–3) as well as cell-penetrating peptides (4–7) lack cell-type specificity, and in many cases are associated with heavy toxicity (8), restricting the successful utilization of such reagents *in vivo*. In contrast, receptor-mediated delivery (RMD) systems (9–11) are conceptually simple and potentially very powerful. By attaching a biomolecular cargo to a natural ligand, receptor-mediated internalization serves as an extremely efficient and selective pathway for delivery of a wide variety of bioactive entities without compromising the integrity of the cell membrane (Fig. 1). Previously described RMD methods rely on cargo attachment to hormones (12) or monoclonal antibodies that recognize a cell-surface receptor (13). However, these large protein ligands can be difficult to isolate or modify in a specific manner.

Here we present a delivery method based on substance P (SP), an 11 amino acid neuropeptide that is rapidly internalized through specific interaction with the neurokinin-1 receptor (NK1R) (14–16). We have engineered a variant of SP (SPv) that enables the specific delivery of proteins to NK1R-expressing cells. Unlike larger ligands, SPv is easily chemically synthesized, allowing the incorporation of various reactive groups to facilitate coupling to myriad biomolecular cargos. Importantly, the NK1R is highly over-expressed in several aggressive tumors (17, 18), particularly astrocytomas and glioblastomas (19, 20), where the level of expression is correlated with the degree of malignancy (21). Consequently, an RMD system using SPv could provide a high level of specificity for the targeting of tumor tissue, with little toxicity for noncancerous cells.

To evaluate the efficacy of the SPv-NK1R delivery system, we used a set of synthetic antibody fragments (sABs) selected from phage display libraries to bind various forms of actin. sABs



**Fig. 1.** Receptor-mediated delivery. A fluorescently-labeled cargo is attached to a natural ligand. Upon binding to the receptor, the cargo is internalized where it can carry out the desired function.

resemble the Fab fragment of an antibody, consisting of a 55-kDa heterodimer of the heavy and light chain variable fragments, forming 1 antigen binding site that can be engineered to bind a specific target protein (22–24). A subset of these sABs can induce changes to actin structure or organization *in vitro*. We demonstrate that sABs conjugated to SPv are rapidly internalized only by cells expressing the NK1R. The internalized sABs appear to escape the endosomes and retain their *in vitro* properties by altering actin organization within the cytoplasm. Additionally, we take advantage of the over-expression of the NK1R in the human glioblastoma cell line U87 (20) to confirm the utility of SPv as a highly specific delivery vehicle that differentiates between tumor and non-tumor cells. Our findings suggest that the SPv-NK1R system could be used in an array of applications, from live cell imaging, mapping signaling pathways, as well as mediating delivery of therapeutic affinity reagents to disrupt intracellular components within tumor cells.

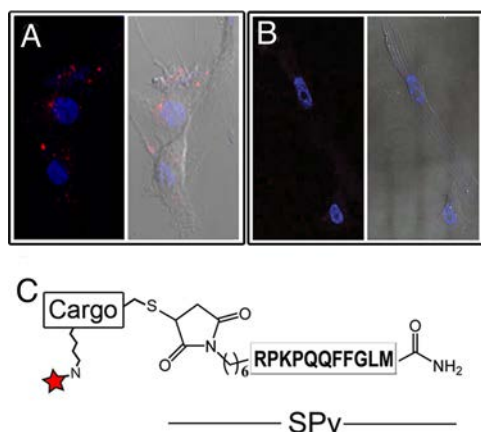
Author contributions: S.S.R. and A.A.K. designed research; S.S.R. and A.L. performed research; S.U. and C.M.B. contributed new reagents/analytic tools; S.S.R., A.L., C.M.B., and R.S.R. analyzed data; and S.S.R. and A.A.K. wrote the paper.

The authors declare no conflict of interest.

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This article contains supporting information online at [www.pnas.org/cgi/content/full/0904907106/DCSupplemental](http://www.pnas.org/cgi/content/full/0904907106/DCSupplemental).





**Fig. 2.** Specific delivery of cargo to tumor cells by SPv. (A) U87 glioblastoma incubated with Alexa555-labeled SP for 2 h followed by fixing and nuclear staining (blue) show internalization of the peptide (red). (B) Normal human astrocytes show no detectable internalization of the peptide. (C) Conjugation of SPv to protein cargo. SPv is synthesized to contain the amino acid sequence of SP (boxed) in addition to an N-terminal maleimide for attachment of protein cargo via a surface cysteine residue. An amine reactive fluorophore (star) is attached to the cargo via surface lysines.

