

## Supplementary information

### Reprogramming Synthetic Cells for Targeted Cancer Therapy

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# B.L. and Y.Y contributed equally to this work.

**Table S1.** Bacterial strains and plasmids used in this study.

Strains	Genotype or description	Source
<i>Escherichia coli</i> DH5 $\alpha$	<i>fhuA2 lac(del)U169 phoA glnV44 <math>\Phi</math>80' lacZ(del)M15 gyrA96 recA1 relA1 endA1 thi-1 hsdR17</i>	Lab collection
<i>E. coli</i> BL21(DE3)	<i>fhuA2 [lon] ompT gal (<math>\lambda</math> DE3) [dcm] <math>\Delta</math>hsdS</i>	Lab collection
<i>E. coli</i> K12 MDS42	<i>fhuACDB(del) endA(del)</i> and deletion of all IS and cryptic prophage <sup>1</sup>	Lab collection
DH5 $\alpha$ pNVC17	BL21(DE3): pNVC17 (for cloning)	This study
BL21(DE3) pNVC17_2	BL21(DE3): pNVC17_2 (low expression profile)	This study
BL21(DE3) pNVC43_2	BL21(DE3): pNVC43_2 (low expression profile)	This study
BL21(DE3) pNVCYT_2	BL21(DE3): pNVCYT_2 (low expression profile)	This study
BL21(DE3) pNVC17_3	BL21(DE3): pNVC17_3 (high expression profile)	This study
BL21(DE3) pNVC43_3	BL21(DE3): pNVC43_3 (high expression profile)	This study
BL21(DE3) pNVCYT_3	BL21(DE3): pNVCYT_3 (high expression profile)	This study
BL21(DE3) control	BL21(DE3): proD_sfGFP (high expression profile control with sfGFP)	This study
BL21(DE3) pNVC17_sfGFP	BL21(DE3): pNVC17_sfGFP (high expression profile with sfGFP downstream)	This study
BL21(DE3) pNVC43_sfGFP	BL21(DE3): pNVC43_sfGFP (high expression profile with sfGFP downstream)	This study
BL21(DE3) pNVCYT_sfGFP	BL21(DE3): pNVCYT_sfGFP (high expression profile with sfGFP downstream)	This study
BL21(DE3) pNVC17_sfGFP_ICeul	BL21(DE3): pNVC17_sfGFP + pRH12x (high expression profile with sfGFP downstream; ICeul endonuclease expression)	This study
BL21(DE3) pNVC43_sfGFP_ICeul	BL21(DE3): pNVC43_sfGFP + pRH12x (high expression profile with sfGFP downstream; ICeul endonuclease expression)	This study
BL21(DE3) pNVC43_SalA_sfGFP_ICeul	BL21(DE3): pNVC43_salA_sfGFP + pRH12x (high expression profile with sfGFP downstream; ICeul endonuclease expression)	This study
BL21(DE3) pNVCYT_sfGFP_ICeul	BL21(DE3): pNVCYT_sfGFP + pRH12x (high expression profile with sfGFP downstream; ICeul endonuclease expression)	This study
BL21(DE3) pSIJ8	BL21(DE3): pSIJ8 (for recombination engineering)	This study
BL21(DE3) $\Delta$ minD	BL21(DE3) $\Delta$ minD::Kan <sup>R</sup> (for minicells generation)	This study
BL21(DE3) $\Delta$ minD pNVC17_sfGFP	BL21(DE3): pNVC17_sfGFP, $\Delta$ minD::Kan <sup>R</sup> (for minicells with high expression profile with sfGFP downstream)	This study
BL21(DE3) $\Delta$ minD pNVC43_sfGFP	BL21(DE3): pNVC43_sfGFP, $\Delta$ minD::Kan <sup>R</sup> (for minicells with high expression profile with sfGFP downstream)	This study

BL21(DE3) $\Delta minD$ pNVCYT_sfGFP	BL21(DE3): pNVCYT_sfGFP, $\Delta minD::Kan^R$ (for minicells with high expression profile with sfGFP downstream)	This study
<b>Plasmids</b>		
pNVgfp	Plasmid with anti-GFP nanobody attached to $\beta$ -intimin, constitutively expressed by the promoter BBa_J23106 and a low strength RBS (12K TIR). $Cm^R$ .	A gift from Dr Karen Polizzi <sup>2</sup>
pNVC17	Anti-GFP nanobody of pNVgfp replaced by anti-CEA nanobody. $Cm^R$ .	This study
pNVC43	Anti-GFP nanobody of pNVgfp replaced by another variant of anti-CEA nanobody. $Cm^R$ .	This study
pNVC17_2	BBa_J23106 of pNVC17 replaced by BBa_J23105. $Cm^R$ .	This study
pNVC43_2	BBa_J23106 of pNVC43 replaced by BBa_J23105. $Cm^R$ .	This study
pNVCYT_2	Anti-CEA nanobody of pNVC17_2 replaced by anti-spike protein nanobody. $Cm^R$ .	This study
pNVC17_3	BBa_J23106 and RBS of pNVC17 replaced by BBa_K1741014 and BBa_K1758100, respectively.	This study
pNVC43_3	Anti-CEA nanobody of pNVC17_3 replaced by another anti-CEA nanobody variant. $Cm^R$ .	This study
pNVCYT_3	Anti-CEA nanobody of pNVC17_3 replaced by anti-spike protein nanobody. $Cm^R$ .	This study
pNVC17_sfGFP	BBa_K1758100 and <i>sfGFP</i> cloned downstream to the anti-CEA nanobody in pNVC17_3. $Cm^R$ .	This study
pNVC43_sfGFP	Anti-CEA nanobody of pNVC17_sfGFP replaced by another anti-CEA nanobody variant. $Cm^R$ .	This study
pNVC43_SalA_sfGFP	<i>salA</i> cloned upstream of <i>sfGFP</i> within pNVC43_sfGFP. $Cm^R$ .	This study
pNVCYT_sfGFP	Anti-CEA nanobody of pNVC17_sfGFP replaced by anti-spike protein nanobody. $Cm^R$ .	This study
prod_sfGFP	A lab plasmid with BBa_K1741014 and BBa_K1758100 upstream of a <i>sfGFP</i> . $Cm^R$ .	Lab collection
pJKR-L-TetR-ICeul	Plasmid with I-Ceul endonuclease controlled by TetR. $Carb^R$ .	<sup>3</sup>
pRH12x	Plasmid with ICeul endonuclease controlled by a crystal violet inducible promoter. $Kan^R$ .	This study
pSIJ8	Plasmid with lambda Red recombineering genes ( <i>exo</i> , <i>bet</i> and <i>gam</i> ) and a flippase (FLP) recombinase. $Carb^R$ .	<sup>4</sup>

