1	In vivo targeted gene delivery using Adenovirus-antibody molecular glue conjugates
2	Paul J. Rice-Boucher ^{1,2} , Elena A. Kashentseva ¹ , Igor P. Dmitriev ¹ , Hongjie Guo ³ , Jacqueline M. Tremblay ⁴ ,
3	Charles B. Shoemaker ⁴ , David T. Curiel ^{1,2} , Zhi Hong Lu ^{1,†}
4	1: Department of Radiation Oncology, Washington University School of Medicine, St. Louis, MO, USA
5	2: Department of Biomedical Engineering, McKelvey School of Engineering, Washington University in
6	Saint Louis, St. Louis, MO, USA
7	3: Tiger Biologics LLC, St. Louis, MO, USA
8	4: Department of Infectious Disease and Global Health, Tufts Cummings School of Veterinary Medicine,
9	North Grafton, MA, USA
10	+: Correspondence should be addressed to Z. H. L. (<u>zhihonglu@wustl.edu</u>)
11	
12	
13	
14	
15	
16	
17	
18	
19	
20	

21 Abstract:

22	Safe and efficient nucleic acid delivery to targeted cell populations remains a significant unmet need in
23	the fields of cell and gene therapy. Towards this end, we pursued Adenoviral vectors genetically
24	modified with the "DogTag" molecular glue peptide, which forms a spontaneous covalent bond with its
25	partner protein, "DogCatcher". Genetic fusion of DogCatcher to single-domain or single-chain antibodies
26	allowed covalent tethering of the antibody at defined locales on the vector capsid. This modification
27	allowed simple, effective and exclusive targeting of the vector to cells bound by the linked antibody. This
28	dramatically enhanced gene transfer into primary B and T cells in vitro and in vivo in mice. These studies
29	form the basis of a novel method for targeting Adenovirus that is functional in stringent in vivo contexts
30	and can be combined with additional well characterized Adenovirus modifications towards applications
31	in cell engineering, gene therapy, vaccines, oncolytics, and others.
32	
33	
34	
35	
36	
37	
38	
39	
40	
41	

42 Introduction:

43 Effective genetic medicine requires the safe and efficient delivery of nucleic acids to cells and tissues of interest. Decades of research has yielded numerous vehicles for delivery of both DNA and 44 45 RNA, including non-viral liposomes, lipid nanoparticles and others, and viral vectors such as adeno-46 associated viruses (AAVs), Adenoviruses (Ads), and lentiviruses¹⁻³. Despite this, a vector capable of 47 efficient and targeted delivery of genes to cells of interest remains elusive. This requirement is of particular importance to support the deployment of gene editors *in vivo*, as expression of the editing 48 49 machinery in off-target regions may lead to undesirable side-effects. Despite these challenges, direct in 50 vivo gene delivery remains an attractive option, potentially circumventing the costly and complicated 51 procedures required for ex vivo cell engineering and enabling novel therapies deliverable to 52 underserved patient populations⁴⁻⁷. Recognition of this has led to extensive research into *in vivo* T cell engineering, and a single report describing *in vivo* B cell engineering⁸⁻¹⁵. Targeted gene transfer into 53 54 these cell types will thus likely be of strong utility. 55 Our group and others have undertaken extensive research into Adenoviral vectors as nucleic 56 acid delivery vehicles. Ads are non-enveloped double-stranded DNA viruses generally responsible for the

common cold and are amongst the best-studied DNA vectors. Ads have a long track-record in the clinic 57 and have been successfully deployed for cancer immunotherapy and vaccines against infectious disease, 58 with estimated doses delivered for COVID-19 ranging in the hundreds of millions^{16,17}. Ads thus represent 59 60 a potentially low-cost and safe vector for *in vivo* cell and gene therapy. Numerous manuscripts have 61 been published on the targeting of Ads both in vitro and in vivo, but these approaches generally rely on 62 the use of small peptides, adaptors, complicated genetic engineering, or the use of bespoke reagents such as DARPINs or single-domain antibodies (sdAbs)¹⁸⁻²⁰. The ideal delivery vehicle would embody a 63 64 single-component vector capable of being grown to high titers and targeted with easily obtained 65 reagents.

66	With these goals in mind, we pursued the development of Ad vectors modified with
67	SpyCatcher/SpyTag family molecular glues. We demonstrate the production of a Human Adenovirus
68	serotype C5 (Ad5) vector with the "DogTag" peptide genetically incorporated in the fiber protein.
69	Through this moiety we covalently attach various antibody species fused with DogCatcher to the virus
70	surface and show that this linkage results in vector retargeting both in vitro and in vivo in primary B and
71	T cells.
72	
73	<u>Results:</u>
74	Development of molecular glue driven antibody conjugation to the virus capsid:
75	SpyCatcher/SpyTag protein-peptide partners spontaneously form a covalent bond under
76	physiological conditions and have been used in numerous protein engineering studies ^{21,22} . Our group
77	previously demonstrated that Simian Adenovirus serotype 36 (SAd36) could be derivatized with SpyTag
78	at each of the major capsid proteins, including fiber, hexon, and pIX – the fiber protein is responsible for
79	the initial binding of the virus to cells, while the hexon is the main structural protein forming the capsid.
80	pIX interlaces the hexon and stabilizes the overall structure ¹⁸ . These derivatives could be linked with a
81	Cas9-SpyCatcher fusion protein to achieve gene editing <i>in vitro</i> ²³ . We thus hypothesized that a similar
82	approach might be useful for targeting – by linking an antibody on the virus capsid through
83	SpyCatcher/SpyTag chemistry, we might be able drive viral uptake through binding of the antibody to
84	the appropriate receptor on a cell surface.
85	We selected B cells as a first choice for targeting due to our previous work on this cell type and
86	standing interest in achieving <i>in vivo</i> engineering for control of infectious diseases ²⁴ . We initially
87	attempted targeting through our existing SpyTag modified SAd36 vectors using an sdAb targeting
88	murine CD40 fused with SpyCatcher (F8SpC), but found these vectors were completely unable to access
89	the immunocyte population in pilot in vivo studies and were thus likely inappropriate for our ultimate

90 aims. Furthermore, concurrent with this work a study describing Ads engineered with the loop-friendly molecular glue "DogTag" was published^{25,26}. Based on our previous positive results with Ad5 based 91 92 vectors for *in vivo* gene delivery to B cells, we thus decided to design an Ad5 vector incorporating 93 DogTag in the Ad5 fiber protein (Ad5FDgT). We selected the HI loop within the fiber for DogTag 94 insertion, as this site has previously been used for insertion of peptides (Fig. 1a and 1b)^{27,28}. We found 95 that this vector was easily upscaled in standard HEK293 cells and yielded titers comparable to the isogenic unmodified Ad5 vector. SDS-PAGE analysis revealed identical protein band patterns between 96 97 Ad5 and Ad5FDgT, with the exception of the expected band shift from the insertion of DogTag in the fiber (Fig. 1d). We note that the wild-type Ad5 fiber and pllla proteins overlap just below the 70kDa 98 99 protein ladder marker, whereas the DogTag modified fiber separates from the pllla protein due to its 100 larger molecular weight.

101 To pair with this vector, we developed several antibodies as DogCatcher fusions. We initially 102 developed the aforementioned anti-mCD40 F8 sdAb as a DogCatcher fusion (F8DgC), but also developed 103 single-chain fragment variable (scFv) fusions of commercial antibodies targeting murine and human B 104 cell markers, including CD19 and CD20 (Fig. 1c). As controls and to assess targeting in an alternate cell 105 type, we also developed scFv fusions targeting the murine T cell marker CD8 α . To assess if these 106 reagents were able to conjugate the virus fiber capsid protein, we co-incubated Ad5FDgT and different 107 antibody fusions at room-temperature for approximately 2 hours, then ran SDS-PAGE gels to assess the 108 degree of shifting in the fiber protein. Similar to a previous study, we found that the DogTag-DogCatcher 109 pair was highly reactive and that all or nearly all of the fiber-DogTag protein reacted with the antibody fusions (Fig. 1e)²⁵. Furthermore, we found these antibodies retained their ability to bind to the 110 111 appropriate cell type, as determined by flow cytometry (Fig. 1f). 112 In vitro and in vivo analysis of Adenovirus-antibody conjugates:

To assess the ability of our Adenovirus-antibody (Ad-Ab) conjugates to achieve targeted gene 113 114 transfer, we infected primary murine B and T cells with Ad-Ab decorated with antibodies targeting B or T 115 cell restricted cell markers (CD40, CD19, CD20 and CD8α). As expected, we found that gene transfer 116 enhancement occurred only when the appropriate antibody was conjugated on the virus – infectivity did 117 not change when B cells were treated with a virus targeting CD8α, while T cell infection rates were not 118 impacted by conjugation of the virus with CD40, CD19, or CD20 targeting antibodies (Fig. 2b). We also 119 assessed this effect in primary human B cells, and similarly found that gene transfer enhancements only 120 occurred in the presence of the appropriate targeting agent. In all cases we tested a variety of molar 121 ratios of antibody-DogCatcher to virus-DogTag, and often observed a bell curve where very high and 122 very low ratios of antibody resulted in lower gene transfer enhancements. This is consistent with the 123 idea that too much antibody fragment may act as a competitive inhibitor, but too little antibody may not 124 saturate the binding sites on the virus. We did observe several notable exceptions to this bell curve rule, 125 especially in the human B cell samples where the highest ratio of antibody-DogCatcher to virus-DogTag 126 resulted in the highest gene transfer enhancement. This may stem from differences in conjugation rates 127 - some antibodies might require higher excesses to achieve full saturation of the virus surface. It is also 128 possible that differences in receptor density on different cell types could play a role. Further work is 129 needed to fully elucidate this effect. Our results were particular striking in murine T cells and human B 130 cells, where Ad5FDgT without enhancing antibody was almost completely incapable of gene transfer. 131 Encouraged by our in vitro results, we decided to determine if antibody conjugation could 132 enhance specific gene transfer in vivo. We injected C57BL/6J mice retro-orbitally with Ad5FDgT alone or 133 Ad5FDgT conjugated with 1D3DgC, an scFv targeting murine CD19, a B cell restricted cell marker 134 (Ad5FDgT-1D3) (Fig. 3a). We scored splenic B and T cells for reporter gene expression three days later, 135 and found a statistically significant enhancement of gene transfer in the B cell compartment, but not in 136 the T cell compartment (Fig. 3b). This enhancement was found across several of the B cell

137	subpopulations we scored, including memory B cells, marginal zone B cells, and follicular B cells (the
138	gating strategy for these subpopulations can be found in Fig. S1). We did not observe enhancement in
139	the plasmablast population, which may be due to lower CD19 expression on this cell type ²⁹ . We also
140	scored eGFP expression per milligram of tissue in several major organs, including the spleen, liver, lung,
141	heart and kidneys. In the main site of Ad5 tropism <i>in vivo</i> , the liver, we observed a non-statistical trend
142	towards decreased gene transfer in the Ad5FDgT-1D3 group, compared to the Ad5FDgT group (Fig. 3c).
143	We also observed the Ad5FDgT-1D3 group showed statistically reduced lung and spleen eGFP
144	expression, potentially indicating virus conjugation may reduce off-target gene expression in these
145	tissues. Further work is required to elucidate the mechanism of these results.
146	Development and analysis of purified single-component Ad-Ab complexes:
147	Taken together, our data suggested that Ad-Ab complexes were a promising platform for
148	controlling gene transfer and expression both in vitro and in vivo. However, for clinical translation and
149	industrial production a single component vector without excess antibody fusion protein is desirable. This
150	would also allow us to remove any potential confounding effects of excess antibody in our experiments.
151	We therefore attempted to develop a workflow to purify and store purified Ad-Ab complexes (Fig 4a). At
152	the lab scale, density gradient centrifugation is often used to prepare purified Ads. From cell lysates, we
153	therefore used a single ultracentrifugation with cesium chloride gradients to purify Ad5FDgT. We then
154	dialyzed this vector against 1X PBS to prevent any negative effects towards the virus conjugation from
155	excess cesium chloride. We split this sample into equal thirds, and treated one fraction each with PBS, α -
156	mCD19 1D3DgC, or α -mCD8 α YTS169DgC. Conjugation was carried out at 25C for 1 hour, and vectors
157	were ultracentrifuged a second time to separate free antibody from the virus conjugates. Importantly,
158	observation of the vector bands did not reveal any obvious differences between virus treated with PBS
159	and virus conjugated with scFvs (Fig. 4b). After spinning vector samples were dialyzed again against our
160	standard 1X PBS with 10% glycerol and aliquoted and frozen at -80C.

161 We carried out several assays to determine the quality of these purified Ad-Ab complexes. 162 Encouragingly, SDS-PAGE analysis revealed that the antibody-conjugated samples retained an identical 163 protein band pattern to the PBS treated vector, with the exception of the expected complete shift in the 164 fiber-DogTag band (Fig. 4c). Critically, we also were unable to observe any signs of free antibody. This 165 result was further confirmed by a western blot analysis of the fiber protein, which revealed that the vast 166 majority of fiber-DogTag reacted with the antibody-DogCatcher fusions (Fig. 4d). We then carried out in 167 vitro analyses of these viruses and found that gene transfer enhancement from the purified and stored 168 viruses was comparable to freshly prepared Ad-Ab complexes (Fig. S2). 169 Interestingly, we observed that different antibodies required different molar ratios of 170 DogCatcher:DogTag to achieve optimal gene transfer in vitro (e.g., 1D3DgC was most potent in the 10X-171 2.5X range while YTS169DgC was most potent at 0.625X – see Fig. 2). We had hypothesized that this 172 could be due to differences in the ability of each antibody fusion to link with the virus. However, with 173 purified Ad5FDgT completely functionalized with YTS169DgC, we observed that gene transfer was lower 174 than in our previous experiments with a 0.625X DogCatcher:DogTag (Fig S2). These results indicate that, 175 at least for mCD8α targeting antibodies, an un-saturated capsid is superior to a saturated one. The Ad 176 fiber presents as a trimer, with three copies of DogTag displayed in close proximity. Full functionalization 177 of this trimer with antibody apparently results in lower gene transfer to murine T cells, potentially 178 indicating excessive cross-linking of CD8 α is sub-optimal for driving viral uptake. Additional work is 179 needed to understand this effect and determine the minimum number of antibodies per capsid to 180 achieve optimal targeting through CD8a. It would also be intriguing to determine if this effect is specific 181 to T cells, or specific to $CD8\alpha$. 182 We next endeavored to assess these purified complexes in vivo (Fig. 4e). As before, we injected 183 naïve C57BL/6J mice retro-orbitally with either non-conjugated Ad5FDgT, Ad5FDgT-1D3, or Ad5FDgT-

184 YTS169. We conducted three separate experiments, first in female mice, then male, and finally a mixed

group. We again found that functionalization of the virus with 1D3DgC resulted in a roughly two-fold 185 186 increase in gene expression to B cells compared to the parent vector (the gating strategy for the cell 187 types described in this section can be found in **Fig. S3**). We found a similar enhancement in the CD8 α + T 188 cell population, with YTS169DgC functionalization resulting in an approximately two-fold increase in 189 gene expression. Critically, neither modification resulted in an increase in gene expression in off-target 190 cell types – 1D3DgC functionalization did not increase gene expression in CD8 α + T cells, and YTS169DgC 191 did not increase gene expression in B cells. We also did not observe statistical differences in groups 192 separated by sex, confirming that our technology is applicable in both male and female mice (Fig. S4). 193 Interestingly, we did observe a slight, non-significant increase in gene expression in CD4+ T cells 194 in mice injected with Ad5FDgT-1D3 and Ad5FDgT-YTS169 compared to Ad5FDgT. The potential 195 mechanism of this result is not clear – activated CD4+ T cells do express an Fc receptor which could 196 conceivably bind to the antibodies used to functionalize our vector, but our study uses scFvs lacking any 197 Fc receptor binding³⁰. Another explanation could be that incorporation of scFvs into the fiber increases 198 the circulation time of the virus and allows slight increases in transduction of off-target cells. There may 199 also be as-of-yet uncharacterized interactions between the various hematopoietic cells and scFvs or 200 DogCatcher itself that are responsible. Additional work is required to understand this effect. 201 We also assessed liver eGFP expression per milligram of tissue as in our previous assay and 202 selected two mice per group for tissue immunohistochemistry. We did again observe a weak, non-203 statistical decrease in liver gene expression in mice injected with Ad5FDgT-1D3, but this result was not 204 true for mice injected with Ad5FDgT-YTS169. We also confirmed that there were not significant 205 differences in liver eGFP expression in male and female mice (Fig. S4). We further confirmed these 206 results using tissue immunohistochemistry and demonstrated that scFv vector functionalization did not 207 result in any radical changes in overall gene expression across all organs (Fig. S5) - as described in

numerous previous studies, the overall *in vivo* distribution of Ad5 to the liver is clearly dominated by
 binding of the liver to blood factors^{31,32}.

