

Supplementary Information

Establishing live-cell single-molecule localization microscopy imaging and single-particle tracking in the archaeon *Haloferax volcanii*

Bartosz Turkowyd*¹, Sandra Schreiber*², Julia Wörtz², Ella Shtifman Segal³, Moshe Mevarech³, Iain G. Duggin⁴, Anita Marchfelder#², Ulrike Endesfelder#^{1,5}

- **Supplementary Text 1**
- **Supplementary Text 2**
- **Supplementary Figure 1**
- **Supplementary Figure 2**
- **Supplementary Figure 3**

Supplementary Text 1

Background sensitivity of single-molecule imaging

Fluorescence microscopy studies can be compromised by autofluorescence background. While conventional fluorescence microscopy in wild-type red *Haloferax* cells is generally possible as already a few GFP together provide sufficient signal above background, single-molecule imaging is severely compromised, especially at high frame rates of dynamic imaging.

The background level is an important factor in the detection and localization of single molecules. If signals are not sufficiently brighter than the background, a large proportion of the weaker signals will remain undetected, and brighter signals that are still visible above the background are localized at lower precision, directly limiting the overall resolution of the super-resolved image.

In our measurements, WR806 cells significantly improve single-molecule imaging conditions for *Haloferax*. **Supplementary Figure 2** shows that our fluorescent proteins yield on average ~ 200-250 photons per image (corresponding to 21.000 to 27.000 AD counts), typical numbers for single-molecule imaging. The pixel size is optimized for the detection and localization of single fluorescent spots and individual spots are measured on an area of about 3x3 pixels. Thus, the average signal of a single fluorescent protein is not much stronger than the average background of 1.300 AD counts per pixel in H119 cells (**Figure 1b**), which results a 9-pixel background signal of 11.700 AD counts (translates into ~ 107 photons for our setup). In contrast, WR806 cells are indistinguishable from the background outside cells and single fluorescent protein signals can be detected above background.

Supplementary Text 2

Sequences of fluorescent proteins

Dendra2 (*E. coli* opt)

ATGAACACCCCGGTATTAACCTGATCAAAGAAGATATGCGCGTGAAGGTGCATATGGAGGGCA
ACGTGAATGGCCACGCCTTCGTGATCGAGGGCGAAGGTAAGGGCAAGCCGTATGAGGGCACCC
AGACAGCCAACCTGACCGTGAAGAGGGGCGACCGCTGCCGTTTACGCTATGACATCCTGACCAC
AGCCGTGCACTACGGTAACCGCGTTTTTACCAAGTACCCGGAGGACATCCCGGACTACTTTAAG
CAGAGCTTCCCTGAGGGCTACAGCTGGGAGCGCACCATGACCTTTGAGGACAAGGGTATCTGCA
CCATCCGCAGCGACATCAGCCTGGAGGGCGACTGTTTCTTCCAGAACGTGCGCTTCAAGGGCAC
CAACTTCCCGCCGAATGGCCCGGTGATGCAGAAAAAGACCCTGAAGTGGGAGCCGAGCACCGA
GAAATTACACGTGCGTGACGGCCTGCTGGTGGGCAACATCAACATGGCCCTGCTGTTAGAGGGC
GGCGGCCACTATCTGTGCGACTTCAAGACCACCTACAAGGCCAAGAAGGTGGTGCAGCTGCCG
GACGCACACTTTGTGGACCACCGCATCGAAATCCTGGGCAACGACAGCGACTACAACAAGGTGA
AGTTATACGAGCACGCAGTTGCCCGCTATAGCCCGCTGCCTAGCCAGGTTTGGTAA