## Results

**Engineering a Delivery Vehicle Based on Substance P.** Substance P (SP) is an 11 amino acid neuropeptide that is specifically internalized through binding to the NK1R, a member of the tachykinin G-coupled receptor family (14–16). The NK1R is highly overexpressed in brain tumor cells (19) compared with normal astrocytes. As a result, U87 glioblastoma cells readily internalize fluorescently-labeled SP (Fig. 2A), while internalization is undetectable in normal human astrocytes (NHA) (Fig. 2B). Additionally, western blot analysis of the NK1R expression confirms significant levels of receptor expression in U87 (19), whereas the receptor is undetectable in NHA (Fig. S1).

We used the selectivity of SP internalization by NK1R-expressing cells to engineer a specific vehicle for the delivery of protein cargo. A useful feature of SP is that only its C-terminal 7 amino acids are essential for binding to the NK1R and subsequent internalization (25), allowing incorporation of reactive groups for cargo attachment at the N terminus without affecting its function. Using standard *t*-Boc chemistry, we synthesized a variant of SP (SPv) with a “built-in” maleimide group at the N terminus. The maleimide moiety can be used to react with free cysteine residues under mild conditions to form a stable thioether bond (Fig. 2C), thus allowing attachment of any protein with free surface cysteines to SPv.

**Specific Delivery of Actin-Binding sABs to NK1R-expressing Cells.** To demonstrate the ability of SPv to deliver fully-functional protein cargo, we chose a set of synthetic antibody fragments (sABs) engineered by phage display selection to bind distinct conformational forms of actin filaments. Characterization of the in vitro actin-binding properties of 3 sABs (sAB-27, sAB-4, and sAB-19) shows that each sAB exhibits binding to the actin filaments. Electron microscopy imaging indicates that sAB-27 reconfigures the overall structure of actin by inducing filament bundling (Fig. 3A). In contrast, sAB-4 was shown to induce rapid severing of the actin filaments by total internal reflection fluorescence (TIRF) microscopy (Fig. 3B). Finally, sAB-19 binds to the end of the filaments without disrupting the overall actin structure (Fig. 3C). To conjugate the sABs to SPv, a cysteine mutation was introduced at position 121 of the heavy chain of each sAB. This position is solvent accessible and lies far from the antigen binding site, allowing conjugation to the SPv

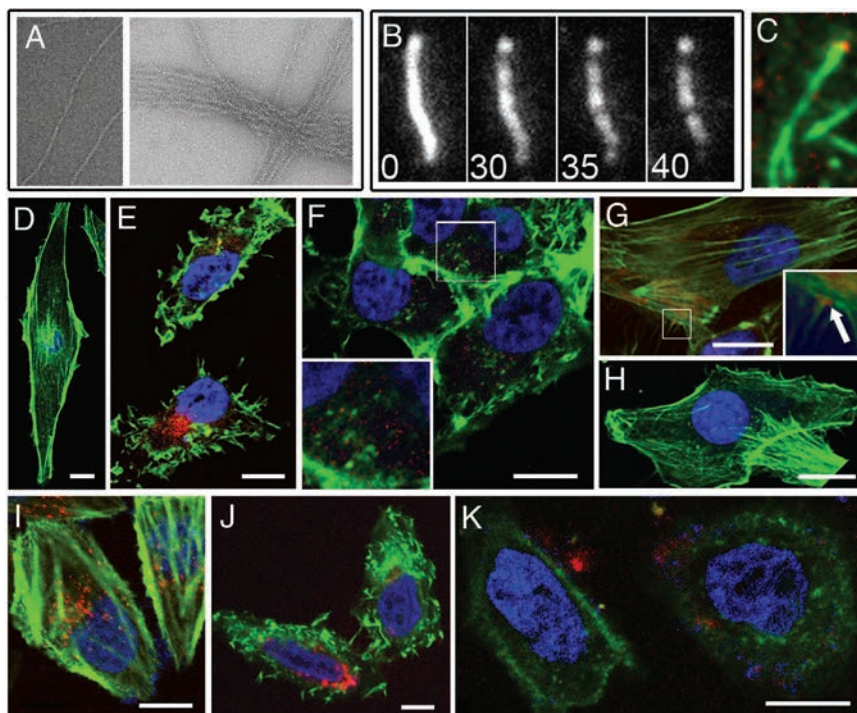
variant via the maleimide moiety without interfering with actin binding. Before attachment to SPv, each sAB was fluorescently labeled via surface lysines using an amine reactive fluorophore to facilitate visualization by confocal microscopy.