210

211 Discussion:

212 Here we describe the development of a novel platform technology for targeting of Adenovirus 213 to defined receptors using antibodies. We pair a simple, easy-to-produce vector with scFv fusions, enabling rapid design of vectors capable of efficient gene transfer into challenging cell types. The Ad-Ab 214 215 system boosts gene transfer into cells moderately-to-completely resistant to Ad infection – in an exemplary test, murine CD8α+ T cells infected with unmodified Ad5FDgT at 25,000 vp/cell showed less 216 217 eGFP expression than those infected with just 40 vp/cell of Ad5FDgT-YTS169 (data not shown). In vitro, 218 Ad-Ab is thus a highly useful platform for gene delivery into primary cell types, especially in contexts 219 where mixed cell populations are present or the cell type is very difficult to infect with conventional 220 vectors.

221 Ad-Ab offers several key advantages over conventional Adenovirus targeting techniques. A 222 classic example is the swapping of fiber knob domains from other Ad serotypes onto the Ad5 fiber tail 223 and shaft, resulting in transfer of tropism from the swapped serotype. This strategy has been extensively 224 used for hematopoietic stem cell (HSC) engineering, wherein swapping the Ad5 fiber for the fiber from Adenovirus serotype 35 results in vector targeting to CD46, which is present on HSCs³³⁻³⁵. Although this 225 226 technique has been highly successful, there is a limited repertoire of serotypes available for fiber 227 swapping, and many bind to proteins such as the coxsackie and adenovirus receptor, CD46, or desmoglein-2 which are expressed in many tissues³⁶. Fiber swapping is thus a limited technique for 228 229 targeting defined cell populations. In comparison, Ad-Ab can theoretically be targeted to any receptor 230 for which an antibody has been described.

231 To overcome these limitations and target specific cell receptors, many groups have genetically engineered the Ad5 fiber to incorporate small peptides. ligands and sdAbs¹⁸⁻²⁰. Although these 232 233 approaches can be successful, extensive editing of the viral capsid often results in reduced titers and 234 infectious particle ratios. In our own hands, we have generated numerous vectors genetically modified 235 to express sdAbs at the fiber^{19,37-39}. Although these efforts have yielded successes, we typically needed 236 to attempt production of many sdAb variants to obtain a vector which was able to be grown to high 237 titers. Additionally, there was no guarantee that successfully produced sdAb modified vectors would 238 bind an accessible epitope in vivo – many vectors which displayed remarkable infectivity enhancements 239 in vitro did not translate. The relative sparsity of available sdAbs also required us to develop novel 240 variants for many antigen targets, a lengthy and costly process involving immunization of large animals. 241 The ability to use the sequences of commercially available and characterized antibodies with validated in 242 vivo effects to generate scFvs is thus a major advantage to the Ad-Ab platform.

243 To circumvent issues with viral titers and infectivity, many groups have attempted the use of 244 adaptors to link antibodies or other targeting agents onto the vector surface – for example, our group 245 has fused the soluble domain of the coxsackie and adenovirus receptor to various targeting agents in the 246 past^{40,41}. This protein is the endogenous receptor for Ad5 and binds the fiber protein with high affinity. A 247 genetic fusion between coxsackie and adenovirus receptor and an antibody or ligand thus allows for 248 vector retargeting. Another similar approach involves the use of a trimerized DARPin which binds the 249 fiber knob fused to targeting DARPins⁴². While these systems show promise *in vitro*, the complex and 250 challenging environment *in vivo* could break the interaction between the vector and the adaptor, 251 leading to a loss of targeting and unexpected effects. Furthermore, such a two-component system may 252 be challenging to translate clinically – a single component, permanently linked system which can be 253 generated and processed in a similar manner to standard Ad vectors is thus a major advantage.

The use of molecular glue technology thus provides a solution to these issues and enables a simple targeting strategy. Other groups have also recognized this utility – SpyTag has been used to target lentivirus and was recently described for Ad targeting of a cancer cell line using an sdAb^{43,44}. Our study validates and builds upon this work by swapping SpyTag for the highly reactive DogTag and demonstrating the use of scFvs for targeting primary cell types *in vitro* and *in vivo*, thus showing the potential of molecular glue driven targeting for clinical applications.

An inherent limitation of Ad5 based vectors, including our Ad-Ab system, is viral particle 260 261 sequestration in the liver. This sequestration, at least for Ad5, has been demonstrated to be largely mediated by binding of Factor X to specific residues in the hexon protein^{31,32}. Our group and others have 262 thus developed numerous vectors capable of escaping liver sequestration through the use of targeted 263 mutations in the hexon⁴⁵⁻⁴⁷. We anticipate the flexibility of the Ad-Ab system will allow us to combine 264 265 the fiber targeting described here with additional capsid mutations – we have already been successful in 266 developing a next-generation Ad-Ab vector with hexon modifications described by Atasheva et al to 267 ablate liver and macrophage sequestration⁴⁷. Similarly, a recent study incorporated DogTag in the hexon to link the virus with SARS-CoV-2 antigens to generate a novel vaccine platform²⁵. Intriguingly, this group 268 269 also reported that hexon functionalization with antigen blocked antibody mediated neutralization of the 270 vector and binding of Factor X. Development of dual-tag modified vectors with DogTag at the hexon and 271 fiber may also be of utility, as placing scFvs at both sites might lead to targeting through the fiber with 272 simultaneous blocking of sequestration factors through the hexon.

Finally, we also previously demonstrated that SAd36 can be derivatized with SpyTag at various locales and used to deliver a SpyCatcher-Cas9 fusion protein to achieve gene editing²³. Delivery of Cas9 as a protein rather than DNA may carry advantages such as reduced off-target effects due to its transient nature. Towards the combination of these systems, we have developed a variant of Ad5FDgT with SpyTag inserted at the C-terminus of the pIX protein, potentially allowing for targeting through

fiber and delivery of Cas9 ribonucleoprotein through pIX. Further characterization of all the described
vectors are of interest.

280 In total we present here a highly flexible and efficient platform for the transfer of genes to 281 precisely targeted cell populations. We believe Ad-Ab adds to the toolbox of novel targeted vectors for 282 gene delivery and may be of use for gene editing, especially in vitro in contexts where specificity and 283 efficiency are paramount. With further modifications to the Ad-Ab capsid, we also anticipate strong 284 utility for *in vivo* precision gene transfer. This system thus represents our initial steps on a path towards 285 low-cost, targeted and safe gene therapy *in vivo* using Ad vectors. 286 287 Methods: Cell Lines: 288 289 HEK293 (ATCC CRL-1573) cells were grown in Dulbecco's Modified Eagle Medium/Ham's F12 1:1 290 mixture supplemented with L-glutamine, 15mM HEPES, 10% fetal bovine serum (FBS) and 100U/mL

291 penicillin-streptomycin. Cells were grown at 37C with 5% CO2 under sterile conditions.

292 <u>Viruses:</u>

293 A previously described first-generation E1/E3 deleted HAdV-C5 vector with the cytomegalovirus 294 (CMV) promoter driving eGFP (Ad5.CMVeGFP) was used to generate $Ad5FDgT^{24}$. The parent plasmid was 295 cut with Barl (Sibenzyme) and BstBl (New England Biolabs) to release the fiber protein. Two PCR 296 fragments were generated encoding the regions upstream and downstream of the HI loop domain with 297 overlaps for the Barl and BstBl cut sites. A synthetic fragment encoding DogTag with short linkers and 298 overlaps for the PCR fragment was synthesized by Integrated DNA Technologies (IDT). These three 299 fragments were assembled with the cut Ad5.CMVeGFP backbone using NEB HiFi DNA Assembly (New 300 England Biolabs), generating Ad5FDgT. This backbone was linearized using PacI and transfected into

301 HEK293 cells for upscale. Viruses were purified and analyzed for viral particle concentration as

302 previously reported⁴⁸.

