

# Chemical Probes for Bioimaging

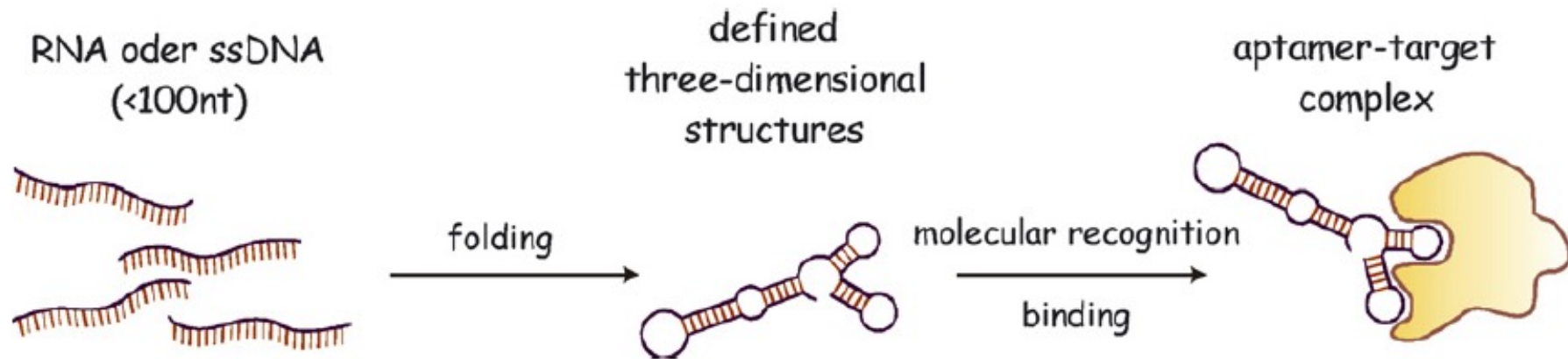
## Part 3: RNA AND DNA IMAGING

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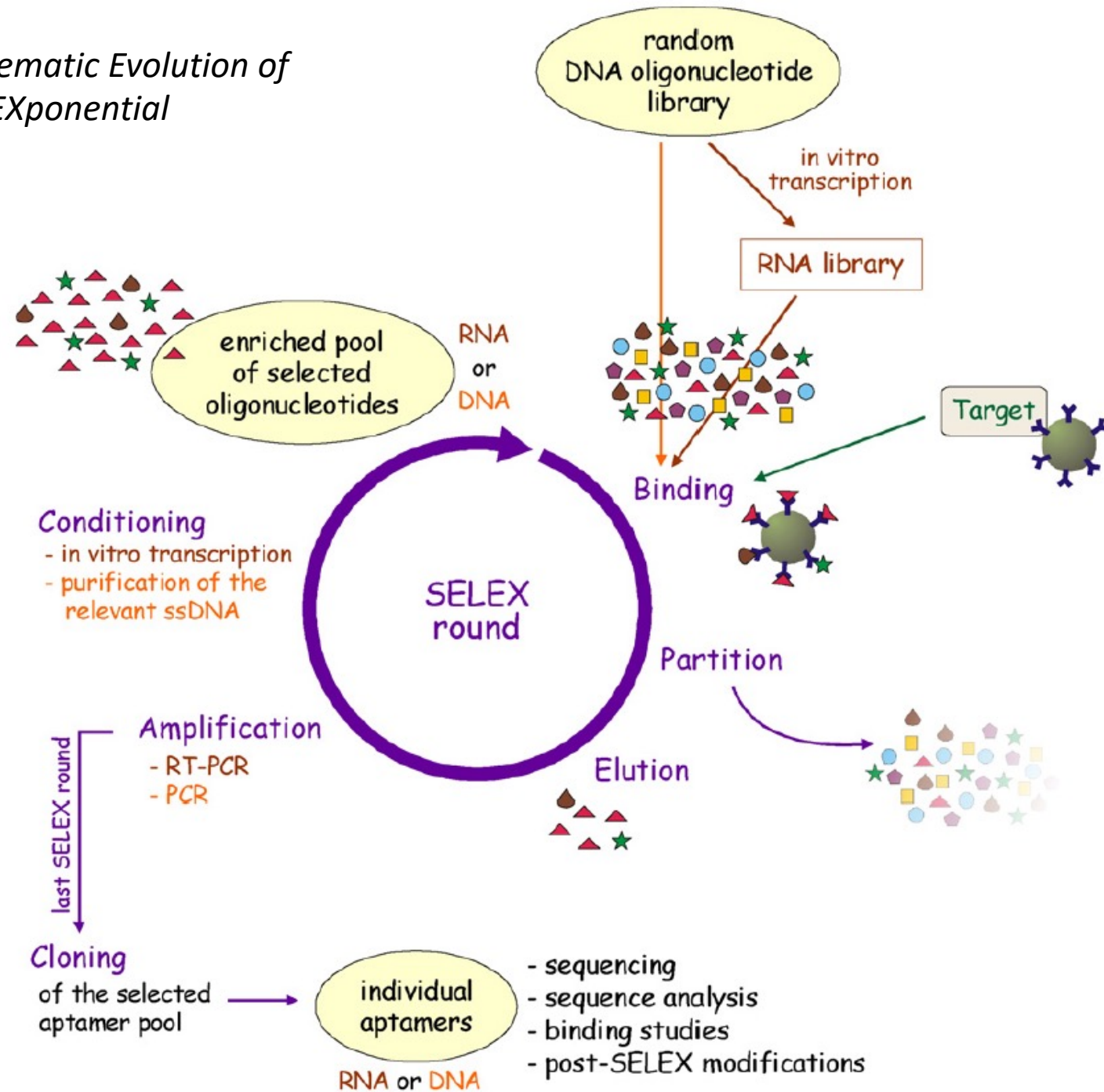
Institute of Chemical Sciences and Engineering  
EPFL, Lausanne