**Table S2.** Primers and gene blocks used in this study.

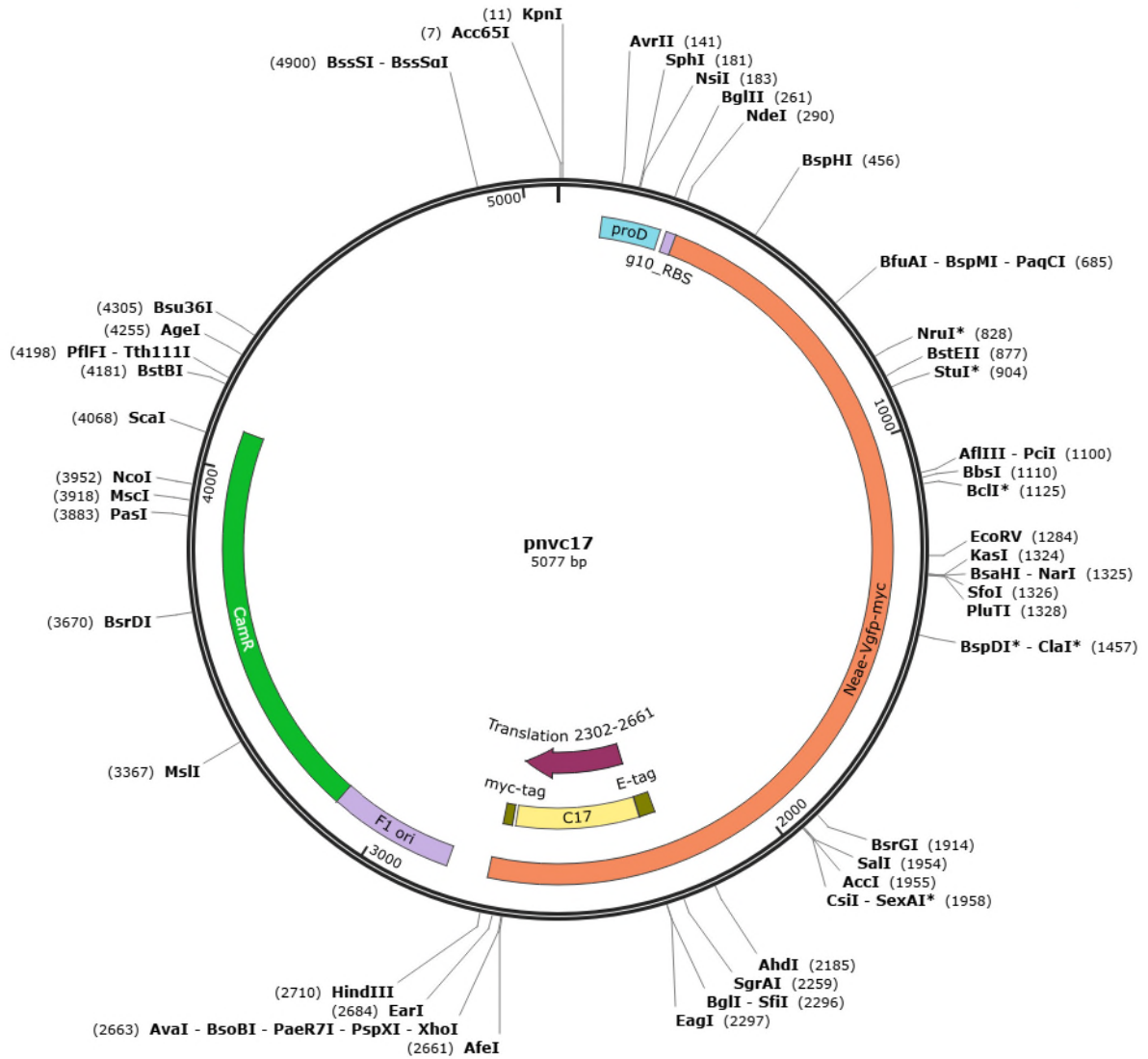
Primers	Sequence (5' → 3')	Part/Purpose	Plasmid
VHH-for	GAACCGGCCAGCCGGCCGAAGTTCAAC TGGTTGAAAG	Anti-CEA nanobody	pNVC17, pNVC43, pNVC17_2, pNVC43_2
VHH-rev	GATGAGTTTTTGTTCGGCGGCCGCGCTG CTAACTGTAACCTG		

pNV-for	GCTCGAGCCGAACAAAACTCATCTCAGA AGAGGATGCAGCTGC	pNVgfp backbone	pNVC17, pNVC43, pNVC17_2, pNVC43_2, pNVCYT_2
pNV-rev	GGCCGGCTGGGCCGGTTCCAGC		
pNV-3- for	ATGATTACTCATGGTTGTTATACCCGG	pNVC17 backbone	pNVC17_3, pNVC43_3, pNVCYT_3
pNV-3- rev	GCACACGGTCACACTGCTTC		
pNV- sfGFP- for	GCATGGATGAGCTCTACAAAAGCTTGAC CTGTGAAGTG	pNVC17_3 backbone	pNVC17_sfGF P, pNVC43_sfGF P, pNVCYT_sfG FP
pNV- sfGFP- rev	ATGGTACCTTTCTCCTCTTTTTATGCAGCT GCATCCTC		
sfGFP- for	AAAGAGGAGAAAGGTACCATGAGCAAAG	sfGFP	pNVC17_sfGF P, pNVC43_sfGF P, pNVCYT_sfG FP
sfGFP- rev	TTTGTAGAGCTCATCCATGCCATGTG		
minD- HR-Kan- for	GTTGGCTGTGTTTTCTTCCGCGAGAGAA AGAAATCGAGTAATGCCATAACTTAGAAA AACTCATCGAGCATCAAATG	Kanamycin resistance gene cassette	-
minD- HR-Kan- rev	CGCTTTGACCGTTCAACCGTTAAATTGAT CCCTTTTTAACAAAGGAATTTCTATGAGCCA TATTCAACGGGAAAC		
HR- check-for	GACCTCAAGAATATCTTTACGCAACTG	Check for success recombinatio n	-
HR- check- rev	CTGTTTGTAATTGTCCTTTTAAACAGCG		
<b>Gene blocks</b>			
VHH- C17	GAAGTTCAACTGGTTGAAAGCGGTGGTG GTTTTGTTTCAGGCAGGCGAAAGTCTGACC CTGAGCTGTACCAGCAGCACCTGACCTT TACACCGTATCGTATGGCATGGTATCGTC AGGCACCGGGTAAACAGCGTGATCTGGT TGCAGATATTAGCAGCGGTGATGGTCGTA CCACCAATTATGCAGATTTTGCCAAAGGT CGTTTTACCATTAGCCGTGACAATATCAA AAACACCGTTTTTCTGCGTATGACCAATC TGAAACCGGAAGATACCGCAGTGTATTAT TGTAATACCTTTGTGAGCTTTGTGGGTATT GCACGTAGCTGGGGTCAGGGCACCCAGG TTACAGTTAGCAGC	Anti-CEA nanobody <sup>5</sup>	
VHH- C43	GAAGTTCAACTGGTTGAAAGCGGTGGTG GTCTGGTTCAGGCAGGCGGTAGTCTGAC CCTGAGCTGTACCAGCAGCACCTGACC TTTACACCGTATCGTATGGGTTGGTATCG TCAGACACCGGGTAAACAGCGTGATCTG GTTGCAGATATTAGCCCTGGTGATGGTAG CACCAAAAATATGCAGGTTTTGCACAGG	Anti-CEA nanobody <sup>6</sup>	