303 Protein Constructs:

304 The single domain antibody JPP-F8 was identified from the blood lymphocytes of two alpacas 305 immunized with murine CD40 protein employing general methods previously reported in detail⁴⁹. The 306 JPP-F8 VHH was purified and evaluated by standard dilution ELISA (cite PMID: 33774040) and shown to bind murine CD40 with a sub-nM EC_{50}^{49} . pDEST14-F8DgC was derived by cloning of a synthesized DNA 307 308 fragment (IDT) containing the camelid JPP-F8 single domain antibody followed by a (G4S)3 flexible linker 309 into the Sfol site of pDEST14-DogCatcher (a gift from Mark Howarth, Addgene plasmid #171772; 310 http://n2t.net/addgene:171772; RRID:Addgene 171772), and the resultant sequences encoded "6xHis-TEV-JPPF8-DogCatcher²⁶. pcDNA3.4-18B12DgC and pcDNA3.4-Rtxv1DgC were derived by cloning of 311 312 synthesized DNA fragments (IDT) containing 18B12 and Rtxv1 scFvs together with a DgC fragment into 313 the EcoRI and HindIII sites of pcDNA3.4-c-Fos scFv [N486/76] (a gift from James Trimmer, Addgene 314 plasmid # 190560 ; http://n2t.net/addgene:190560 ; RRID:Addgene 190560), and the resultant 315 sequences encoded "IL-2 signal sequence-18B12 scFV-(G4S)3-DgC-6xHis" and "IL-2 signal sequence-316 Rtxv1 scFV-(G4S)3-DgC-6xHis⁷⁵⁰. pcDNA3.4-1D3DgC and pcDNA3.4-FMC163DgC were derived by cloning 317 of synthesized DNA fragments containing 1D3 scFv and FMC163 scFv into the EcoRI and BamHI sites of 318 pcDNA3.4-Rtxv1DgC, and the resultant sequences encoded "IL-2 signal sequence-1D3 scFv-(G4S)3-DgC-319 6xHis" and "IL-2 signal sequence-FMC63 scFv-(G4S)3-DgC-6xHis". pcDNA3.4-HA22DgC, pcDNA3.4-320 1YTS169DgC pcDNA3.4-2.43DgC were derived by cloning of synthesized DNA fragments containing 321 IGHV1-46 signal sequence followed by HA22 svFv, YTS169 scFv, and 2.43 scFv into Xbal and BamHI sites 322 of pcDNA3.4-1D3DgC, and resultant sequences encoded "IGHV1-46 signal sequence-HA22 scFv-(G4S)3-323 DgC-6xHis", "IGHV1-46 signal sequence-YTS169 scFv-(G4S)3-DgC-6xHis", and "IGHV1-46 signal 324 sequence-2.43 scFv-(G4S)3-DgC-6xHis".

325 <u>Recombinant Protein Production:</u>

326	The plasmids pDEST14-F8DgC and pDEST14-DogCatcher were introduced into protein expression
327	BL21(DE3)-RIPL E. coli cells. Single colonies were used to inoculate 25 mL starter LB containing 100
328	μ g/mL carbenicillin and 50 μ g/ml chloramphenicol grown at 37 °C overnight. The starter cultures were
329	added to 500 ml fresh media without antibiotics, and cultures were grown at 37 $^\circ$ C with shaking at 250
330	rpm for 2.5 hours. Protein expression was induced with 1 mM IPTG, and the cultures were incubated at
331	30 °C with shaking at 250 rpm for 4 hours. Cultures were centrifuged, and cell pellets were resuspended
332	in lysis buffer (0.5 mM Tris, 0.3 M NaCl, 10 mM imidazole, 0.2% Triton X-100, 1 mg/ml lysozyme, 20
333	units/ml DNase I, 1 mM PMSF, and one complete mini EDTA-free protease inhibitor cocktail tablet per
334	10 ml) and incubated at 37 $^\circ$ C for 30 minutes. The cell lysates were clarified by centrifugation at 32,000
335	rcf at 4° C for 30 minutes.
336	For mammalian recombinant protein production, 80-90% confluent 293T cells were transfected
337	with pcDNA3.4-based protein expression plasmids in the presence of transporter 5 reagent
338	(Polysciences, Inc.). The transfected cells were cultivated in DMEM medium containing 10% fetal bovine
339	serum for 4 to 6 hours and switched to FreeStyle 293 medium for additional 4 to 5 days. The culture
340	supernatants were collected and concentrated with Amicon Ultra-15 with 10K NMWL.
341	The 6xHis-tagged recombinant proteins produced in bacterial and mammalian systems were purified
342	using a HisPur Ni-NTA column with 20 to 40 mM imidazole washing buffer and 300 mM imidazole
343	elution buffer, and eluted proteins were dialyzed in 10% glycerol in PBS with three buffer changes using
344	3.5KDa molecular weight cut-off Slide-A-Lyzer Dialysis Cassettes. Protein concentration was measured
345	using BCA according to the manufacturer's instructions (Thermo Scientific).
346	SDS-PAGE and Western Blot:

Whole virus protein analysis and conjugation analysis was carried out using SDS-PAGE. Viruses
and proteins were incubated at a 2:1 ratio with 3x SDS sample buffer containing 187.5 mM Tris-HCl, 6%

SDS, 30% glycerol, 0.125M dithiothreitol and 0.03% bromophenol blue at pH 6.8 for 15 minutes at 100°
C. Samples were then loaded on a 4-15% gradient gel and resolved using a Criterion electrophoresis
system (Bio-Rad). Staining was carried out using GelCode Blue according to the manufacturer's protocol
(Thermo Scientific).

353 Western blot analysis was carried out to detect the Ad5FDgT fiber protein and its 354 modified/conjugated derivatives. The samples containing about 6×10^{11} viral particles were mixed 1:1 355 with the 2X Laemmli SDS-PAGE loading buffer containing 2-mercaptoethanol (Sigma) and heated in 356 boiling water for 5 minutes to denature the viral proteins. The denatured samples were run on 4 - 20% 357 Tris-Glycine Mini protein gel (Invitrogen) using Novex Tris-Glycine SDS Running Buffer (Invitrogen) as 358 recommended by the manufacturer. The iBlot 2 Dry Blotting System (Invitrogen) was used to transfer 359 electrophoretically resolved viral proteins from the gel to PVDF membrane (Invitrogen) as 360 recommended by the manufacturer. We employed the iBind Western System (Invitrogen) for 361 immunodetection of Ad5FDgT fiber proteins using the primary mouse monoclonal antibody 4D2 against 362 the N-terminal fiber tail domain. 4D2 was a kind gift from Jeff Engler and was produced by the 363 Hybridoma and Monoclonal Immunoreagent Core at the University of Alabama, Birmingham⁵¹. Secondary detection was carried out using anti-mouse IgG conjugated with Alkaline Phosphatase (AP) 364 365 (Sigma). The protein bands bound with both primary and secondary antibody were developed with 366 colorimetric AP substrate reagent kit (Bio-Rad) as recommended by the manufacturer. 367 Primary Cell Culture: 368 Primary mouse and human B cell culture and infection was carried out as previously described

by our group²⁴. Briefly, mouse B cells were magnetically isolated from splenocytes and cultured for 30h
 in RPMI 1640 supplemented with 10% FBS, 1X Nonessential Amino Acids, 1X sodium pyruvate and 1X 2 mercapto-ethanol. 50 µg/mL LPS was used as an activation agent. Infections were carried out overnight
 with 5x10⁵ cells in 50 µL LPS-free media with 5% FBS. Infected cells were then returned to 500 µL total

volume with complete media and incubated for a total of 48 hours after infection prior to flow

374 cytometry analysis.

375	Human B cells were magnetically isolated from healthy donor peripheral blood mononuclear
376	cells and cultured for approximately 72 hours prior to infection according to previously described
377	protocols ^{24,52-54} . For infections 2.5x10 ⁵ cells were infected in a total volume of 40 μ L STEMMacs HSC
378	Expansion media (Stem Cell Technologies) with 0.2% FBS for 3 hours, then transferred to 1mL complete
379	media. Flow cytometry analysis was carried out 48 hours after infection.
380	Mouse T cells were isolated from splenocytes using the Miltenyi Pan T Cell Isolation Kit II
381	according to the manufacturer's instructions. Cells were activated for 1 hour prior to infection using the
382	Miltenyi T Cell Activation/Expansion kit according to the manufacturer's instructions. 1-2x10 ⁶ T cells
383	were incubated with 200 μ L anti-CD3/CD28 microbeads in 10mL RPMI 1640 supplemented as above and
384	with 10ng/mL mouse IL-4. For infections $5x10^5$ cells were resuspended in 100 μ L culture media and
385	infected for 16-20 hours. An additional 150 μL complete media was then added and cells were analyzed
386	via flow cytometry 48 hours after infection.
387	Virus Conjugation:
388	For in vitro experiments, virus was prepared at 2X the concentration required for infection and
389	mixed with an equal volume of the required amount of antibody fusion protein diluted in 1X PBS. This
390	mixture was incubated at room temperature or 25° C for about 2 hours prior to infection.
391	For in vivo experiments, the required amount of virus for all injections was incubated with the required
392	amount of antibody without further dilution for 2 hours at room temperature or 25 $^\circ$ C. 1X PBS was then
393	added to bring the conjugated virus to the appropriate final volume.
394	Flow Cytometry:

For analysis of antibody-DogCatcher fusions, splenocyctes isolated from C57BI/6J mice (about 1
 million cells) were stained in 100 μL FACS buffer (1X PBS containing 0.5% BSA) with 0.5 μg of DgC,