Dendra2Hfx

ATGAACACGCCGGGCATCAACCTCATCAAAGAGGACATGCGCGTCAAGGTCCACATGGAAGGCA
ACGTCAACGGCCACGCGTTCGTATCGAAGGCGAAGGCAAGGGCAAGCCGTACGAAGGCACGC
AGACGGCGAACCTCACGGTCAAAGAAGGCGCCCCGCTCCCGTTCTCGTACGACATCCTCACGAC
GGCGGTCCACTACGGCAACCGCGTGTTCACGAAGTACCCGAGGACATCCCGGACTACTTCAAG
CAGTCGTTCCCGGAGGGCTACTCGTGGGAGCGCACGATGACGTTGAGGACAAGGGCATCTGC
ACGATCCGCTCGGACATCTCGCTCGAAGGCGACTGCTTCTTCCAGAACGTCCGCTTCAAGGGCA
CGAACTTCCCGCCGAACGGCCCGGTATGCAGAAAAAGACGCTCAAGTGGGAGCCGTGACCCG
AGAAGCTCCACGTCCGCGACGGCCTGCTCGTCCGCAACATCAACATGGCGCTGCTCCTCGAAG
GCGGCGGCCACTACCTCTGCGACTTCAAGACGACGTACAAGGCGAAGAAGGTGCTCCAGCTCC
CGGACGCGCACTTCGTGACACCAGCAATCGAGATCCTCGGCAACGACTCGGACTACAACAAGT
CAAGCTCTACGAGCACGCGGTGCGCGATACTCGCCGCTCCCGTCCGCAAGTCTGGTAA

| | | |
|------------|-----|--|
| Dendra2 | 1 | ATGAACACCCCGGGTATTAACCTGATCAAAGAAGATATGCGCGTGAAGGTGCATATGGAGGGCA |
| Dendra2Hfx | 1 | ATGAACACGCCGGGCATCAACCTCATCAAAGAGGACATGCGCGTCAAGGTCCACATGGAAGGCA |
| Dendra2 | 61 | GGCAACGTGAATGGCCACGCCTTCGTGATCGAGGGCGAAGGTAAGGGCAAGCCGTATGAGGGCACCC |
| Dendra2Hfx | 61 | GGCAACGTCAACGGCCACGCGTTCGTGATCGAAGGCGAAGGCAAGGGCAAGCCGTACCGAA |
| Dendra2 | 121 | GGCACCAGACAGCCAACCTGACCGTGAAGAGGGGCGAACCGCTGCCGTTTACGCTATGACATCCTGACCAC |
| Dendra2Hfx | 121 | GGCACGCAGACGGCGAACCTCACGGTCAAAGAAGGCGCCCCGCTCCCGTTCTCGTACGACATCCTCACGAC |
| Dendra2 | 181 | ATCCTGACCCACAGCCGTGCACTACGGTAACCCGCGTTCACCAAGTACCCGGAGGACATCCTGACATCCTGACCAC |
| Dendra2Hfx | 181 | ATCCTCACGACGGCGGTCCACTACGGCAACCCGCGTTCACCAAGTACCCGGAGGACATCCTGACATCCTGACCAC |
| Dendra2 | 241 | CCGGACTACTTTAAGCAGAGCTTCCCTGAGGGCTACAGCTGGGAGCGCACCATGACCTTTTCCGCTTCAAGCAGT |
| Dendra2Hfx | 241 | CCGGACTACTTCAAGCAGTTCGTTCCCGGAGGGCTACTCGTGGGAGCGCACCATGACCTTTTCCGCTTCAAGCAGT |
| Dendra2 | 301 | GAGGACAAGGGTATCTGCACCATCCGCAAGCGACATCAGCCTGGAGGGCGACTGTTTCTTCCGCTTCAAGCAGT |
| Dendra2Hfx | 301 | GAGGACAAGGGCATCTGCACGATCCGCTCGGACATCAGCCTCGAAGGGCGACTGCTTCTTCCGCTTCAAGCAGT |
| Dendra2 | 361 | CAGAACGTGCGCTTCAAGGGCACCAACTTCCCGCCGAATGGCCCGGTGATGCAGAAAAAG |
| Dendra2Hfx | 361 | CAGAACGTCCGCTTCAAGGGCACCAACTTCCCGCCGAATGGCCCGGTGATGCAGAAAAAG |
| Dendra2 | 421 | ACCCTGAAGTGGGAGCCGAGCACCGAGAAATTAACACGTGCGTGACGGCCTGCTGGTGGGC |
| Dendra2Hfx | 421 | ACCCTCAAGTGGGAGCCGTCGACCGAGAAATTAACACGTGCGTGACGGCCTGCTGGTGGGC |
| Dendra2 | 481 | AACATCAACATGGCCCTGCTGTAGAGGGCGGGCCACTATCTGTGCGACTTCAAGACC |
| Dendra2Hfx | 481 | AACATCAACATGGCCCTGCTGTGAGGGCGGGCCACTATCTGTGCGACTTCAAGAGG |
| Dendra2 | 541 | ACCTACAAGGCCAAGAAGGTGGTGCAGCTGCCGGACGCAACTTTGTTGGACCACCGCATC |
| Dendra2Hfx | 541 | ACGTACAAGGCCAAGAAGGTGCTCCAGCTGCCGGACGCAACTTTGTTGGACCACCGAATC |