Single-stranded RNA and DNA can fold into defined three-dimensional structures that can bind molecules and catalyze reactions



<http://www.annualreviews.org/doi/pdf/10.1146/annurev.biochem.68.1.611>

**SELEX: Systematic Evolution of Ligands by EXponential Enrichment**



## How to make RNA libraries

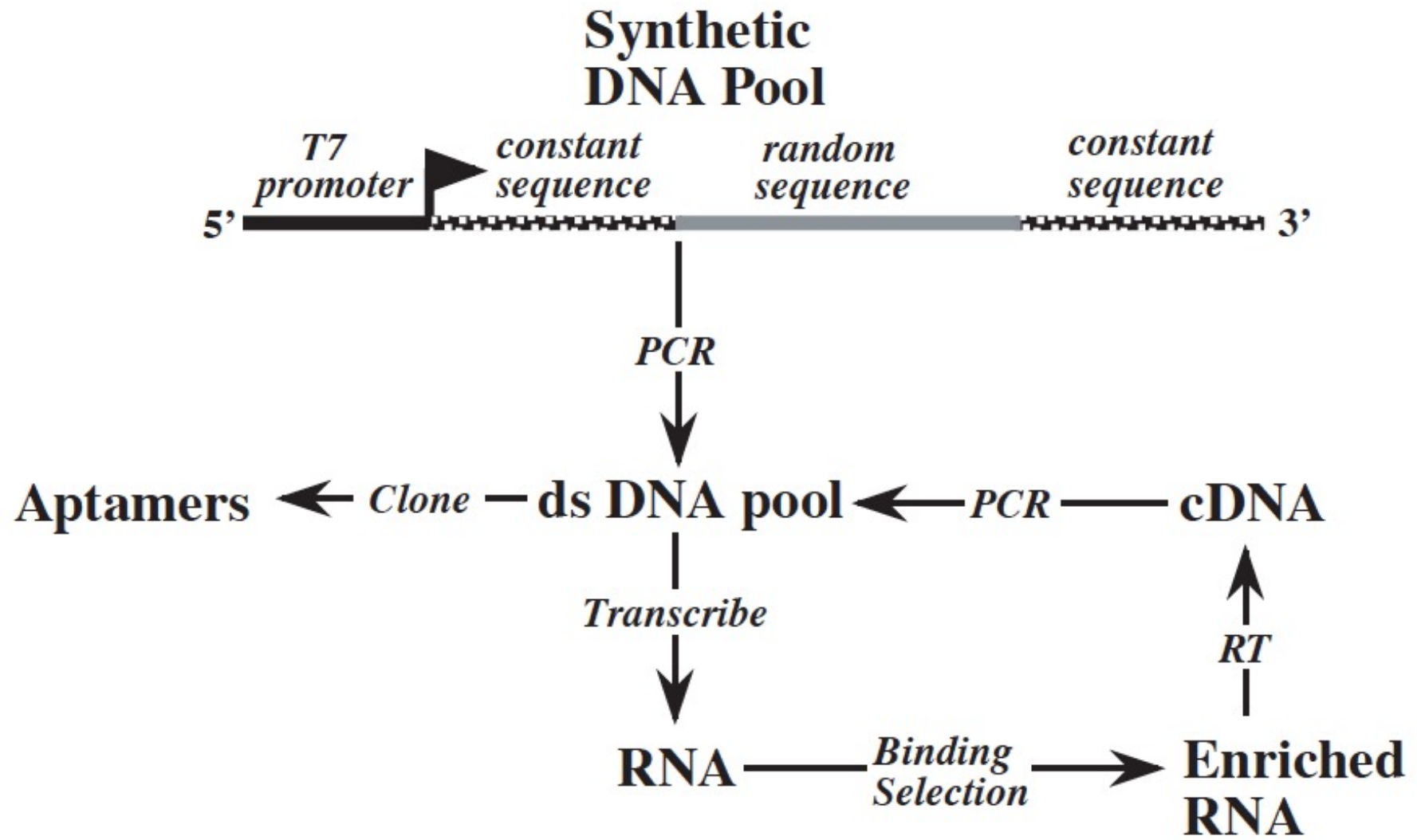




Table 1 (Continued)