397 1D3DgC, 18B12DgC, 2.43DgC, or PBS control at 4° C for 30 minutes. Samples were then washed with 2 ml FACS buffer and resuspended in 100 µl FACS buffer containing 0.5 µl FITC anti-His Tag antibody 398 399 (BioLegend) and 0.5 µl Alexa Fluor[®] 594 anti-mouse B220 antibody (BioLegend) or 0.5 µl Alexa Fluor[®] 400 594 anti-mouse CD8a antibody (BioLegend). Samples were incubated at 4° C for 30 minutes, washed 401 again in 2 ml FACS buffer then resuspended in 100 µl FACS buffer. For blocking studies, splenocytes 402 were pretreated with 7.5 µg or 15.0 µg full length 1D3 antibody (BioLegend) at 4° C for 30 minutes 403 before the addition of 1D3DgC. 404 For analysis of *in vitro* infected murine B cells, samples were stained with anti-CD19 (Invitrogen 405 #RM7717) and Fixable Far Red live/dead dye (Invitrogen). Samples were gated as singlets/live/CD19+. 406 Human B cells were stained with anti-CD19 (BioLegend) and Fixable Far Red dye. Samples were gated as 407 above. Mouse T cells were stained with anti-CD8 α (BioLegend) and Fixable Far Red dye or Sytox Red dye 408 (Invitrogen). Samples were gated as singlets/live/CD8 α +. 409 For *in vivo* studies, spleens were processed into single cell suspensions and resuspended in 1mL 410 1X PBS with 2% FBS. 100uL of the suspension was used for analysis and was pre-incubated with 10 µL Fc 411 Blocking Reagent (Miltenyi) for 10 minutes prior to staining. In all cases live/dead discrimination was 412 carried out using Fixable Far Red dye. For the first in vivo assay (Fig. 3), B cells were gated as 413 singlets/live/CD19+/CD3-, while T cells were gated as singlets/live/CD3+/CD19-. Germinal center B cells 414 were gated as GL7+/CD95+/IgDlo/CD38lo, marginal zone as IgM+/IgDlo, follicular as IgMlo/IgD+, 415 memory as IgDlo/GL7-/CD38+, and plasmablasts as CD138+/IgD-. All B cell subsets were first passed through a singlets/live/CD19+ gate prior to further analysis. 416 417 For the second in vivo assay (Fig. 4), B cells were gated as singlets/live/CD19+/CD3-/CD22+/CD4-418 /CD8-. CD8a+ T cells were gated as singlets/live/CD8a+/CD19-/CD3+/CD22-/CD4-. CD4+ T cells were 419 gated as singlets/live/CD4+/CD19-/CD3+/CD22-/CD8α-.

- 420 Mouse antibodies used were as follows: CD19 (Invitrogen or Miltenyi), CD8α (BioLegend), CD3
- 421 (Invitrogen), GL7 (BioLegend), CD95 (Invitrogen) IgD (Invitrogen), CD38 (Miltenyi), IgM (Invitrogen),
- 422 CD138 (Invitrogen), CD22 (BioLegend), CD4 (Invitrogen).
- 423 All analysis was carried out using FlowJo 10.10.0.
- 424 Animal Studies:

425 C57BI/6J mice were acquired from the Jackson Laboratory and housed in a pathogen-free environment. Mice aged 6-9 weeks were injected retroorbitally with 5×10^{10} viral particles in 150 μ L total 426 427 volume and sacrificed roughly 72 hours later via anesthetization with Avertin followed by cervical 428 dislocation. In the first in vivo study, one-half of each spleen was used for flow cytometry analysis, while 429 the remaining half, liver, lungs, kidneys and heart were snap frozen in liquid nitrogen. Organs were 430 thawed on ice and processed for eGFP quantification using a Fluorimetric GFP Quantitation Kit (Cell 431 Biolabs) per the manufacturer's instructions. Tissues were homogenized in the included lysis buffer 432 supplemented with 10% Proteinase Inhibitor Cocktail (Sigma) using an Omniprep 96 automated 433 homogenizer. Tissue lysated were then centrifuged at 4000-5000 rpm to remove cell debris and 434 supernatants were transferred to clean tubes. Supernatants were diluted as appropriate then aliquoted into 96-well plates for eGFP quantification. Total solution eGFP was normalized against total solution 435 protein from a BCA assay of the lysates performed according to the manufacturer's instructions (Thermo 436 437 Scientific).

In the second *in vivo* study, the whole spleen of each animal was used for flow cytometry and whole livers were used for tissue eGFP quantitation as above. All experiments were approved by the Institutional Animal Care and Use Committee of the Washington University in St. Louis School of Medicine (Protocol #22-0360) and were performed in accordance with the National Institutes of Health Guide for Care and Use of Laboratory Animals. All efforts were made to minimize suffering and the total number of animals used in the study.

444 <u>Purification of Ad-Ab complexes:</u>

445	Cells from 20X infected T175 flasks were harvested after the development of cytopathic effect
446	and centrifuged at 1200 rpm for 5 minutes. Supernatants were removed and cell pellets were frozen at -
447	80° C. Samples were subjected to 3 freeze-thaw cycles in room-temperature water and dry ice to lyse
448	the cells, then centrifuged at 4000 rpm for 10 minutes to clarify the lysates. Supernatants were then
449	ultracentrifuged on cesium chloride gradients for 2 hours at 25,000 rpm at 4 $^\circ$ C. Viral bands were
450	harvested and dialyzed twice against 1X PBS. Virus was removed from dialysis and viral particles were
451	quantified as above. Virus was then split into three equal aliquots and treated with either PBS, two-fold
452	molar excess of 1D3DgC, or two-fold molar excess of YTS169DgC for 1 hour at 25 $^\circ$ C. Conjugated viruses
453	were ultracentrifuged again on cesium chloride gradients for 1.5 hours at 25,000 rpm at 4 $^\circ$ C. Viral
454	bands were harvested and dialyzed three times against 1X PBS with 10% glycerol, then frozen at -80 $^\circ$ C.
455	Statistical Analyses:
456	Specific methods used for analysis are noted in the corresponding Figure legends. In all cases
457	statistical analysis was carried out using GraphPad Prism 10. A p value of < 0.05 was used and
458	significance is indicated as *: p < 0.0322, **: p < 0.0021, ***: p < 0.0002, **** p < 0.0001, ns: not
459	significant. In all cases error bars correspond to standard deviation.
460	
461	Data availability statement:
462	Plasmids used in this study are to be deposited in AddGene. Flow cytometry data is available
463	upon reasonable request. All other data can be found in the main text or Supplemental.
464	
465	Acknowledgements:
466	The authors would like to thank Mark Selby, Rosa Romano, and Hongil Park of Walking Fish

467 Therapeutics, Inc for their contributions to human B cell culturing and analysis. This work was supported

468	by National Institute of Health grants 1R21EB033459-01A1 and 1R01AI174270-01A1 awarded to David T			
469	Curiel, 1R21HL166887-01A1 awarded to Zhi Hong Lu, and T32HL007088-45 awarded to Stephen Oh.			
470	Additional funding for this work was provided by Walking Fish Therapeutics, Inc through award P21-			
471	04949 to Zhi Hong Lu.			
472				
473	CrediT author contributions:			
474	Paul J. Rice-Boucher: Conceptualization, formal analysis, investigation, visualization,			
475	methodology, writing – original draft. Elena A. Kashentseva: Methodology, investigation. Igor P.			
476	Dmitriev: Methodology. Hongjie Guo: Methodology, investigation. Jacqueline M. Tremblay: Resources.			
477	Charles B. Shoemaker: Resources. David T. Curiel: Conceptualization, supervision, writing – review and			
478	editing. Zhi Hong Lu: Supervision, conceptualization, formal analysis, visualization, investigation,			
479	methodology, writing – review and editing.			
480				
481	Declaration of interests:			
482	Hongjie Guo is the founder and Chief Scientific Officer of Tiger Biologics, LLC, a protein			
483	production company. Paul J. Rice-Boucher. David T. Curiel, and Zhi Hong Lu are co-inventors on a patent			
484	application describing the use of the Ad-Ab system for B cell targeting and engineering.			
485				
486	Keywords:			
487	Gene delivery, adenovirus, molecular glue, SpyTag, SpyCatcher, DogTag, DogCatcher			
488				
489	References:			
490 491	Bulcha, T. J., Wang, Y., Ma, H., Tai, L. W. P. & Gao, G. Viral vector platforms within the gene therapy landscape. <i>Signal Transduction and Targeted Therapy</i> 6 (2021).			