Dendra2 601 GAAATCCTGGGCAACGACAGCGACTACAACAAGGTGAAGTTATACGAGCACGCAGTTGCC
Dendra2Hfx 601 GACATCCTCGGCAACGACTCGGACTACAACAAGTCAAGCTCTACGAGCACGCCGTCGCG

Dendra2 661 CGCTATAGCCCGCTGCC TAGCCAGGTTTGGTAA
Dendra2Hfx 661 CGATACTCGCCGCTCCCGTCAAGTCTGGTAA

PAmCherry1 (E. coli opt)

ATGGTAAGCAAAGGCGAAGAGGACAACATGGCGATCATTAAAGGAGTTCATGCGCTTTAAGGTTACAT
GGAGGGTAGCGTGAATGGTTCATGTGTTTAAAATTGAGGGTGAGGGCGAGGGTCGTCCGTATGAAGGT
ACCCAAACGCCAAACTGAAGGTCACGAAAGGTGGCCCGCTGCCGTTCACTTGGGACATCTTGAGCC
CGCAATTCATGTACGGCTCCAACGCATACGTCAAACACCCAGCAGACATTCCGGACTACTTTAAACTG
AGCTTTCCGGAAGGCTTTAAGTGGGAGCGCGTTATGAAATTCGAAGATGGTGGTGTGTGTCACCGTCAC
GCAGGACAGCAGCCTGCAGGATGGTGAAGTTCATCTATAAAGTGAAGTGCAGCGGTACCAACTTCCCG
TCTGATGGCCCGTTATGCAAAAAGAAAACCATGGGCTGGGAGGCACTGTCCGAGCGTATGTACCCGG
AAGATGGCGCGCTGAAAGGTGAGGTTAAGCCTCGTGTGAAACTGAAAGATGGCGGTCACTATGACGC
GGAAGTTAAGACGACCTATAAGGCTAAGAAGCCGGTCCAAGTCCCGGTGCGTACAATGTGAATCGT
AAGTTGGACATCAGGAGCCATAACGAGGACTACACCATTGTGGAACAGTATGAGCGTGCCGAAGGTC
GTCACAGCACCGGTGGCATGGATGAACTGTACAAGTAA

PAmCherry1Hfx

ATGGTCAGCAAGGCGAAGAGGACAACATGGCGATCATCAAAGAATTCATGCGCTTCAAGGTCCACA
TGGAAGGCTCGGTCAACGGCCACGTGTTTCGAGATCGAGGGCGAGGGCGAAGGCCGACCGTACGAAG
GCACGCAGACGGCGAAGCTCAAAGTCACGAAGGTGGCCCGCTCCCGTTCACGTGGGACATCCTCT
CGCCGCAGTTCATGTACGGCTCGAACGCGTACGTCAAGCACCCGGCGGACATCCCGGACTACTTCAA
GCTCTCGTTCGCCGAGGGCTTCAAGTGGGAGCGCGTTCATGAAGTTCGAGGACGGCGGGCGTTCGTCAC
GGTCACGCAGGACTCGTCGCTGCAGGACGGCGAGTTCATCTACAAAGTCAAGTCCGCGGCACGAA
CTTCCCGTTCGACGGCCCGGTTCATGCAGAAAAGACGATGGGCTGGGAGGCGCTCTCGGAGCGCAT
GTACCCCGAGGACGGGGCGCTCAAAGGCGAAGTCAAGCCGCGCGTCAAGTCAAAGACGGCGGCCA
CTACGACGCGGAAGTCAAGACGACGTACAAGGCGAAGAAGCCGGTCCAGCTCCCGGTGCCTACAA
CGTCAACCGCAAGCTCGACATCACGTGCGACAACGAGGACTACACGATCGTCGAGCAGTACGAGCGC
GCGGAAGGCCGACACTCGACGGGCGGCATGGACGAACTCTACAAATGA