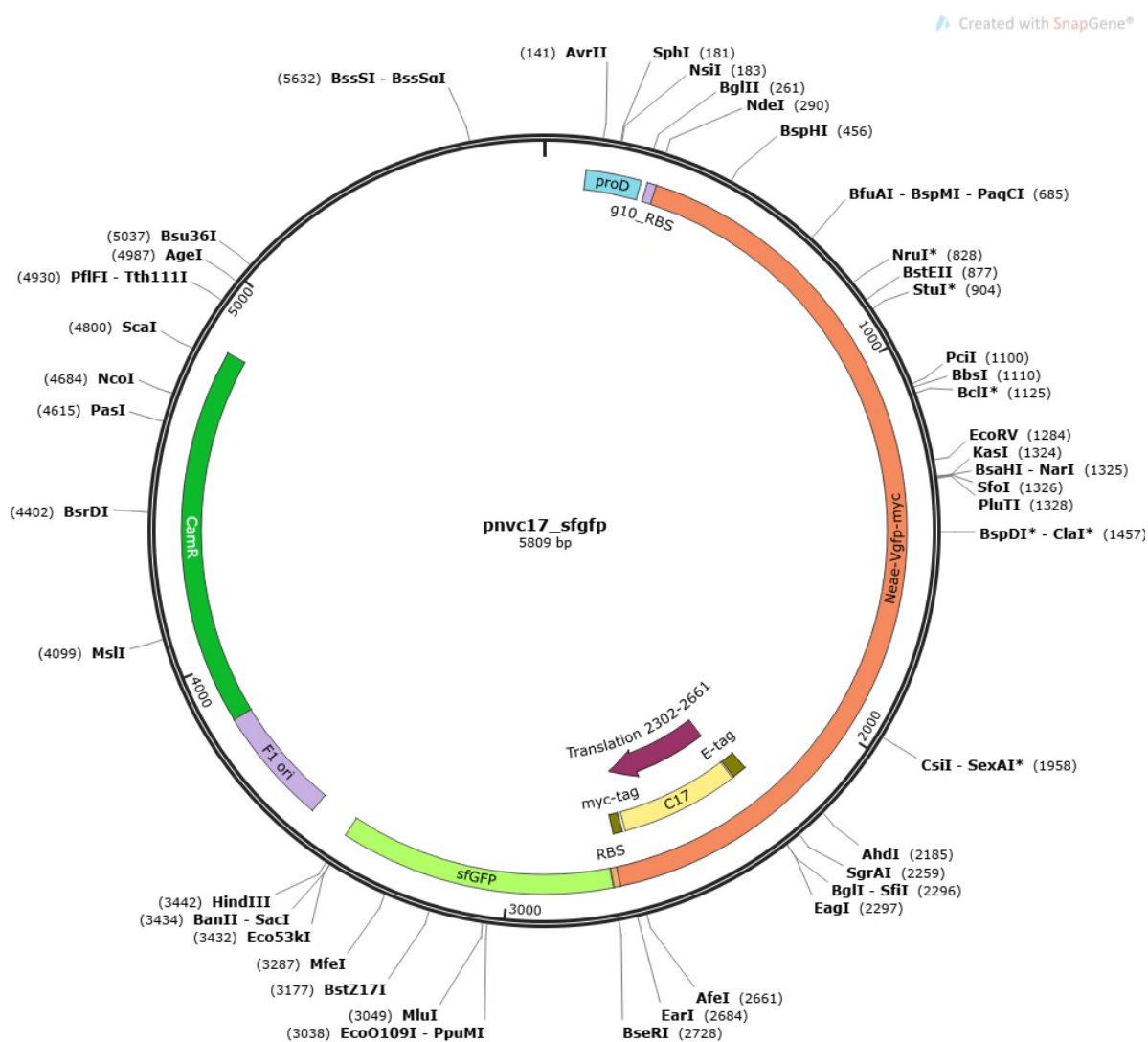
	GTCGTTTTACCATTAGCCGTGATAATATCA AAAACACCGTT TACCTGCAGATGAACGATCTGAAACCGGA AGATACCGCAGTGTATTATTGTAATACCTA TGTTGCCTTTGTTGGTCGTGCACGTACCT GGGGTCAGGGCACCCAGGTTACAGTTAG CAGC	
VHH- CYT	CAGGTGCAGCTCGTGGAGACGGGGGGA GGCTTGGTGCAGCCTGGGGGGTCTCTGA GACTCTCCTGTGCAGCCTCTGGATTCACC TTCAGTAGCGTCTACATGAACTGGGTCCG CCAGGCTCCAGGGAAGGGGCCCGAGTG GGTCTCGCGTATTAGTCCGAATAGTGGTA ATATTGGGTATACAGACTCCGTGAAGGGC CGATTCACCATCTCCAGAGACAACGCCAA GAACACACTGTATCTGCAAATGAATAACC TGAAACCTGAGGACACGGCCCTGTATTAC TGTGCGATTGGTTTGAATTTGAGTAGTAG CTCCGTTAGGGGCCAGGGGACCCAGGTC ACCGTCTCCTCA	Anti-Spike protein nanobody <sup>7</sup>
proD- g10	CACAGCTAACACCACGTCGTCCCTATCTG CTGCCCTAGGTCTATGAGTGGTTGCTGGA TAACTTTACGGGCATGCATAAGGCTCGTA TAATATATTCAGGGAGACCACAACGGTTT CCCTCTACAAATAATTTTGTTTAACTTTGA ATTCAAAGATCTTTAACTTTAAGAAGGAG ATATACAT	BBa_K1741014 and BBa_K1758100