As the structure and location of intracellular actin are readily identifiable by phalloidin staining, the ability of SPv to deliver functional sABs can be rigorously tested by the ability of the internalized sABs to disrupt or localize with actin filaments. U87 cells were incubated for 2 h with 20 nM SPv conjugated to sAB-4, sAB-27, or sAB-19. The cells were then fixed, stained with phalloidin, and analyzed using confocal microscopy. A z-series of cells treated with the sAB-SPv conjugates confirms internalization and localization of the conjugate with actin filaments (Movie S1). Additional analysis of the confocal microscopy images indicates that the SPv-attached sABs internalized by U87 cells retain their in vitro properties by co-localization or reorganization of the cellular actin structure. Compared with untreated cells (Fig. 3D), SPv-mediated delivery of sAB-27 induces a conformational change to the actin filaments that stimulates bundling, drastically changing the overall shape of the actin filaments and, consequently, cell morphology (Fig. 3E). Similar results are obtained at very low concentrations (2 nM) of sAB-27-SPv. Cells treated with the sAB-4-SPv conjugate lack prominent stress fibers or actin-based protrusive structures (Fig. 3F), suggesting that sAB-4 induces severing of the filaments in the cytoplasm as observed in vitro (Fig. 3B). In contrast, the internalized sAB-19-SPv conjugate does not affect the overall structure of the actin filaments (Fig. 3G). In some cases, sAB-19 was found to localize to the base of filopodia (Fig. 3G, inset), presumably at the so-called pointed end of the actin filaments. A similar localization pattern was observed by immunostaining of fixed cells using fluorescently-labeled sAB-19 (Fig. S2). No internalization was observed after treatment of cells with up to 200 nM fluorescently-labeled sAB-4 alone (Fig. 3H). While in some cases the internalized sABs localize to peri-nuclear vesicles, the significant reorganization of intracellular actin indicates that a considerable fraction of the internalized sABs escapes the endosome in an active form.

Unlike U87 cells, HeLa cells lack significant amounts of the NK1 receptor. We therefore transfected HeLa cells with a plasmid coding for the NK1R, which then show the ability to internalize fluorescently-labeled SP alone (Fig. 3I) or SPv-conjugated sABs (Fig. 3J and K). As observed in U87 cells and in vitro, sAB-27 disrupts both the structure of the actin filaments and the overall cell morphology (Fig. 3K), while sAB-4 results in severing of actin filaments (Fig. 3J), compared to cells treated with SP only (Fig. 3I). The ability of the SPv-delivered sABs to affect the actin structure in transfected HeLa cells suggests that introduction of the NK1R is sufficient for the delivery of active cargo using SPv. Thus, this delivery technology has promise for a wide variety of transfectable mammalian cell lines.

**The Effect of sAB-27 on Cell Viability.** The interaction of actin filaments with many cellular components is essential to the normal function of cells. Thus, the actin-bundling activity of sAB-27, which significantly disrupts actin structure, would presumably have a drastic effect on the survival of cells. Therefore, we investigated the effect of sAB-27 on the viability of U87 glioblastoma cells following SPv-mediated delivery. U87 cells were treated with either 10 nM or 20 nM of SPv alone, sAB-27 alone, or the SPv-sAB-27 conjugate, and an MTS cell viability assay was carried out. Incubation of U87 cells with the SPv-sAB-27 conjugate showed a concentration-dependent decrease in cell viability compared with treatment with SPv or sAB-27 alone (Fig. 4A). A time-course of the viability of cells treated with the SPv-sAB-27 conjugate was carried out up to 48 h post-treatment. A drastic decrease in the viability of the glioblastoma cells was observed within hours following treatment with