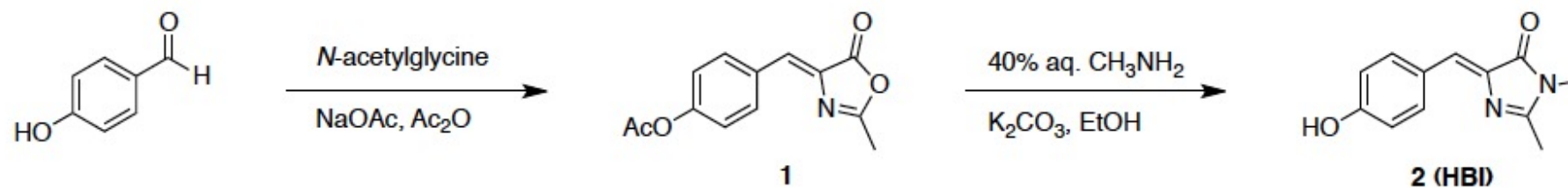
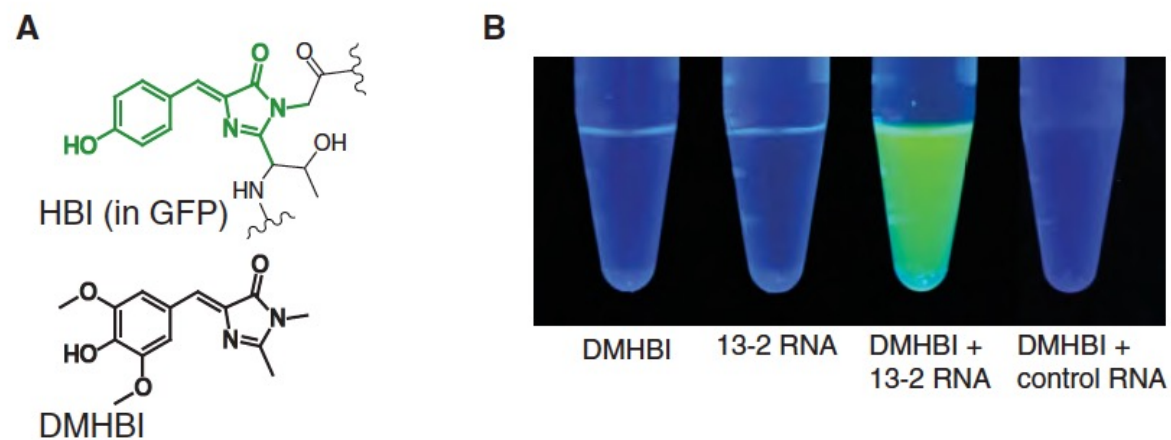
| Target for aptamer selection | Type of aptamer | $K_D$               | References                                         |
|------------------------------|-----------------|---------------------|----------------------------------------------------|
| Ricin toxin                  | DNA             | 58–105 nmol/L       | Tang et al. (2006)                                 |
| Cholic acid                  | DNA             | 5–67.5 $\mu$ mol/L  | Kato et al. (2000)                                 |
| 4,4'-Methylenedianiline      | RNA             | 0.45–15 $\mu$ mol/L | Brockstedt et al. (2004)                           |
| Dopamine                     | RNA             | 2.8 $\mu$ mol/L     | Mannironi et al. (1997)                            |
| Cocaine                      | DNA             | n.s. <sup>a</sup>   | Stojanovic et al. (2000)                           |
| Nucleotides and derivatives  |                 |                     |                                                    |
| Adenine                      | RNA             | 10 $\mu$ mol/L      | Meli et al. (2002)                                 |
| ATP (adenosine)              | RNA             | 0.7–50 $\mu$ mol/L  | Sassanfar and Szostak (1993)                       |
| Adenosine/ATP                | DNA             | 6 $\mu$ mol/L       | Huizenga and Szostak (1995)                        |
| ATP                          | RNA             | 4.8–11 $\mu$ mol/L  | Sazani et al. (2004)                               |
| Xanthine                     | RNA             | 3.3 $\mu$ mol/L     | Kiga et al. (1998)                                 |
| cAMP                         | RNA             | 10 $\mu$ mol/L      | Koizumi and Breaker (2000)                         |
| Cofactors                    |                 |                     |                                                    |
| Coenzyme A                   | RNA             | n.s. <sup>a</sup>   | Saran et al. (2003)                                |
| Cyanocobalamin               | RNA             | 88 nmol/L           | Lorsch and Szostak (1994)                          |
| Riboflavin                   | RNA             | 1–5 $\mu$ mol/L     | Lauhon and Szostak (1995)                          |
| FMN                          | RNA             | 0.5 $\mu$ mol/L     | Burgstaller and Famulok (1994)                     |
| FAD                          | RNA             | 137–273 $\mu$ mol/L | Burgstaller and Famulok (1994)                     |
| NAD                          | RNA             | n.s. <sup>a</sup>   | Burgstaller and Famulok (1994)                     |
|                              | RNA             | 2.5 $\mu$ mol/L     | Lauhon and Szostak (1995)                          |
| S-adenosyl methionine        | RNA             | n.s. <sup>a</sup>   | Burke and Gold (1997)                              |
| S-adenosyl homocysteine      | RNA             | 0.1 $\mu$ mol/L     | Gebhardt et al. (2000)                             |
| Biotin                       | RNA             | 5 $\mu$ mol/L       | Wilson et al. (1998) and Wilson and Szostak (1995) |
| Nucleic acids                |                 |                     |                                                    |
| TAR RNA element of HIV-1     | DNA             | 50 nmol/L           | Boiziau et al. (1999)                              |