PAmCherry1 1 ATGGTAAGCAAAGGCGAAGAGGACAACATGGCGATCATTAAGGAGTTCATGCGCTTTAAG
PAmCherry1Hfx 1 ATGGTCAGCAAGGCGAAGAGGACAACATGGCGATCATCAAAGAATTCATGCGCTTCAAG

PAmCherry1 61 GTTTCACATGGAGGGTAGCGTGAATGGTTCATGTGTTTAAAATTGAGGGTGAGGGCGAGGGT
PAmCherry1Hfx 61 GTCCACATGGAAAGGCTCGGTCACCGCCACGTGTTTCGAGATCGAGGGCGAGGGCGAAGG

PAmCherry1 121 CGTCCGTATGAAGGTACCCAAACCGCCAAACTGAAAGTTCACGAAAAGGTGGCCCGCTGCCG
PAmCherry1Hfx 121 CGACCGTACGAAGGCACGCAGACGGCGAAGCTCAAAGTTCACGAAAAGGTGGCCCGCTCCCG

PAmCherry1 181 TTCACTTGGGACATCTTTGAGCCCGCAGTTCATGTACGGTCCAACGCAATACGTCAAACAC
PAmCherry1Hfx 181 TTCACGTGGGACATCTCTCGCCCGCAGTTCATGTACGGTCCAACGCAATACGTCAAACAC

PAmCherry1 241 CCAGCAGACATTCCGGACTACTTTAAACTGAGCTTTCGGGAAGGCTTTAAGTGGGAGCGC
PAmCherry1Hfx 241 CCGGCGGACATCCCGGACTACTTCAAAGCTCTCGTTCGCCGAGGCTTCAAGTGGGAGCGC

PAmCherry1 301 GTTATGAAATTCGAAGATGGTGGTGTGTGTCACCGTCACGCAGGACAGCAGCCTGCAGGAT
PAmCherry1Hfx 301 GTCATGAAGTTCGAGGACGGCGGCGTTCGTCACGGTTCACGCAGGACTCGTTCGCTGCAGGAC

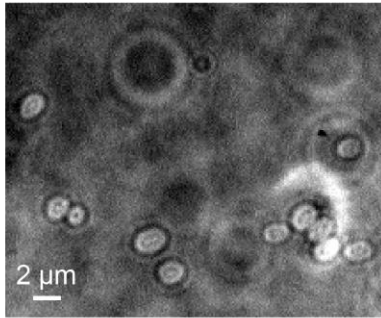
PAmCherry1 361 GGTGAGTTCATCTATAAAGTGAAGTTCGCGGGTACCAACTTCCCGTCTGATGGCCCGGTT
PAmCherry1Hfx 361 GGCAGGTTTCATCTACAAGTCAAGCTCCGCGGCACGAAGTTCCCGTCGACGGCCCGGTT

PAmCherry1 421 ATGCAAAAAGAAAACCATGGGCTGGGAGGCACTGTCCGAGCGTATGTACCCGGAAGATGGC
PAmCherry1Hfx 421 ATGCAGAAAAGAACGATGGGCTGGGAGGCGCTTCGCGAGCGCATGTACCCGGAAGATGGC