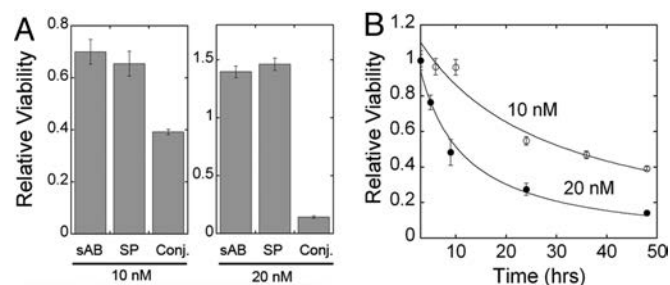


**Fig. 3.** Delivery of actin-binding sABs to cells expressing the NK1 receptor. (A) Electron micrographs of the bundling effect of sAB-27; left panel: actin only; right panel: actin in the presence of  $6.4 \mu\text{M}$  sAB-27. (B) Severing of TMR-labeled actin filaments by sAB-4 viewed by TIRF microscopy. Numbers indicate time in seconds following addition of  $0.5 \mu\text{M}$  sAB-4. (C) TIRF microscopy imaging of TMR-labeled actin filaments (green) in the presence of  $0.25 \mu\text{M}$  cy5-labeled sAB-19 (red). (D) Phalloidin staining of untreated U87 glioblastoma cell. (E) The actin-bundling effect in U87 cells treated with  $20 \text{ nM}$  cy5-labeled sAB-27-SPv conjugate (red) followed by fixing and staining with TMR-phalloidin (green). (F) U87 cells treated with  $20 \text{ nM}$  Alexa488-labeled sAB-4-SPv conjugate (red), followed by fixing and staining with TMR-phalloidin (green), inset: enlarged box showing severed actin filaments. (G) U87 cells treated with  $20 \text{ nM}$  Alexa488-labeled sAB-19-SPv conjugate (red) followed by fixing and staining with TMR-phalloidin (green); arrow in inset indicates an example of sAB-19 localizing to the base of filopodia. (H) U87 cells treated with  $200 \text{ nM}$  sAB-4 only exhibiting normal actin filaments. (I–K) HeLa cells transiently transfected with the NK1 receptor. (I) HeLa cells were treated with  $100 \text{ nM}$  SP-Alexa555 (red), then fixed and stained with FITC-phalloidin (green). (J) HeLa cells treated with  $20 \text{ nM}$  cy5-labeled sAB-27-SPv conjugate (red), followed by fixing and staining with TMR-phalloidin (green). (K) HeLa cells treated with  $20 \text{ nM}$  cy5-labeled sAB-4-SP conjugate (red) followed by staining with TMR-phalloidin (green). (Scale bar,  $10 \mu\text{m}$ .)

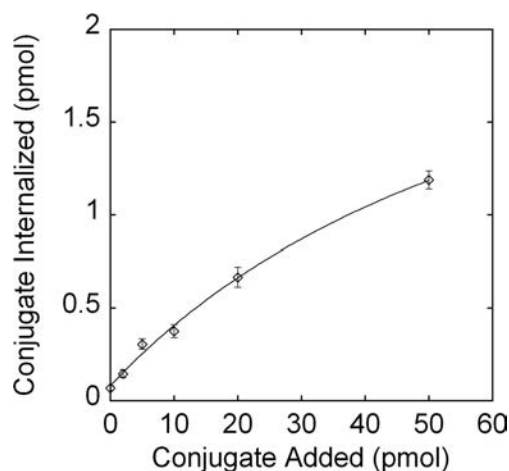
SPv-sAB-27 (Fig. 4B), suggesting that the SPv-NK1R system may be useful in the treatment of several cancers.

**Quantitative Analysis of SPv-mediated Internalization of sAB-27.** To determine the efficiency of SPv-mediated delivery of proteins, U87 cells were incubated with increasing concentrations of a cy5-labeled SPv sAB-27 conjugate for 24 h in a 96-well plate. The cells were washed extensively with PBS and residual fluorescence was determined, reflecting the amount of internalized conjugate. The results show a correlation between the amount of conjugate added and the concentration of the internalized conjugate that fits a hyperbolic response (Fig. 5). Addition of as little as 2 pmol

of the conjugate results in the retention of approximately 0.15 pmoles over roughly 5,000 cells in each well. Taking into account the average volume of U87 cells ( $500\text{--}1,000 \mu\text{m}^3$ ), we estimate that the intracellular concentration of the conjugate is in the mid-micromolar range ( $30\text{--}60 \mu\text{M}$ ). This indicates that even a



**Fig. 4.** Cell viability assay of U87 glioblastoma cells. (A) Forty-eight hour treatment with sAB27, SP or sAB-27-SPv conjugate; left panel,  $10 \text{ nM}$ ; right panel,  $20 \text{ nM}$ . (B) Time course of inhibition of cell viability by the sAB27-SPv conjugate; open circles,  $10 \text{ nM}$ ; closed circles,  $20 \text{ nM}$ .