a

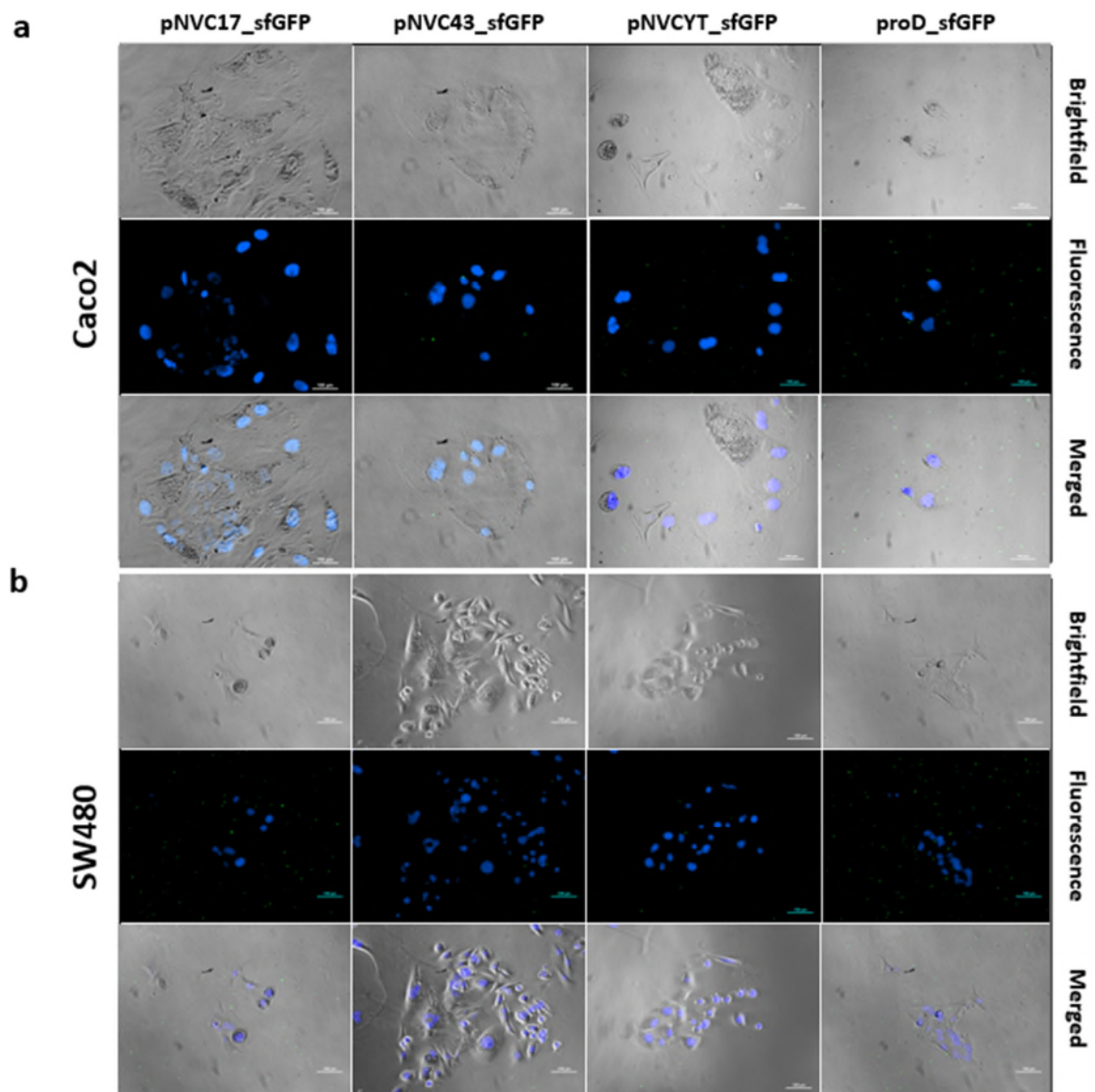
Created with SnapGene®



**b**

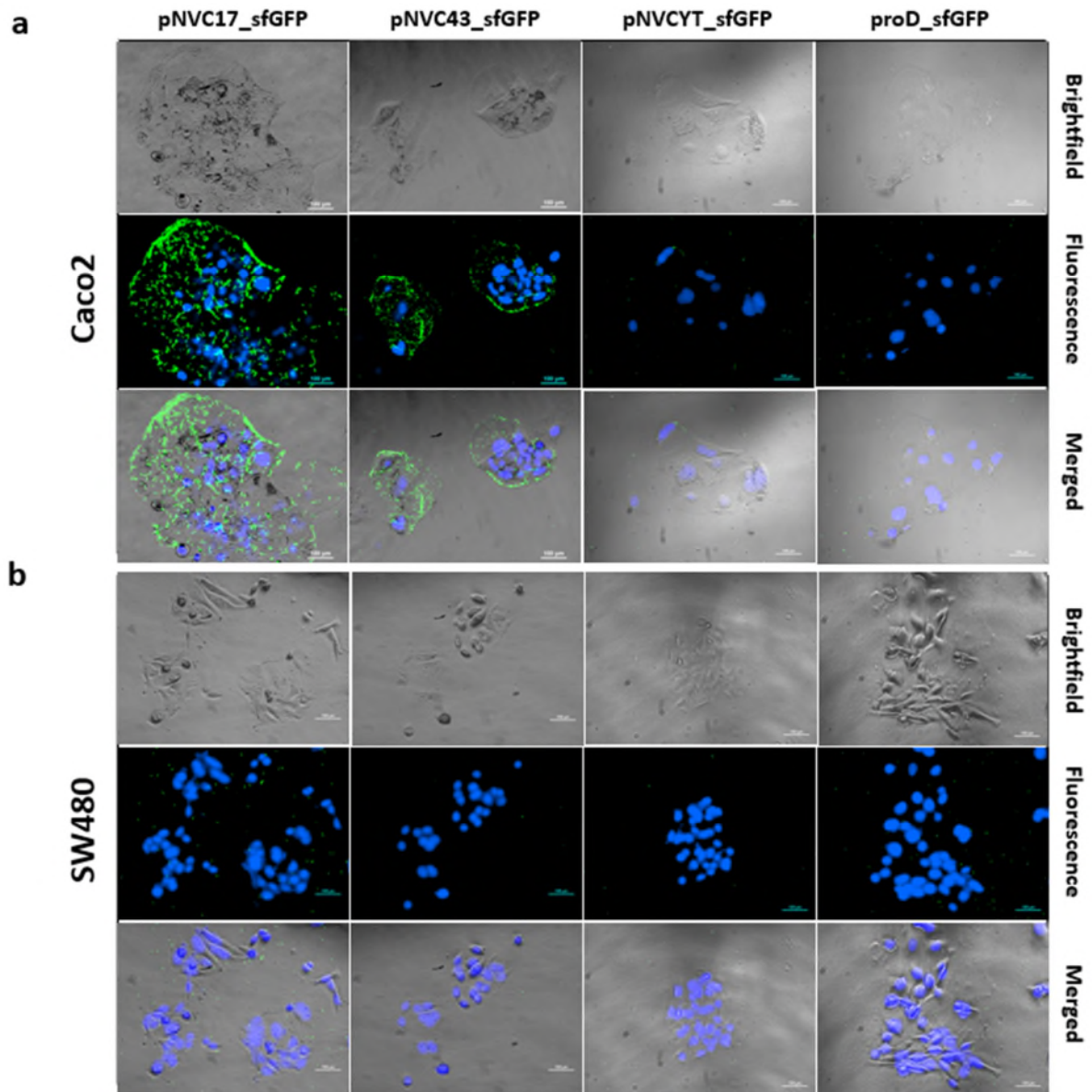


**Figure S1.** Plasmid map of (a) pNVC17 and (b) pNVC17\_sfGFP plasmids. The promoter (proD), ribosome binding site (g10\_RBS) and nanobody (C17) are interchangeable with other parts listed in Table S2.