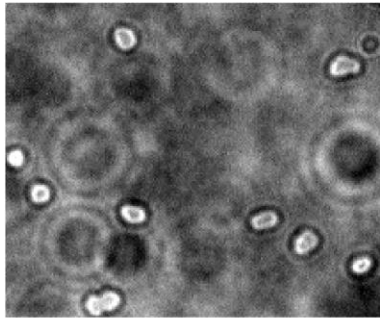
PAmCherry1 481 GCGCTGAAAGGTGAGGTTAAGCCTCGTGTGAAACTGAAAGATGGCGGTCACTATGACGCG
PAmCherry1Hfx 481 GCGCTCAAAGGCGAAGTCAAGCCGCGCGTCAAAGCTCAAAGACGGCGGCCTACTACGACGCG

| | | | | | | | | | | | | | | |
|---------------|-----|-------|-----------|----------|--------|-----------------|-----------------|-----|----------------|----------|--------|---------|--------|-----|
| PAmCherry1 | 541 | GAAGT | T | AAGACGAC | CTAT | AAGGCT | AAGAAGCCGGTCCAA | CT | G | CCGGGTGG | TACAAT | GTG | | |
| PAmCherry1Hfx | 541 | GAAGT | CAAGACGAC | GTAC | AAGGCG | AAGAAGCCGGTCCAA | CT | CCC | GGTGG | TACAAC | GTG | | | |
| PAmCherry1 | 601 | AAT | CGT | AAGT | TG | GACATCACG | AGC | CAT | AACGAGGACTACAC | CAT | TGT | GAA | CAGTAT | GAG |
| PAmCherry1Hfx | 601 | AAC | CGC | AAGCT | CG | GACATCACG | TGC | CAC | AACGAGGACTACAC | GAT | CGT | CGA | CAGTAC | GAG |
| PAmCherry1 | 661 | CGT | GCC | GAAGGT | CGT | CACAGCAC | CGG | T | GGCATGGAT | GAACT | G | TACAACT | AA | |
| PAmCherry1Hfx | 661 | CGC | GCG | GAAGGC | CGA | CACACTCG | ACG | GGC | GGCATGGAC | GAACT | C | TACAA | TGA | |