**Fig. 5.** Quantification of sAB-SPv conjugate internalization by U87 cells. Cells plated on a 96-well plate were treated with increasing amounts of sAB-27-SPv conjugate fluorescently labeled with cy5. After 24 h incubation, the cells were extensively washed, and the amount of the internalized conjugate was calculated from fluorescence values using a standard curve.

small fraction ( $\approx 1\%$ ) of the conjugate escaping the endosome would be comparable to the concentrations used for the *in vitro* studies, and would therefore, be sufficient to elicit a similar effect. It is important to note that the fit suggests an upper limit to the ability of U87 cells to internalize the SPv-conjugates with saturation at high concentrations of added conjugate. This apparent limit may be due to downregulation of the NK1R upon internalization of as a result of SPv uptake.

## Discussion

We have developed an efficient and robust methodology that facilitates the delivery of protein cargos into the cytoplasm of live cells. The platform for our approach has been built such that it can be ported to the research community and used “out of the box.” The method utilizes the receptor-mediated internalization of SPv as an efficient transmembrane delivery vehicle, circumventing many of the problems associated with traditional transfection approaches, such as cytotoxicity. While ligand-induced receptor internalization is a ubiquitous mechanism of receptor regulation, the SP-NK1R system offers several distinct advantages over other systems. SP is a short peptide with subnanomolar affinity to the NK1R. Because only the C-terminal portion of SP is required for binding to the receptor, a wide variety of cargos can be linked to the N terminus.

A crucial feature of this RMD system is that enough of the SPv conjugates appear to readily escape the endosome without loss of function. We have tested other ligand-receptor systems that are efficiently internalized, but in many cases the cargo appears to remain trapped in the endosome and are thus biologically inactive. Although it has been suggested that SP is quickly degraded after internalization (16), it is not clear why SPv-conjugated cargos escape the endosomal machinery more efficiently than in other systems. It is possible that by linking cargo to SP the peptide is sufficiently modified in character that its fate is effectively altered. How and why this occurs is beyond the scope of this study. However, essentially all protein-based cargo that we have conjugated to SPv have been successfully delivered to the cytoplasm without loss of function. In complementary studies, we have determined that our engineered SPv can deliver a wide variety of bioactive cargo including nucleic acid-based molecules such as siRNA, shRNA, and DNA, as well as imaging agents and proteins up to at least 400 kDa. While the delivery of RNA/DNA molecules appears to be promising, further work is needed to rigorously profile their bioactivity in live cells.

Combining our SPv delivery technology with the ability to introduce functional sABs into live cells could be a transformative advancement for studying signaling pathways and other intracellular functions. In extensive studies, we have determined that phage display library-sorting protocols can be designed to produce high affinity sABs that recognize a specific protein surface, a particular conformational state or a multicomponent complex. These attributes, combined with the ability to fine-tune the affinity to the target, result in a class of “designer” affinity reagents that can be programmed to carry out a wide range of directed functions within the cell following internalization. Examples of this include the sABs engineered against different forms of actin described here. Importantly, we have observed that sABs retain their function within the cytoplasm for at least several days after internalization. Thus, sABs engineered for live cell imaging, or inhibition or activation of specific junctions in signaling pathways will have an extended functional lifetime in a live cell system.

Another notable feature of the system is its exquisite specificity for targeting certain types of cancer cells. We have shown that internalization of SPv is restricted to cells expressing the NK1R. This receptor is overexpressed in many types of tumors and primary cancers, including breast carcinomas, adenocarcinomas of the colon, astrocytomas, and glioblastomas (17). This ability to discriminate between cancer cells expressing the NK1R

and normal cells suggests that receptor-mediated delivery using SPv as a Trojan Horse for bioactive cargo may have great therapeutic potential. Further, the specific delivery of sABs directed toward cytoplasmic targets within tumor cells potentially changes the paradigm for antibody-based therapies. These therapies may no longer be limited to the current extracellular targets. Thus, using sABs designed to inhibit intracellular signaling nodes, the possibility certainly exists for focusing future antibody therapies toward a much richer set of cancer targets.