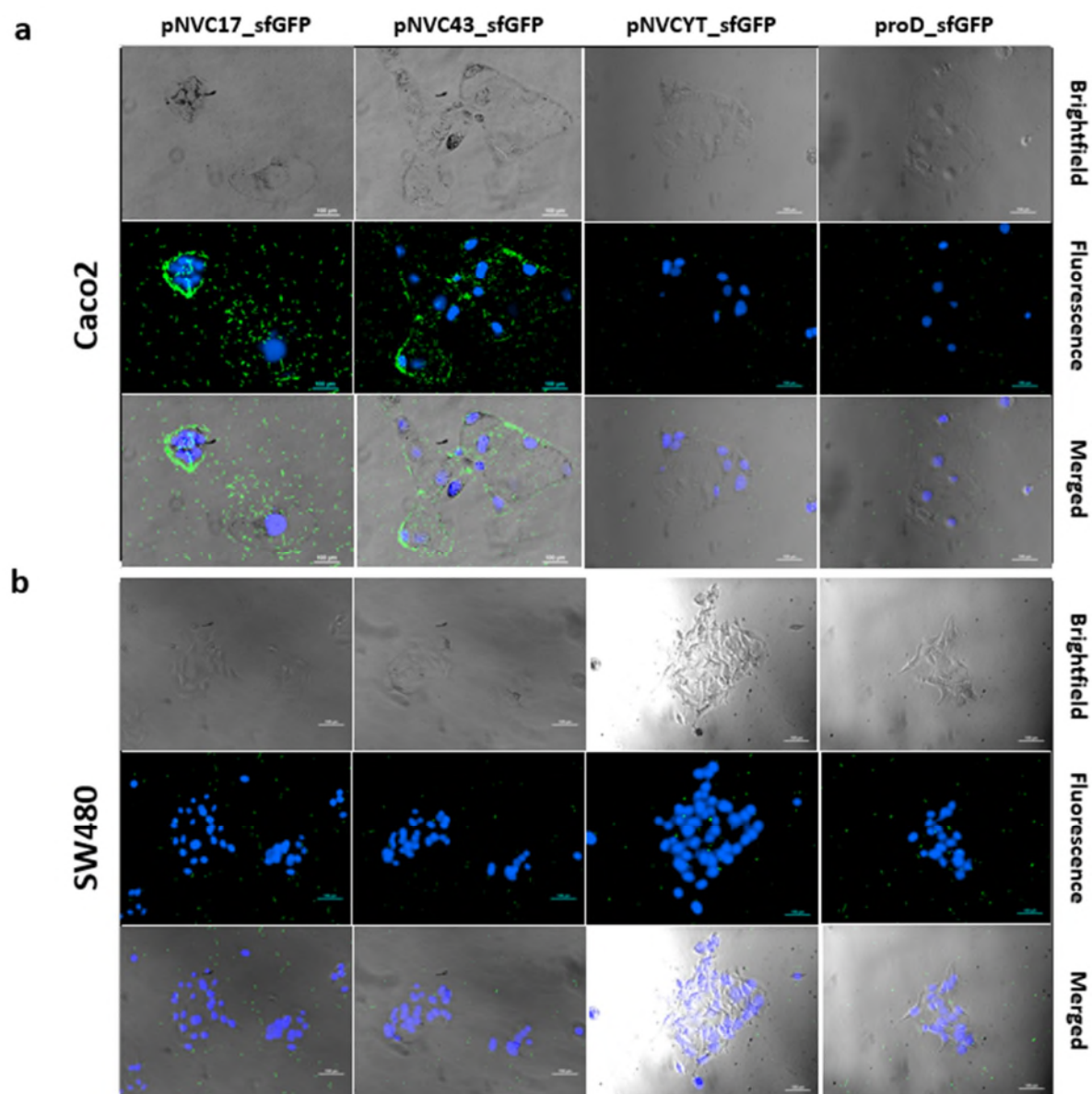


**Figure S2.** Time 0 hour (within ~5 minutes) incubation of engineered *E. coli* carrying different pNV plasmids with sfGFP (green) with two different colorectal cancer cell lines: **(a)** Caco2, a high CEA expressing cell line and **(b)** SW480, a low CEA expressing cell. Nuclei of cancer cells are stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.

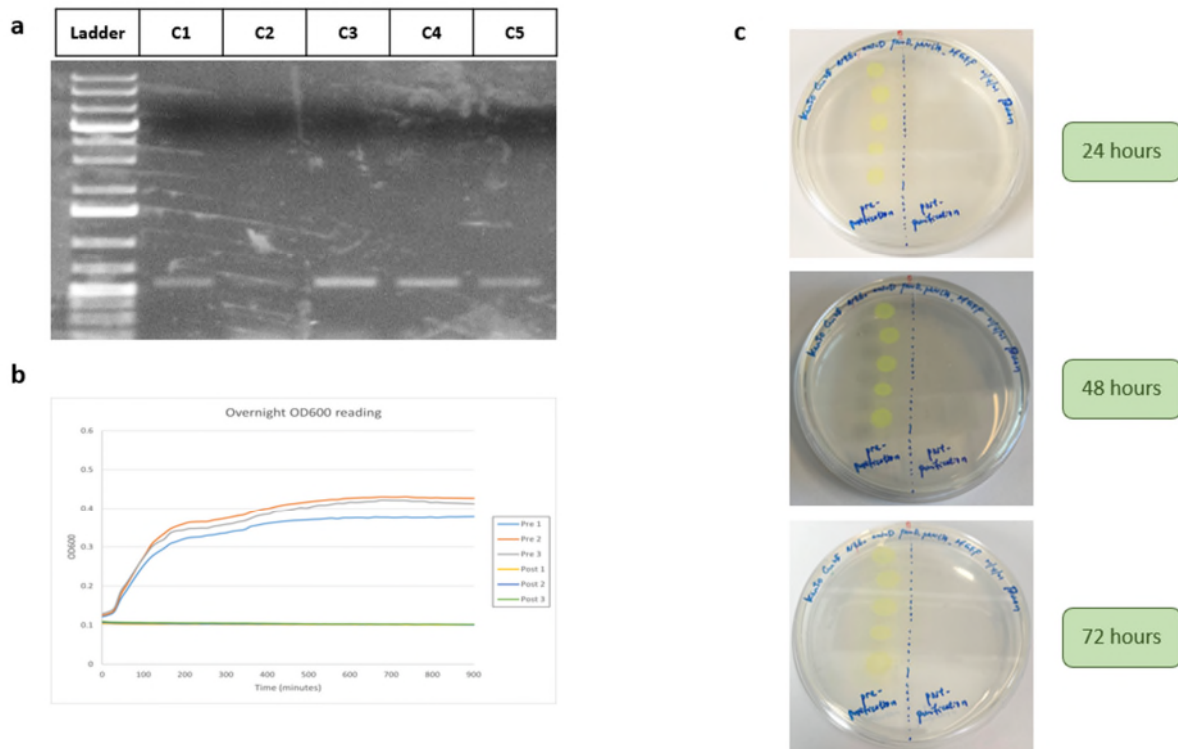




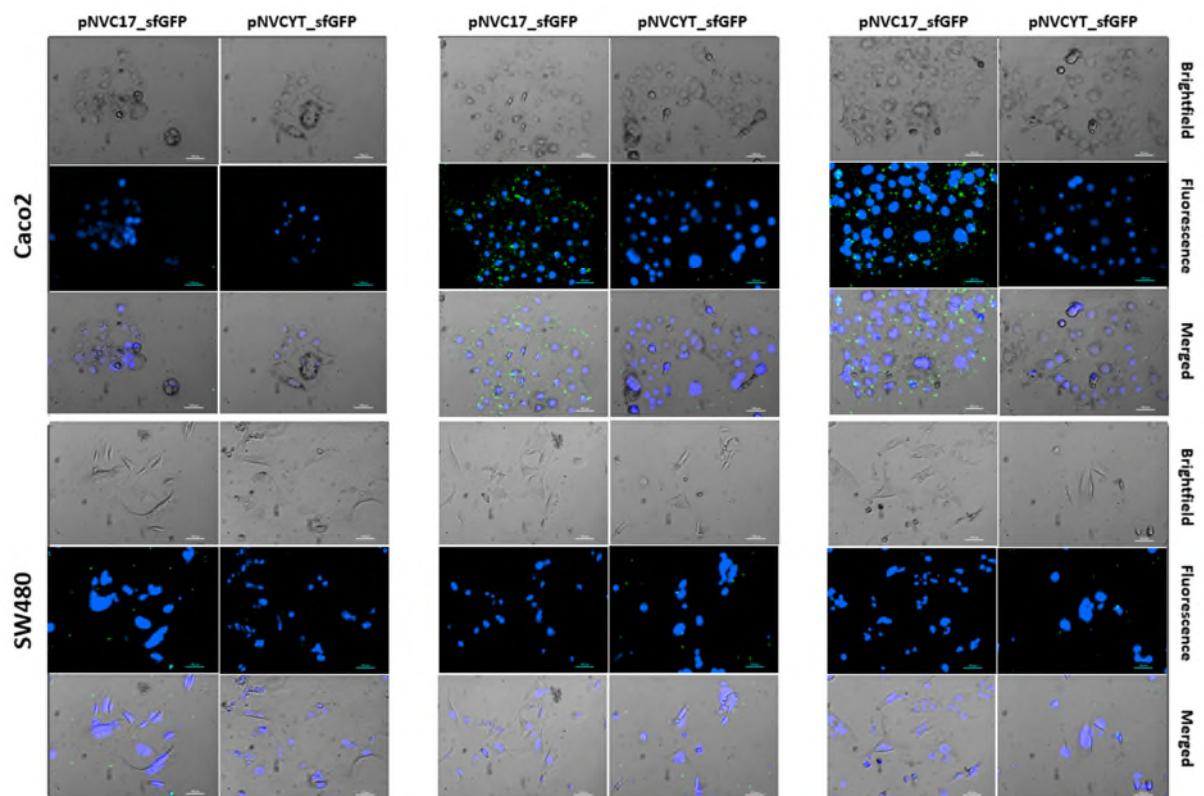
**Figure S3.** Time 4 hours incubation of engineered *E. coli* carrying different pNV plasmids with sfGFP (green) with two different colorectal cancer cell lines: **(a)** Caco2, a high CEA expressing cell line and **(b)** SW480, a low CEA expressing cell. Nuclei of cancer cells are stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.



