



**Supplementary: A strategy to optimize the generation of stable chromobody cell lines for visualization and quantification of endogenous proteins in living cells**

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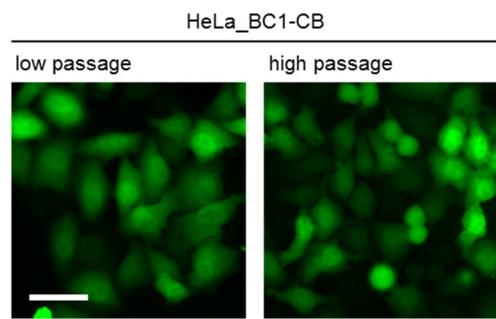
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AAVS1-CB-donor fragment (gBlock® gene fragment)	2	GCAGGCATGCTGGGATGCGGTGGCTATGGACTAGTATTAAATCTAGAGGC GGCAATTGTTCACTCCTCAGGTGCAGGC AAATCTAGAGGCAGGAATTGTTCACTCCTCAGGTGCAGGCTGCCATCAGAAGGGTGGTGGCTGGTGGCAATGCCCTGGCTCACAAATACCACGTGAGATCTTTTCCCTCTGCCAAAATTATGGGGACATCATGAAGGCCCTGAGCAGTCAGCTGACTTCGGCTAACAAAGGAAATTATTTCTATTGCAATAGTGTGTTGGAATTTTTGTGTCCTCACTCGGAAGGACATATGGGAGGGCAAATCATTTAAACATCAGAATGAGTATTGGTTAGAGTTGGCAACATATGCCCATATGCTGGCTGCCATGAACAAAGGTGGCTATAAGAGGTCACTAGTATATGAAACAGCCCCCTGCTGTCCATTCTTATTCCATAGAAAAGCCTGACTTGAGGTTAGATTTTTTATATTGTTTGTGTTATTCTTAAACATCCCTAAATTTCTTAGATGTTTACTAGCCAGATTTCCTCCTCTCCTGACTACTCCCAGTCAGCTGTCCTCTCTCGT CGACAGTACTAAGCTTGACAGAAAAGCCCACCTTCTAGGCTCCTCTCTAGTCTCCTGATATTGGGCTAACCCCCACCTCCCTGAGCAGATTCTTATCTGGTACACACCCCCATTCTCTGGAGCCATCTCTCTGCCAGAACCTCTAACGGTTGCTTACGATGGAGCCAGAGAGCATCCTGGGAGGGAGAGCTTGGCAGGGGTGGAGGGAAAGGGGGGATGCGTACCTGCCGTTCTCAGTGGCACCCCTGCGTACCCCTCTCCAGAACCTGAGCTGCTCTGACGCCGCTGTGCTGGTCTACTGATCCTGGTCTCCAGCTTCTTACACTTCCAAGAGGAGAACGAGTTGGAAAAACAAAATCAGAATAAGTGGCTCTGAGTTCTAACTTGGCTCTCACCTTCTAGTCCCCAATTATATTGTTCTCGTGCCTAGTTACCTGAGATAAGCCCAGTAGCCAGCCCCCTCCTGCCAGGGCTGTGGTACGGAGGGGGGTGTCCTCTCTGAGGAAATGGTCTTGTGAGAATGGTGTGCTTAGGTGTTACCGAGTCCTGGCCCTACTCTCTTCTCTCCATCCTTCTTAAAGAGTCCCCAGTGTCTGGAGCATATTCTCCGCCAGAGCAGGGTCCCGCTCCCTAACGGCCCTGCTCTGCTCTGCTCTGCTCAATTGTTGTTAACCTGTTATTGCAAGCTTATAATGG

Table S1: List of DNA oligonucleotides and synthesized gene fragments used in this study

	Expression construct
1.	pEGFP-N1 (Clontech)
2.	Ub-M-BC1-CB [1]
3.	Ub-R-BC1-CB [1]
4.	LMN-CB [2]
5.	Ub-M-LMN-CB (this study)
6.	Ub-R-LMN-CB (this study)
7.	(EF1 $\alpha$ )-Ub-R-BC1-CB (this study)
8.	( $\beta$ -actin)-Ub-R-BC1-CB (this study)
9.	mCherry [3]
10.	mCherry-CTNNB1 [1]
11.	Cas9_gRNA [4]
12.	AAVS1_Ub-R-BC1-CB (this study)
13.	Actin-CB [5]
14.	AAVS1_Ub-R-ACT-CB (this study)

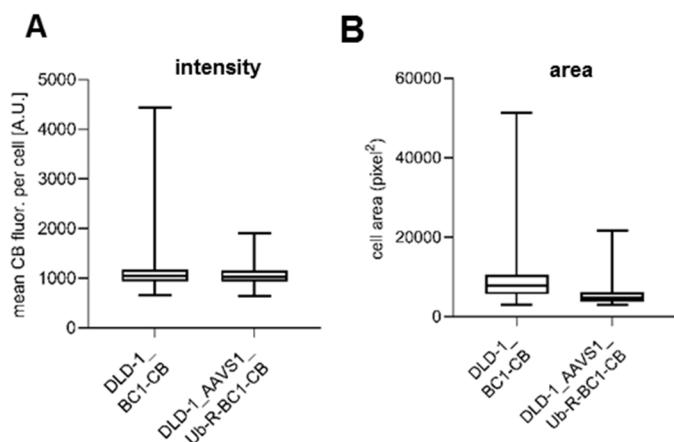
Table S2: List of mammalian expression constructs used in this study