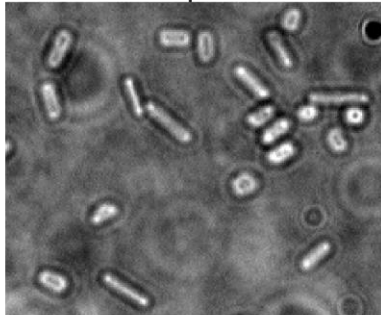
H119



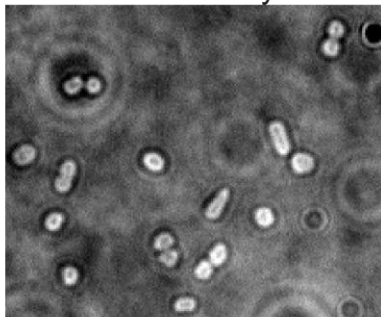
WR806



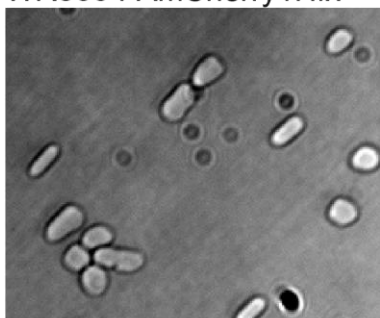
WR806 mMaple3



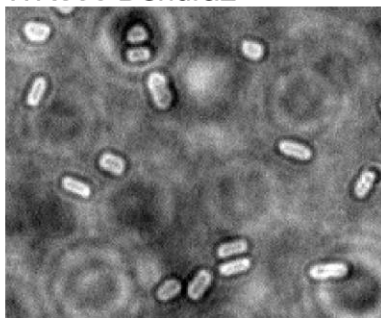
WR806 PAmCherry1



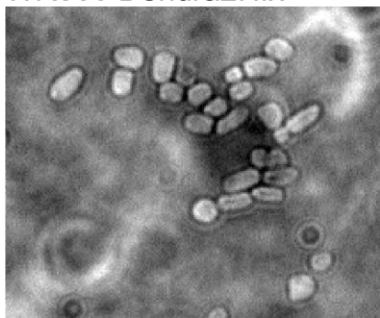
WR806 PAmCherry1Hfx



WR806 Dendra2

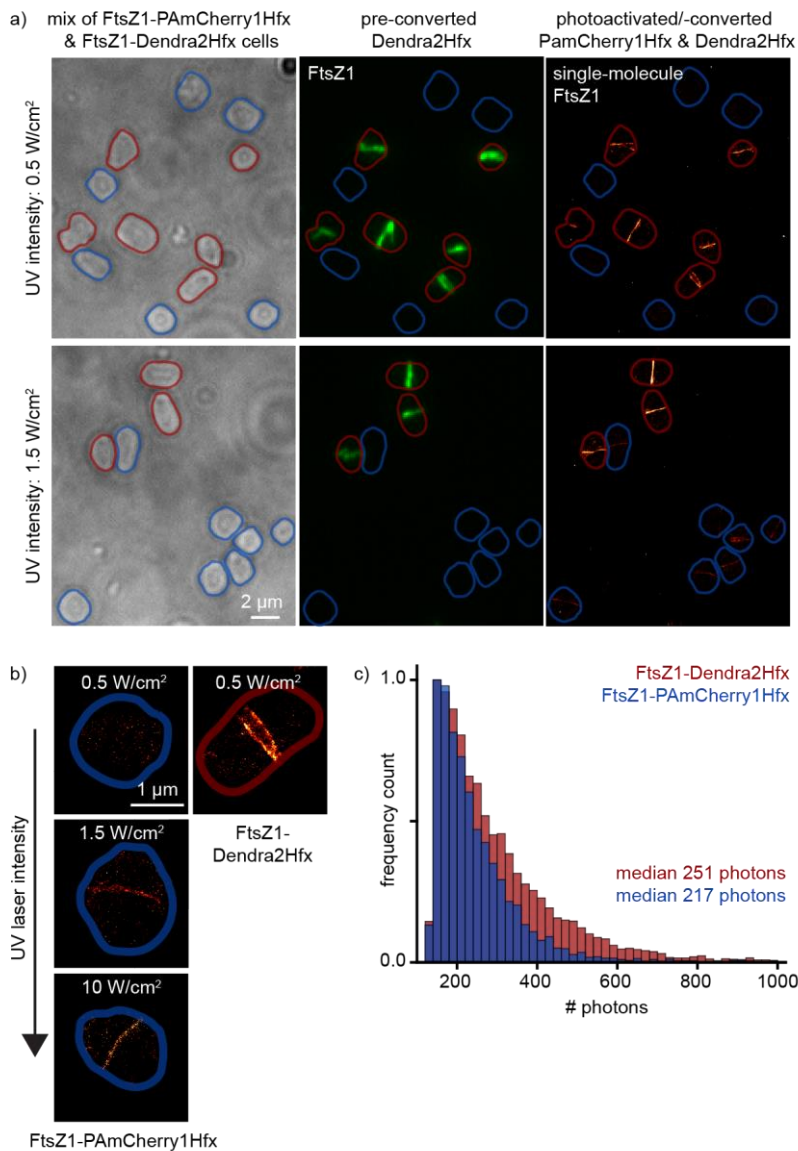


WR806 Dendra2Hfx



Supplementary Figure 1. Cell phenotype varies under the production of different fluorescent proteins

All strains producing fluorescent proteins which showed efficient photoactivation/-conversion (**Figure 1c**) showed similar phenotypes which are comparable to the parental strains H119 and WR806. In contrast, the strain producing mMaple3 shows strongly altered morphology. This and the fact that mMaple3 showed no fluorescence (**Figure 1c**) suggests improper protein folding that might cause increased cytotoxicity leading to a different growth phenotype.