492 https://doi.org/https://doi.org/10.1038/s41392-021-00487-6

493 494 495	2	Lostalé-Seijo, I. & Montenegro, J. Synthetic materials at the forefront of gene delivery. <i>Nature Reviews Chemistry</i> 2 , 258-277 (2018). <u>https://doi.org/https://doi.org/10.1038/s41570-018-0039-1</u>
496 497	3	Dunbar, E. C. <i>et al.</i> Gene therapy comes of age. <i>Science</i> 359 , eaan4672 (2018). <u>https://doi.org/https://doi.org/10.1126/science.aan4672</u>
498 499 500	4	Wang, D., Zhang, F. & Gao, G. CRISPR-Based Therapeutic Genome Editing: Strategies and In Vivo Delivery by AAV Vectors. <i>Cell</i> 181 , 136-150 (2020). <u>https://doi.org/doi</u> : 10.1016/j.cell.2020.03.023
501 502 503	5	Wilbie, D., Walther, J. & Mastrobattista, E. Delivery Aspects of CRISPR/Cas for in Vivo Genome Editing. <i>Accounts of Chemical Research</i> 52 , 1555-1564 (2019). <u>https://doi.org/https://doi.org/10.1021/acs.accounts.9b00106</u>
504 505 506	6	Papathanasiou, M. M. <i>et al.</i> Autologous CAR T-cell therapies supply chain: challenges and opportunities? <i>Cancer Gene Therapy</i> 27 , 799-809 (2020). <u>https://doi.org/https://doi.org/10.1038/s41417-019-0157-z</u>
507 508	7	Li, C. & Lieber, A. Adenovirus vectors in hematopoietic stem cell genome editing. <i>FEBS Lett</i> 593 , 3623-3648 (2019). <u>https://doi.org/10.1002/1873-3468.13668</u>
509 510	8	Hamilton, R. J. <i>et al.</i> In vivo human T cell engineering with enveloped delivery vehicles. <i>Nature Biotechnology</i> (2024). <u>https://doi.org/https://doi.org/10.1038/s41587-023-02085-z</u>
511 512 513	9	Nahmad, D. A. <i>et al.</i> In vivo engineered B cells secrete high titers of broadly neutralizing anti-HIV antibodies in mice. <i>Nature Biotechnology</i> 40 , 1241-1249 (2022). <u>https://doi.org/doi</u> : 10.1038/s41587-022-01328-9
514 515	10	Agarwal, S. <i>et al.</i> In Vivo Generation of CAR T Cells Selectively in Human CD4+ Lymphocytes. <i>Molecular Therapy</i> 28 , 1783-1794 (2020). <u>https://doi.org/doi</u> : 10.1016/j.ymthe.2020.05.005
516 517 518	11	Agarwal, S., Weidner, T., Thalheimer, B. F. & Buchholz, J. C. In vivo generated human CAR T cells eradicate tumor cells. <i>Oncolmmunology</i> 8 , e1671761 (2019). https://doi.org/https://doi.org/10.1080/2162402X.2019.1671761
519 520 521	12	Ho, N. <i>et al.</i> In vivo generation of CAR T cells in the presence of human myeloid cells. <i>Molecular Therapy - Methods & Clinical Development</i> 26 , 144-156 (2022). https://doi.org/DOI:https://doi.org/10.1016/j.omtm.2022.06.004
522 523 524	13	Michels, R. K. <i>et al.</i> Preclinical proof of concept for VivoVec, a lentiviral-based platform for in vivo CAR T-cell engineering. <i>Journal for ImmunoTherapy of Cancer</i> 11 , e006292 (2023). <u>https://doi.org/10.1136/jitc-2022-006292</u>
525 526 527	14	Pfeiffer, A. <i>et al.</i> In vivo generation of human CD19-CAR T cells results in B-cell depletion and signs of cytokine release syndrome. <i>EMBO Molecular Medicine</i> 10 , e9158 (2018). https://doi.org/https://doi.org/10.15252/emmm.201809158
528 529	15	Rurik, G. J. <i>et al.</i> CAR T cells produced in vivo to treat cardiac injury. <i>Science</i> 375 , 91-96 (2022). <u>https://doi.org/10.1126/science.abm0594</u>
530 531	16	Baker, A., Aguirre-Hernández, C., Halldén, G. & Parker, A. Designer Oncolytic Adenovirus: Coming of Age. <i>Cancers</i> 10 , 201 (2018). <u>https://doi.org/doi</u> : 10.3390/cancers10060201

532 533 534	17	Coughlan, L., Kremer, J. E. & Shayakhmetov, M. D. Adenovirus-based vaccines—a platform for pandemic preparedness against emerging viral pathogens. <i>Molecular Therapy</i> 30 , 1822-1849 (2022). https://doi.org/https://doi.org/10.1016/j.ymthe.2022.01.034
535	18	Beatty, S. M. & Curiel, T. D. in <i>Applications of viruses for cancer therapy</i> 39-67 (2012).
536 537 538	19	Kaliberov, A. S. <i>et al.</i> Adenoviral targeting using genetically incorporated camelid single variable domains. <i>Laboratory Investigation</i> 94 , 893-905 (2014). <u>https://doi.org/https://doi.org/10.1038/labinvest.2014.82</u>
539 540 541	20	Smith, N. S. <i>et al.</i> The SHREAD gene therapy platform for paracrine delivery improves tumor localization and intratumoral effects of a clinical antibody. <i>Proceedings of the National Academy of Sciences</i> 118 , e2017925118 (2021). <u>https://doi.org/doi</u> : 10.1073/pnas.2017925118
542 543	21	Keeble, H. A. & Howarth, M. Power to the protein: enhancing and combining activities using the Spy toolbox. <i>Chemical Science</i> 11 , 7281-7291 (2020). <u>https://doi.org/DOI</u> : 10.1039/D0SC01878C
544 545 546	22	Zakeri, B. <i>et al.</i> Peptide tag forming a rapid covalent bond to a protein, through engineering a bacterial adhesin. <i>Proceedings of the National Academy of Sciences</i> 109 , E690-E697 (2012). https://doi.org/https://doi.org/10.1073/pnas.1115485109
547 548 549 550	23	Lu, H. Z., Li, J., Dmitriev, P. I., Kashentseva, A. E. & Curiel, T. D. Efficient Genome Editing Achieved via Plug-and-Play Adenovirus Piggyback Transport of Cas9/gRNA Complex on Viral Capsid Surface. <i>ACS Nano</i> 16 , 10443-10455 (2022). <u>https://doi.org/DOI</u> : 10.1021/acsnano.2c00909
551 552	24	Rice-Boucher, J. P. <i>et al.</i> Adenoviral vectors infect B lymphocytes in vivo. <i>Molecular Therapy</i> 31 , 2600-2611 (2023). <u>https://doi.org/DOI:https://doi.org/10.1016/j.ymthe.2023.07.004</u>
553 554 555	25	Dicks, D. J. M. <i>et al.</i> Modular capsid decoration boosts adenovirus vaccine-induced humoral immunity against SARS-CoV-2. <i>Molecular Therapy</i> 30 , 3639-3657 (2022). https://doi.org/DOI:https://doi.org/10.1016/j.ymthe.2022.08.002
556 557 558	26	Keeble, H. A. <i>et al.</i> DogCatcher allows loop-friendly protein-protein ligation. <i>Cell Chemical Biology</i> 29 , 339-350.e310 (2022). https://doi.org/https://doi.org/10.1016/j.chembiol.2021.07.005
559 560 561 562	27	Dmitriev, I. <i>et al.</i> An Adenovirus Vector with Genetically Modified Fibers Demonstrates Expanded Tropism via Utilization of a Coxsackievirus and Adenovirus Receptor-Independent Cell Entry Mechanism. <i>Journal of Virology</i> 72 , 9706-9713 (1998). <u>https://doi.org/DOI</u> : <u>https://doi.org/10.1128/jvi.72.12.9706-9713.1998</u>
563 564 565	28	Krasnykh, V. <i>et al.</i> Characterization of an Adenovirus Vector Containing a Heterologous Peptide Epitope in the HI Loop of the Fiber Knob. <i>Journal of Virology</i> 72 , 1844-1852 (1998). <u>https://doi.org/DOI</u> : <u>https://doi.org/10.1128/jvi.72.3.1844-1852.1998</u>
566 567 568	29	Tellier, J. & Nutt, L. S. Standing out from the crowd: How to identify plasma cells. <i>European Journal of Immunology</i> 47 , 1276-1279 (2017). <u>https://doi.org/https://doi.org/10.1002/eji.201747168</u>
569 570	30	Chauhan, K. A. Human CD4+ T-Cells: A Role for Low-Affinity Fc Receptors. <i>Frontiers in Immunology</i> 7 (2016). <u>https://doi.org/doi</u> : 10.3389/fimmu.2016.00215