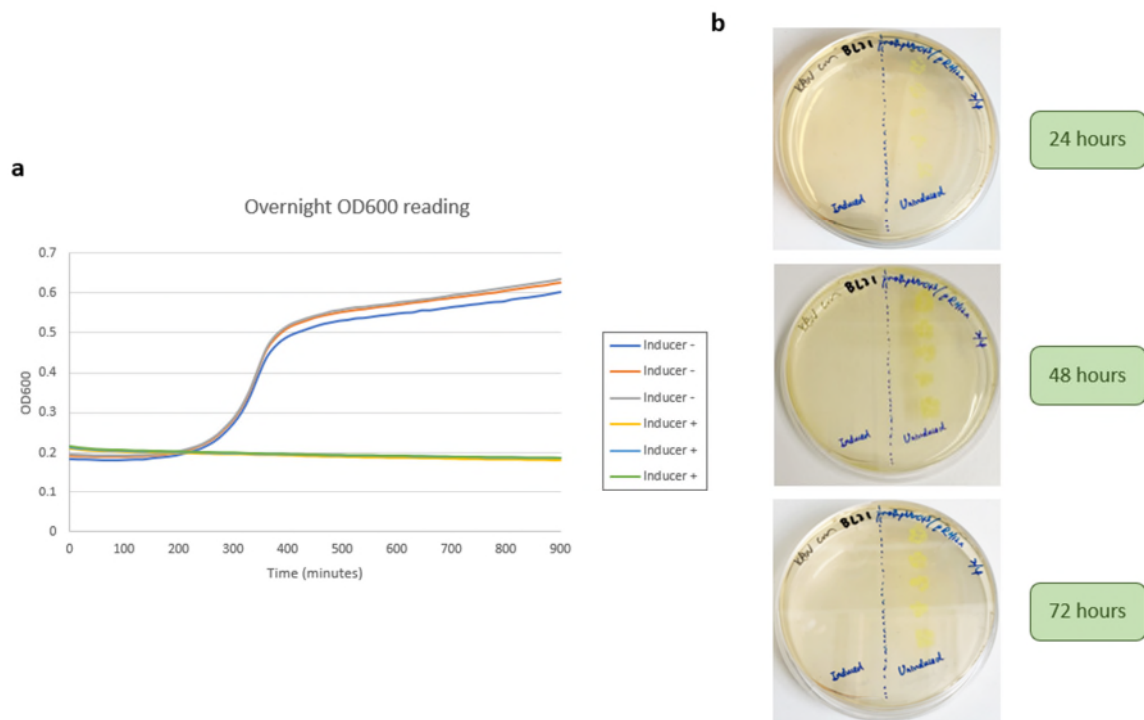
**Figure S4.** Time 8 hours incubation of engineered *E. coli* carrying different pNV plasmids with sfGFP (green) with two different colorectal cancer cell lines: **(a)** Caco2, a high CEA expressing cell line and **(b)** SW480, a low CEA expressing cell. Nuclei of cancer cells are stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.



**Figure S5. (a)** Gel electrophoresis of PCR products from five different colonies (C1-C5) with successful *minD* knock out. The primers used were *minD*-HR-Kan-for and *minD*-HR-Kan-rev (Table S2), which anneal to the inserted Kanamycin cassette and the *minE* gene within the bacterial genome, respectively. The length of PCR product is 563 bp. The ladder used is a 1kb plus DNA ladder from NEB. **(b)** Overnight OD<sub>600</sub> reading of pre-purification culture (pre 1, 2, 3) and post-purification culture (post 1, 2, 3) of minicells. Each line represents a single biological repeat. **(c)** Pre-purification and post-purification minicell culture plated on LB-Kan-Cm agar for 24 hours (top), 48 hours (mid) and 72 hours (bottom) at 37 °C.