## Methods

**Peptide Synthesis of SPv Variants.** All protected amino acids and resin for peptide synthesis were purchased from Peptides International, solvents from Fisher, salts and buffers from Sigma Aldrich. SPv-maleimide synthesis was carried out manually by standard t-Boc methods using *p*-methyl-benzhydrylamine resin to produce a C-terminal amide after HF cleavage. 6-Maleimido-hexanoic acid (Sigma) was used to introduce an N-terminal maleimide with a 6-carbon linker. An additional variant of SP was synthesized that lacks the maleimide moiety but includes a cysteine mutation at position 3 for attachment of thiol-reactive Alexa-555 and was used as a positive control for NK1 receptor-mediated internalization. The peptides were cleaved from the resin using HF and extracted in 50% acetonitrile plus 0.1% trifluoroacetic acid aqueous solution for lyophilization. The purity of the crude peptides was determined using an analytical C-18 reversed-phase column on a Shimadzu 10A-vp and their masses were confirmed by MALDI MS.

**Fluorescence Labeling and SPv Conjugation to sABs.** Actin-binding sABs were engineered by phage display and purified as described (22). Each sAB contained a C-terminal 6-his tag and a cysteine mutation at position Ala-121 of the heavy chain. The purified sABs were dialyzed into 50 mM sodium borate buffer, pH 7.4, for fluorescent conjugation. NHS-ester fluorophores (Cy3 or Cy5; GE Healthcare, Alexa-488 or Alexa-647; Invitrogen) were added to each sAB in a 5-fold excess from a 10 mM stock solution in DMSO. Reactions were carried out at room temperature for 1 h, followed by gel filtration using a PD-10 column (GE Healthcare) equilibrated with 20 mM Mops, 100 mM NaCl pH 6.9 to remove excess fluorophore. The fluorescently-labeled sABs were reacted with a 10-fold excess of SPv-maleimide from a 5 mM stock in DMSO. The reaction was carried out overnight at 4 °C in 20 mM Mops, 100 mM NaCl, pH 6.9, followed by gel filtration using a PD-10 column to remove excess SPv.

**TIRF Microscopy.** Images were collected with a custom-built total internal reflection fluorescence microscope. A 100 $\times$ , 1.45 NA objective (Olympus) and an EMCCD camera (Ixon, Andor Technologies) were used (*SI Methods*).

**Electron Microscopy.** Electron Micrographs were obtained on a FEI Tecnai F30 transmission electron microscope (*SI Methods*).

**Cell Lines, Western Blots, and Transfection.** See *SI Methods*.

**Confocal Microscopy.** For each sample,  $10^4$  U87 cells were seeded on an 18 mm cover-slip in MEM with 10% FBS overnight in a 12-well plate, then starved in MEM with 0.1% BSA for 3 h. The reagent (SPv-sAB conjugate, SP-Fluorophore, or sAB only) was filter-sterilized, diluted to the desired final concentration in MEM 0.1% BSA and added directly to the live cells in each well for 2 h. Untreated cells or cells treated with sAB only served as a negative control. Following incubation with each reagent, the cells were washed 3 times with PBS, fixed in 4% *p*-formaldehyde, permeabilized in TBST and blocked with 10% normal goat serum for 1 h. Actin filaments were stained with FITC-phalloidin (Invitrogen, 1:250 dilution), or TMR-phalloidin (Sigma-Aldrich, 1:3,000 dilution) for 1 h at room temperature. Nuclei were stained with either DAPI (10 mg/mL) or Hoechst reagent (Invitrogen, 1:2,000) for 5 min at room temperature, and samples were mounted using ProLong Gold reagent (Invitrogen). Confocal microscopy images were collected using a Leica SP2 confocal microscope with FITC and Alexa-488 excited by a 488 nm laser, TMR and Cy3 by a 543 nm laser, and Alexa 574 and Cy5 by a 633 laser. Multicolored images were overlaid using the ImageJ software.

**MTS proliferation assay.** CellTiter 96 AQueous assay (Promega) was used to determine cell viability (see *SI Methods*).

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