571 572 573	31	Kalyuzhniy, O. <i>et al.</i> Adenovirus serotype 5 hexon is critical for virus infection of hepatocytes in vivo. <i>Proceedings of the National Academy of Sciences</i> 105 , 5483-5488 (2008). <u>https://doi.org/DOI</u> : 10.1073/pnas.0711757105
574 575	32	Waddington, N. S. <i>et al.</i> Adenovirus Serotype 5 Hexon Mediates Liver Gene Transfer. <i>Cell</i> 132 , 397-409 (2008). <u>https://doi.org/DOI</u> : 10.1016/j.cell.2008.01.016
576 577	33	Li, C. <i>et al.</i> In vivo HSC prime editing rescues Sickle Cell Disease in a mouse model. <i>Blood</i> (2023). <u>https://doi.org/10.1182/blood.2022018252</u>
578 579 580	34	Li, C. <i>et al.</i> Safe and efficient in vivo hematopoietic stem cell transduction in nonhuman primates using HDAd5/35++ vectors. <i>Molecular Therapy - Methods & Clinical Development</i> 24 , 127-141 (2022). <u>https://doi.org/DOI:https://doi.org/10.1016/j.omtm.2021.12.003</u>
581 582 583	35	Wang, H. <i>et al.</i> Curative in vivo hematopoietic stem cell gene therapy of murine thalassemia using large regulatory elements. <i>JCI Insight</i> 5 (2020). <u>https://doi.org/doi</u> : 10.1172/jci.insight.139538
584 585 586	36	Arnberg, N. Adenovirus receptors: implications for targeting of viral vectors. <i>Trends in Pharmacological Sciences</i> 33 , 442-448 (2012). https://doi.org/https://doi.org/10.1016/j.tips.2012.04.005
587 588 589 590	37	Erp, V. A. E., Kaliberova, N. L., Kaliberov, A. S. & Curiel, T. D. Retargeted oncolytic adenovirus displaying a single variable domain of camelid heavy-chain-only antibody in a fiber protein. <i>Molecular Therapy - Oncolytics</i> 2 , 15001 (2015). https://doi.org/DOI:https://doi.org/10.1038/mto.2015.1
591 592 593	38	Sharma, K. P. <i>et al.</i> Development of an adenovirus vector vaccine platform for targeting dendritic cells. <i>Cancer Gene Therapy</i> 25 , 27-38 (2018). <u>https://doi.org/https://doi.org/10.1038/s41417-017-0002-1</u>
594 595 596	39	Lee, M. <i>et al.</i> Advanced genetic engineering to achieve in vivo targeting of adenovirus utilizing camelid single domain antibody. <i>Journal of Controlled Release</i> 334 , 106-113 (2021).
597		https://doi.org/https://doi.org/10.1016/j.jconrel.2021.04.009
598 599 600 601	40	https://doi.org/https://doi.org/10.1016/j.jconrel.2021.04.009 Dmitriev, I., Kashentseva, E., Rogers, E. B., Krasnykh, V. & Curiel, T. D. Ectodomain of Coxsackievirus and Adenovirus Receptor Genetically Fused to Epidermal Growth Factor Mediates Adenovirus Targeting to Epidermal Growth Factor Receptor-Positive Cells. <i>Journal of</i> <i>Virology</i> 74 , 6875-6884 (2000). https://doi.org/DOI: https://doi.org/10.1128/jvi.74.15.6875- 6884.2000
598 599 600 601 602 603 604	40 41	https://doi.org/https://doi.org/10.1016/j.jconrel.2021.04.009 Dmitriev, I., Kashentseva, E., Rogers, E. B., Krasnykh, V. & Curiel, T. D. Ectodomain of Coxsackievirus and Adenovirus Receptor Genetically Fused to Epidermal Growth Factor Mediates Adenovirus Targeting to Epidermal Growth Factor Receptor-Positive Cells. <i>Journal of</i> <i>Virology</i> 74, 6875-6884 (2000). <u>https://doi.org/DOI</u> : <u>https://doi.org/10.1128/jvi.74.15.6875-6884.2000</u> 624. Adenovirus Targeting to c-erbB-2 Oncoprotein by Single-Chain Antibody Using Recombinant Adapter Protein or Genetic Modification of Viral Capsid. <i>Molecular Therapy</i> 5, S203-S204 (2002). <u>https://doi.org/DOI:https://doi.org/10.1016/S1525-0016(16)43454-4</u>
598 599 600 601 602 603 604 605 606 607	40 41 42	 https://doi.org/https://doi.org/10.1016/j.jconrel.2021.04.009 Dmitriev, I., Kashentseva, E., Rogers, E. B., Krasnykh, V. & Curiel, T. D. Ectodomain of Coxsackievirus and Adenovirus Receptor Genetically Fused to Epidermal Growth Factor Mediates Adenovirus Targeting to Epidermal Growth Factor Receptor-Positive Cells. <i>Journal of</i> <i>Virology</i> 74, 6875-6884 (2000). https://doi.org/DOI: https://doi.org/10.1128/jvi.74.15.6875- 6884.2000 624. Adenovirus Targeting to c-erbB-2 Oncoprotein by Single-Chain Antibody Using Recombinant Adapter Protein or Genetic Modification of Viral Capsid. <i>Molecular Therapy</i> 5, S203-S204 (2002). https://doi.org/DOI:https://doi.org/10.1016/S1525-0016(16)43454-4 Dreier, B. <i>et al.</i> Development of a generic adenovirus delivery system based on structure-guided design of bispecific trimeric DARPin adapters. <i>Proceedings of the National Academy of Sciences</i> 110, E869-E877 (2013). https://doi.org/DOI: 10.1073/pnas.1213653110
598 599 600 601 602 603 604 605 606 607 608 609 610	40 41 42 43	https://doi.org/https://doi.org/10.1016/j.jconrel.2021.04.009 Dmitriev, I., Kashentseva, E., Rogers, E. B., Krasnykh, V. & Curiel, T. D. Ectodomain of Coxsackievirus and Adenovirus Receptor Genetically Fused to Epidermal Growth Factor Mediates Adenovirus Targeting to Epidermal Growth Factor Receptor-Positive Cells. <i>Journal of</i> <i>Virology</i> 74, 6875-6884 (2000). https://doi.org/DOI: https://doi.org/10.1128/jvi.74.15.6875- 6884.2000 624. Adenovirus Targeting to c-erbB-2 Oncoprotein by Single-Chain Antibody Using Recombinant Adapter Protein or Genetic Modification of Viral Capsid. <i>Molecular Therapy</i> 5, S203-S204 (2002). https://doi.org/DOI:https://doi.org/10.1016/S1525-0016(16)43454-4 Dreier, B. <i>et al.</i> Development of a generic adenovirus delivery system based on structure-guided design of bispecific trimeric DARPin adapters. <i>Proceedings of the National Academy of Sciences</i> 110, E869-E877 (2013). https://doi.org/DOI: 10.1073/pnas.1213653110 Kadkhodazadeh, M. <i>et al.</i> Fiber manipulation and post-assembly nanobody conjugation for adenoviral vector retargeting through SpyTag-SpyCatcher protein ligation. <i>Frontiers in Molecular</i> <i>Biosciences</i> 9 (2022). https://doi.org/https://doi.org/10.3389/fmolb.2022.1039324

611 612 613	44	Kasaraneni, N., Chamoun-Emanuelli, M. A., Wright, G. & Chen, Z. Retargeting Lentiviruses via SpyCatcher-SpyTag Chemistry for Gene Delivery into Specific Cell Types. <i>mBio</i> 8 (2017). <u>https://doi.org/doi</u> : 10.1128/mBio.01860-17
614 615 616	45	Kaliberov, A. S. <i>et al.</i> Retargeting of gene expression using endothelium specific hexon modified adenoviral vector. <i>Virology</i> 447 , 312-325 (2013). <u>https://doi.org/doi</u> : 10.1016/j.virol.2013.09.020
617 618 619	46	Alba, R. <i>et al.</i> Identification of coagulation factor (F)X binding sites on the adenovirus serotype 5 hexon: effect of mutagenesis on FX interactions and gene transfer. <i>Blood</i> 114 , 965-971 (2009). https://doi.org/https://doi.org/10.1182/blood-2009-03-208835
620 621 622	47	Atasheva, S. <i>et al.</i> Systemic cancer therapy with engineered adenovirus that evades innate immunity. <i>Science Translational Medicine</i> 12 , eabc6659 (2020). <u>https://doi.org/DOI</u> : 10.1126/scitranslmed.abc6659
623 624 625	48	Lorincz, R. <i>et al.</i> In vivo editing of the pan-endothelium by immunity evading simian adenoviral vector. <i>Biomedicine & Pharmacotherapy</i> 158 , 114189 (2023). <u>https://doi.org/DOI</u> : 10.1016/j.biopha.2022.114189
626 627 628 629	49	Jaskiewicz, J. J., Tremblay, M. J., Tzipori, S. & Shoemaker, B. C. Identification and characterization of a new 34 kDa MORN motif-containing sporozoite surface-exposed protein, Cp-P34, unique to Cryptosporidium. <i>International Journal for Parasitology</i> 51 , 761-775 (2021). <u>https://doi.org/doi</u> : 10.1016/j.ijpara.2021.01.003
630 631 632	50	Mitchell, G. K. <i>et al.</i> High-volume hybridoma sequencing on the NeuroMabSeq platform enables efficient generation of recombinant monoclonal antibodies and scFvs for neuroscience research. <i>Scientific Reports</i> 13 (2023). <u>https://doi.org/DOI</u> : 10.1038/s41598-023-43233-4
633 634	51	Hong, S. J. & Engler, A. J. Domains required for assembly of adenovirus type 2 fiber trimers. Journal of Virology 70 , 7071-7078 (1996). <u>https://doi.org/doi</u> : 10.1128/jvi.70.10.7071-7078.1996
635 636 637	52	Luo, B. <i>et al.</i> Engineering of α-PD-1 antibody-expressing long-lived plasma cells by CRISPR/Cas9- mediated targeted gene integration. <i>Cell Death & Disease</i> 11 (2020). <u>https://doi.org/DOI</u> : 10.1038/s41419-020-03187-1
638 639 640	53	Hung, L. K. <i>et al.</i> Engineering Protein-Secreting Plasma Cells by Homology-Directed Repair in Primary Human B Cells. <i>Molecular Therapy</i> 26 , 456-467 (2018). <u>https://doi.org/DOI</u> : 10.1016/j.ymthe.2017.11.012
641 642 643	54	Cheng, YH. R. <i>et al.</i> Ex vivo engineered human plasma cells exhibit robust protein secretion and long-term engraftment in vivo. <i>Nature Communications</i> 13 (2022). <u>https://doi.org/https://doi.org/10.1038/s41467-022-33787-8</u>
644 645	55	Jumper, J. <i>et al.</i> Highly accurate protein structure prediction with AlphaFold. <i>Nature</i> 596 , 583-589 (2021). <u>https://doi.org/https://doi.org/10.1038/s41586-021-03819-2</u>
646 647	56	Meng, C. E. <i>et al.</i> UCSF ChimeraX: Tools for structure building and analysis. <i>Protein Science</i> 32 (2023). <u>https://doi.org/https://doi.org/10.1002/pro.4792</u>
648		