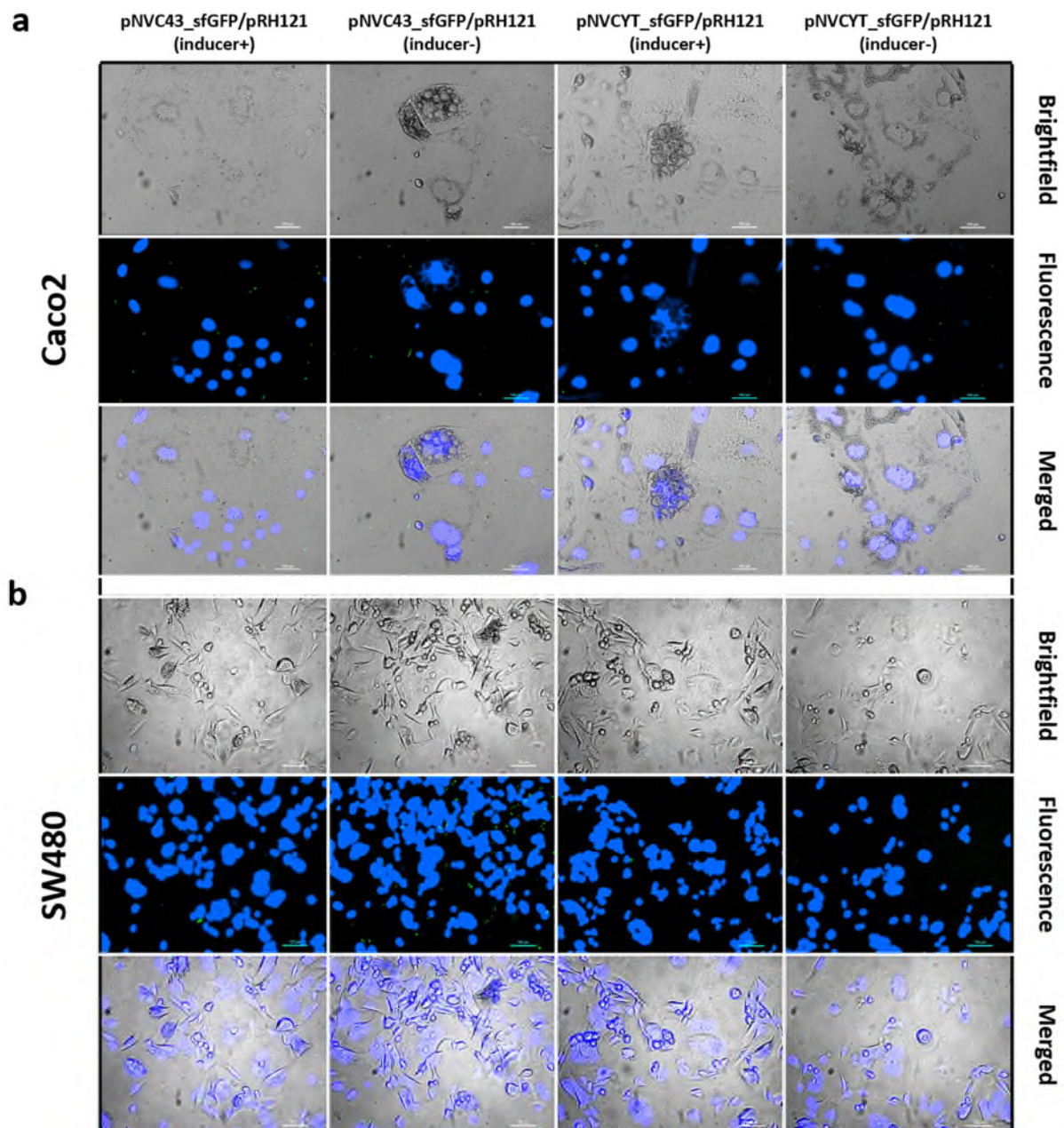


**Figure S6.** Time at 0 (left), 4 (mid) and 8 (right) hours incubation of engineered minicells carrying pNVC17\_sfGFP or pNVCYT\_sfGFP plasmids (green) with two different colorectal cancer cell lines: **(a)** Caco2, a high CEA expressing cell line and **(b)** SW480, a low CEA expressing cell. Nuclei of cancer cells are stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.

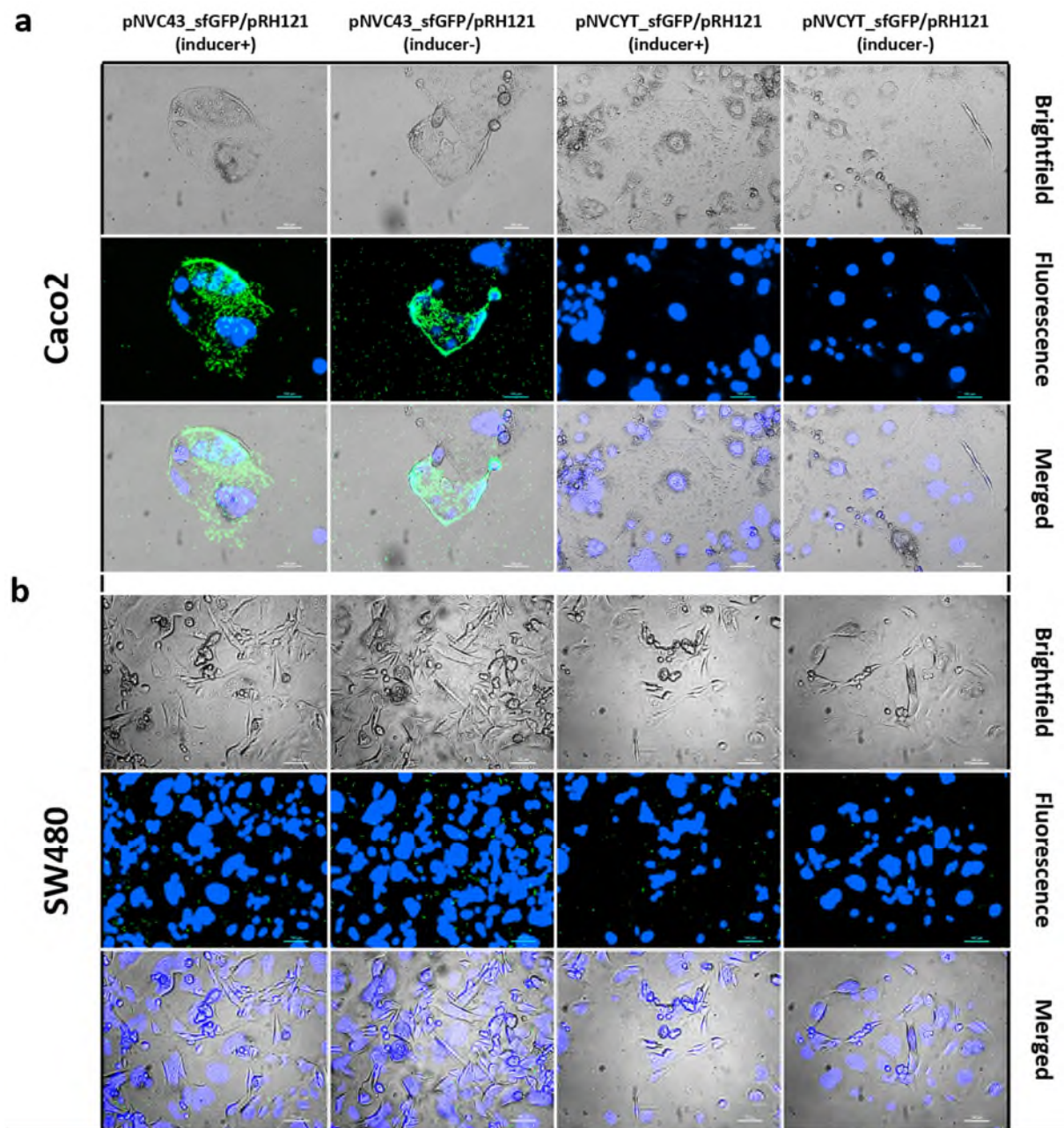


**Figure S7. (a)** Overnight OD600 reading of SimCell pYB1 carrying pNVC43\_sfGFP after two hours induction of inducer cocktail X (inducer +) or without inducer (inducer -). Each line represents a single biological repeat. **(b)** Induced and uninduced SimCell pYB1 culture plated on LB-Kan-Cm agar for 24 hours (top), 48 hours (mid) and 72 hours (bottom) at 37 °C.