## Supplementary Figures



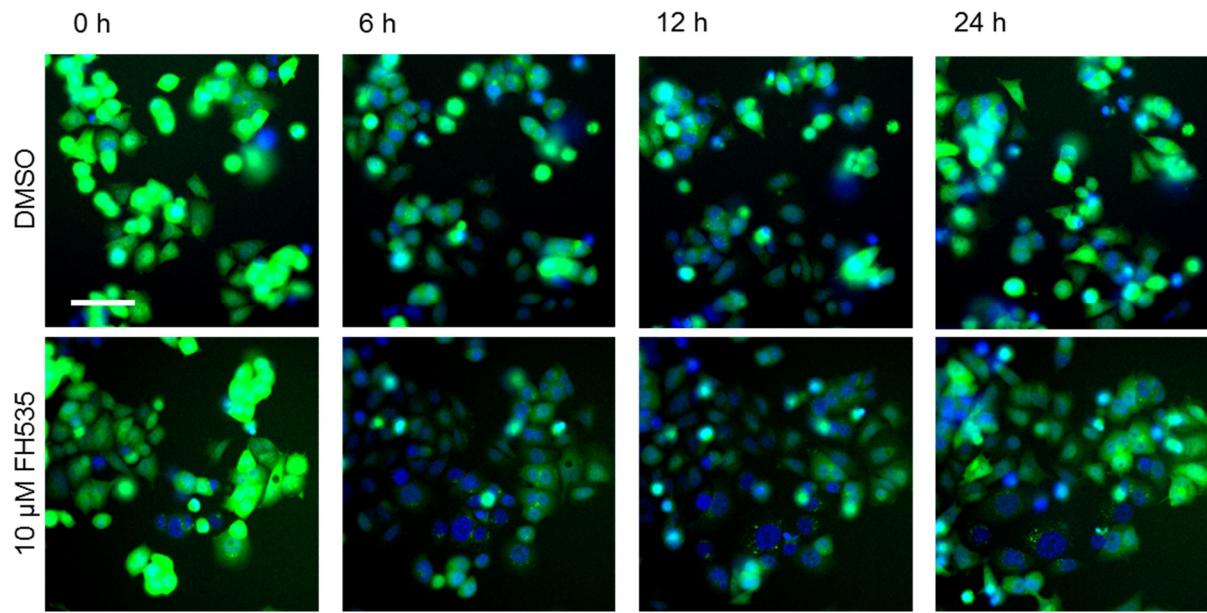
**Figure S1 Stable HeLa\_BC1-CB cell line displays heterogeneity in CB expression at high cell passages.**

Illustration of HeLa cells stably expressing the BC1-CB driven by the CMV promoter at an early passage (passage number 4, left side) and at a high passage (passage number 20, right side). Scale bar: 50  $\mu$ m.



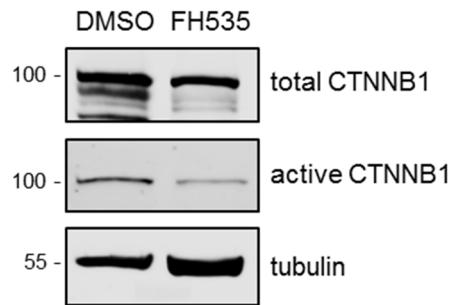
**Figure S2 Population-wide analysis of BC1-CB signal in DLD-1\_BC1-CB cells and DLD-1\_AAVS1\_Ub-R-BC1-CB.**

Fluorescence images of DLD-1\_BC1-CB cells and DLD-1\_AAVS1\_Ub-R-BC1-CB (as illustrated in **Figure 6C**) were subjected to quantitative image analysis. Using a customized image segmentation algorithm (see Material and Methods), the mean fluorescence intensity of the CB (A) and the cell area (B) were determined for each individual cell imaged. Number of analyzed cells: DLD-1\_BC1: n=2478; DLD1\_AAVS1\_Ub-R-BC1-CB: n=2090.



**Figure S3** Fluorescence images of DLD-1\_AAVS1\_Ub-R-BC1-CB cells upon compound treatment.

DLD-1\_AAVS1\_Ub-R-BC1-CB were either treated with 10  $\mu$ M FH535 or 0.01 % DMSO as control and were continuously imaged every 2 h for up to 24 h. Representative fluorescence images are shown after 0, 6, 12 and 24 h, scale bar: 50  $\mu$ M.



**Figure S4: FH535 reduces the amount of active CTNNB1 in DLD-1\_AAVS1\_Ub-R-BC1-CB cells.**

DLD-1\_AAVS1\_Ub-R-BC1-CB were either treated with 10  $\mu$ M FH535 or 0.01 % DMSO as control for 24 h. Cells were lysed using lysis buffer comprising 0.5% NP-40 and equal protein amounts of the soluble fraction were subjected to SDS-PAGE followed by immunoblot analysis using antibodies specific for total CTNNB1, active CTNNB1 and tubulin as loading control.

## References

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