649 Figure Legends:

650	Figure 1: Development and characterization of Ad-Ab targeting. a. Schematic overview of system design.
651	DogTag is genetically inserted into the Ad fiber knob (Ad5FDgT), while DogCatcher is fused to antibody
652	species. Mixing of these reagents results in permanent linkage of the virus and antibody at the fiber
653	knob locale. Fiber model generated with AlphaFold2 and visualized with UCSF ChimeraX ^{55,56} . b. Ad5FDgT
654	genome overview. Ad5FDgT is based on an E1/E3 deleted Ad5 with the CMV promoter driving eGFP
655	expression from the E1 region. DogTag is inserted with minimal flex linkers at the HI loop of the fiber
656	knob domain. c. Antibody-DogCatcher fusion designs. Antibody domains are separated from DogCatcher
657	by a flexible (G4S)₃ linker d. SDS-PAGE analysis of control Ad5 and Ad5FDgT. In all cases the molar
658	amount of antibody and virus refers to DogCatcher and DogTag, respectively. e . Gel shift analysis of
659	Ad5FDgT binding to DogCatcher and antibody-DogCatcher fusions. Left, binding to bacterially produced
660	DogCatcher and an sdAb-DogCatcher fusion. Middle and right, binding to representative scFv-
661	DogCatcher fusions. f. Flow cytometry analysis of representative scFvs in murine splenocytes. 1D3DgC
662	and 18B12DgC stains were gated on B220+ populations, while 2.43DgC staining was gated on the CD8+
663	population. 1D3DgC specificity was assessed by blocking with excess full length 1D3 antibody.
664	Figure 2: In vitro characterization of Ad-Ab targeting. a. Conceptual workflow. Primary lymphocytes are
665	magnetically isolated from mouse splenocytes or human PBMCs then cultured with activating agents.
666	On the day of infection Ad5FDgT is conjugated with the appropriate antibody at the indicated
667	DogCatcher:DogTag molar ratios, then used to infect cells. b. Ad-Ab infectivity enhancement in primary
668	murine B cells (top right), murine T cells (bottom left), and human B cells (bottom right). For each
669	antibody, the group with the maximum mean infectivity was compared to the no antibody group using a
670	standard unpaired two-tailed t-test. Welch's correction was used in cases where the F-test revealed
671	significant differences in variances. DgC = DogCatcher , n=9 replicates from 3 experiments in mouse B
672	cells, n=6 from 2 experiments in mouse T cells. YTS169DgC = α -mCD8 α scFv, n=4-5 from 2 experiments

673	in mouse B cells, n=6 from 2 experiments in mouse T cells, n=5 from 3 experiments in human B cells.
674	2.43 = α -mCD8 α scFv, n=4-5 from 2 experiments in mouse B cells, n=6 from 2 experiments in mouse T
675	cells. F8DgC = α -mCD40 sdAb, n=9 from 3 experiments in mouse B cells, n=6 from 2 experiments in
676	mouse T cells. 1D3DgC = α -mCD19 scFv, n=4 from 2 experiments in mouse B cells, n=6 from 2
677	experiments in mouse T cells. 18B12DgC = α -mCD20 scFv, n=4-5 from 2 experiments in mouse B cells,
678	n=6 from 2 experiments in mouse T cells. FMC63DgC = α -hCD19 scFv, n=5 from 3 experiments in human
679	B cells. Rtxv1DgC (Rituximab) = α-mCD20 scFv, n=4 from 2 experiments in human B cells. HA22DgC = α-
680	hCD22 scFv, n=5 from 3 experiments in human B cells. Created with BioRender.com.
681	Figure 3: In vivo characterization of Ad-Ab targeting. a. Experiment design. C57BL/6J mice were injected
682	RO with 5x10 ¹⁰ vp/mouse of either Ad5FDgT or Ad5FDgT conjugated with 1D3DgC. Major organs were
683	harvested 72h later for eGFP expression analyses. b. Flow cytometry results. Left, eGFP expression in B
684	and T cells. Right, eGFP expression in B cell subsets. GC = germinal center, MEM = memory, MZ =
685	marginal zone, FO = follicular, PB = plasmablast. In all cases non-conjugated and conjugated groups were
686	compared in each population using a standard unpaired two-tailed t-test. Welch's correction was used
687	in cases where the F-test revealed significant differences in variances. eGFP expression in mice injected
688	with PBS was used to normalize the data in all cases. c. Quantitative eGFP analysis in major tissues.
689	Tissues were homogenized in lysis buffer and eGFP expression was analyzed via fluorimetry. eGFP was
690	normalized to total tissue protein content as assessed by BCA. In all cases ordinary one-way ANOVA with
691	Tukey's correction for multiple comparisons was used to compare groups where n=5 male mice. In the
692	liver, lung and kidney values were log transformed to normalize variances. Created with BioRender.com.
693	Figure 4: Development of purified Ad-Ab complexes. a. Purification workflow. Virus particles are purified
694	from cell lysates via cesium chloride (CsCl) ultracentrifugation, then briefly dialyzed against 1X PBS to
695	remove excess salts. Conjugation is then carried out, followed by a second CsCl purification, dialysis and
696	final storage. b. Viral band images in CsCl gradients after second ultracentrifugation. c. SDS-PAGE

697	analysis of purified Ab-Ab complexes. d. Western blot against the HAdV-C5 fiber tail of purified Ad-Ab
698	complexes. e. In vivo analysis of purified Ad-Ab complexes. C57BL/6J mice aged 6-9 weeks were injected
699	on three separate occasions with $5x10^{10}$ of the indicated vectors. In the first experiment n=2 female
700	mice per group were used. In the second experiment n=2 male mice per group were used, and a single
701	Ad5-FDgT-YTS169 sample was removed from analysis due to very low viability found during flow
702	analysis. In the fourth experiment n=4 mice per group were used split equally between male and female
703	mice. Three days later splenocytes were assessed for eGFP expression using flow cytometry. Livers were
704	assessed for tissue eGFP expression as well. Data from all experiments were combined and analyzed
705	using ordinary one-way ANOVA with Tukey's correction for multiple comparisons. Created with
706	BioRender.com.
707	
700	
708	
709	
710	
711	
712	
713	
714	
715	
/15	
716	
717	

718 Figures:





725





Prepare CsCl gradient Dialysis Conjugation CsCl gradient cell lysate centrifugation centrifugation b. c. Reagent pmol 1D3DgC -25 25 --25 25 YTS169DgC --Ad5FDgT+ Ad5FDgT+ Ad5FDgT+ 2.5 2.5 121 2.5 -Ad5FDgT YTS169DgC PBS 1D3DgC 180 130 FDgT-scFv 100 70 🗲 FDgT 55 ✓ scFv-DgC 40 35 25 15 10 kDa



Aliquot and

Dialysis