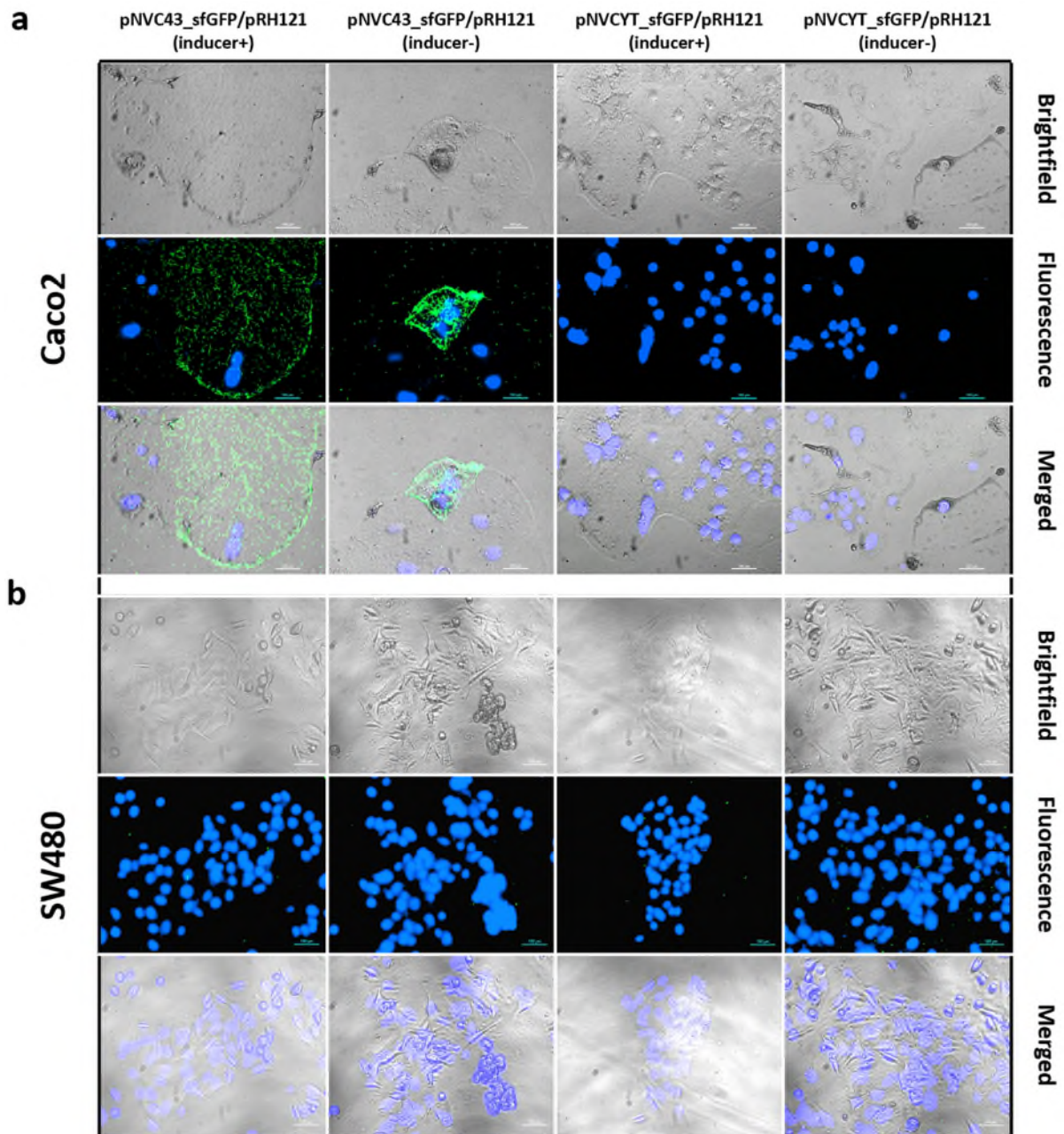


**Figure S8.** 0 hour (~within 5 minutes) incubation of programmed SimCell pYB1 carrying pNVC43\_sfGFP or pNVCYT\_sfGFP with (+) and without (-) the addition of inducer cocktail X (green). Bacterial cells under inducer+ are genome less and non-dividing SimCells. **(a)** Caco2 and **(b)** SW480 are the high CEA expressing and low CEA expressing colorectal cancer cell lines, respectively. Nuclei were stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.



**Figure S9.** Time 4 hours incubation of programmed SimCell pYB1 carrying pNVC43\_sfGFP or pNVCYT\_sfGFP with (+) and without (-) the addition of inducer cocktail X (green). Bacterial cells under inducer+ are genome less and non-dividing SimCells. **(a)** Caco2 and **(b)** SW480 are the high CEA expressing and low CEA expressing colorectal cancer cell lines, respectively. Nuclei were stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.



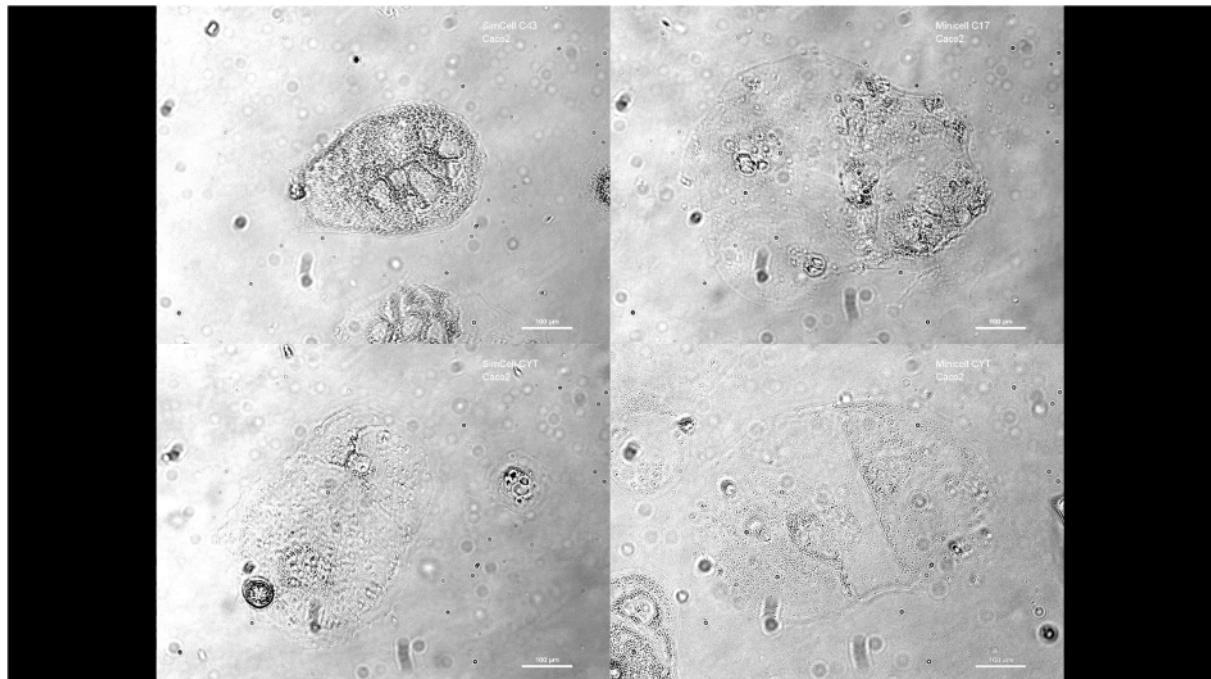


**Figure S10.** 8 hours incubation of programmed SimCell pYB1 carrying pNVC43\_sfGFP or pNVCYT\_sfGFP with (+) and without (-) the addition of inducer cocktail X (green). Bacterial cells under inducer+ are genome less and non-dividing SimCells. **(a)** Caco2 and **(b)** SW480 are the high CEA expressing and low CEA expressing colorectal cancer cell lines, respectively. Nuclei were stained with Hoechst dye (blue). Scale bar is 100  $\mu$ m.



(Please access the video here:

<https://1drv.ms/v/s!Atd6LwNkjPrwx158BYJ5FBiO6hxn?e=aladcC>)



**Movie 1.** Time lapse video of Caco2 after 2 hours incubation with SimCell + pNVC43\_sfGFP (top left); SimCell + pNVCYT\_sfGFP (bottom left); minicell + pNVC17\_sfGFP (top right); minicell + pNVCYT\_sfGFP (bottom right). All cultures were washed thrice and added with fresh media supplemented with Ethidium homodimer prior to imaging. The duration of the experiment is labelled in hh:mm:ss format on the top left. SimCell and minicell are in green and the nuclei of Caco2 with compromised membrane were stained by Ethidium homodimer (red). Scale bar is 100  $\mu$ m.

## References

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