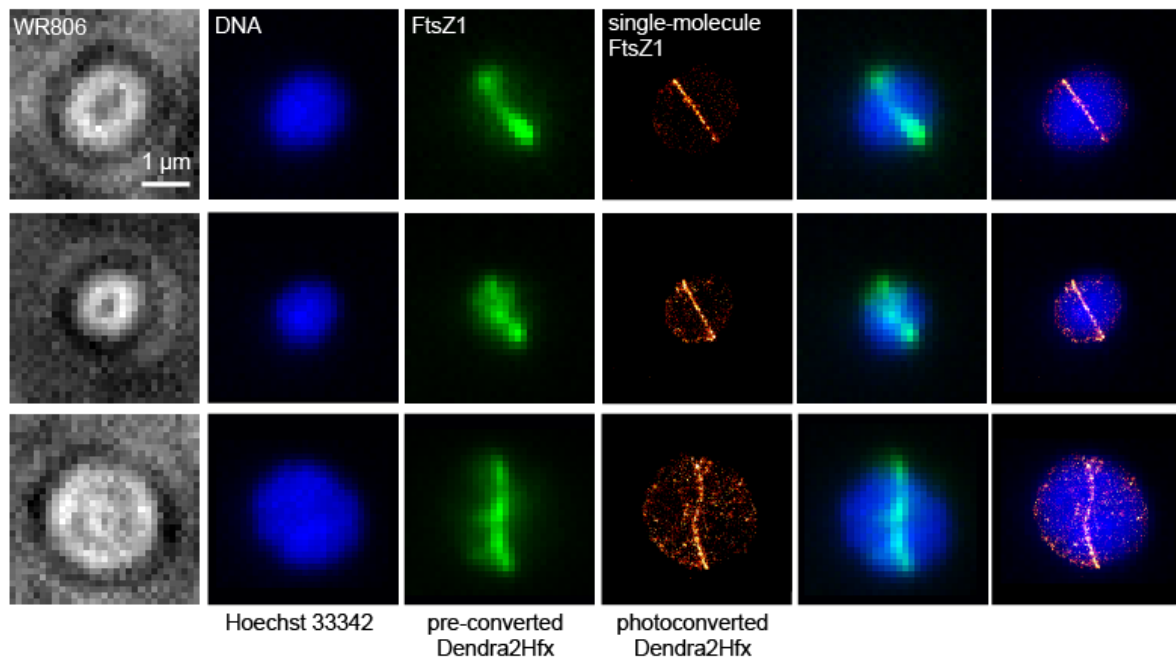


Supplementary Figure 2. Dendra2Hfx performs superior to PAmCherry1Hfx.

(a) WR806 cells producing either FtsZ1-Dendra2Hfx or FtsZ1-PAmCherry1Hfx were mixed on the same agarose pad and imaged under different UV laser intensities. When 0.5 W/cm² of the UV light was applied, FtsZ1-Dendra2Hfx revealed good photoconversion which results in dense SMLM ring structures. In contrast, FtsZ1-PAmCherry1Hfx was not photoactivated. A read-out of FtsZ1-PAmCherry1Hfx became only possible when increasing the UV light intensity to 1.5 W/cm².

(b) To provide an image quality comparable to FtsZ1-Dendra2Hfx when imaging FtsZ1-PAmCherry1Hfx, the UV light intensity had to be significantly increased. 10 W/cm² was barely sufficient to achieve the same quality as for the FtsZ1-Dendra2Hfx structure when applying 0.5W/cm².

(c) Under imaging conditions as in (a), Dendra2Hfx emits on average more photons (median of 251 photons) than PAmCherry1Hfx (median of 217 photons) per fluorescent emission in a single imaging frame.



Supplementary figure 3. Dual color imaging of FtsZ1-Dendra2Hfx and DNA stained with Hoechst 33342. As FtsZ1-Dendra2Hfx can be photoconverted by primed photoconversion, which does not rely on UV light illumination, it is possible to use the DNA-intercalating dye Hoechst 33342 as a co-stain to obtain a DNA-reference image.