|                              | RNA             | 20–50 nmol/L        | Duconge and Toulme (1999)                          |
|                              | DNA             | 50 nmol/L           | Sekkal et al. (2002)                               |
| Yeast phenylalanine tRNA     | RNA             | 12–26 nmol/L        | Scarabino et al. (1999)                            |
| <i>E.coli</i> 5S RNA         | RNA             | 6–12 $\mu$ mol/L    | Ko et al. (1999)                                   |
|                              | RNA             | 3 $\mu$ mol/L       | Ko et al. (2001)                                   |
| Amino acids                  |                 |                     |                                                    |
| L-Arginine                   | RNA             | 330 nmol/L          | Geiger et al. (1996)                               |
|                              | DNA             | ~2.5 nmol/L         | Harada and Frankel (1995)                          |
| L-Citrulline                 | RNA             | 62–68 $\mu$ mol/L   | Famulok (1994)                                     |
| L-Valine                     | RNA             | 12 nmol/L           | Majerfeld and Yarus (1994)                         |
| L-Isoleucine                 | RNA             | 1–7 nmol/L          | Lozupone et al. (2003)                             |
|                              | RNA             | 200–500 $\mu$ mol/L | Majerfeld and Yarus (1998)                         |
| D-Tryptophan                 | RNA             | 18 $\mu$ mol/L      | Famulok and Szostak (1992)                         |
| L-tyrosinamide               | DNA             | 45 $\mu$ mol/L      | Vianini et al. (2001)                              |
| L-histidine                  | RNA             | 8–54 $\mu$ mol/L    | Majerfeld et al. (2005)                            |
| Carbohydrates                |                 |                     |                                                    |
| Cellobiose                   | DNA             | 600–nmol/L          | Yang et al. (1998)                                 |
| Sialyl Lewis X               | RNA             | 0.085–10 nmol/L     | Jeong et al. (2001)                                |
| Chitin                       | DNA             | n.s. <sup>a</sup>   | Fukusaki et al. (2000)                             |
| Sialyllactose                | DNA             | 4.9 $\mu$ mol/L     | Masud et al. (2004)                                |
| Sephadex                     | DNA             | n.s. <sup>a</sup>   | Srisawat et al. (2001)                             |
| Antibiotics                  |                 |                     |                                                    |
| Kanamycin A                  | RNA             | ≤300 nmol/L         | Lato et al. (1995)                                 |
| Kanamycin B                  | RNA             | 180 nmol/L          | Kwon et al. (2001)                                 |
| Streptomycin                 | RNA             | n.s. <sup>a</sup>   | Wallace and Schroeder (1998)                       |
| Neomycin                     | RNA             | ~100 nmol/L         | Wallis et al. (1995)                               |
| Tobramycin                   | RNA             | 2–3 nmol/L          | Wang and Rando (1995)                              |
| Lividomycin                  | RNA             | ≤300 nmol/L         | Lato et al. (1995) and Lato and Ellington (1996)   |
| Moenomycin A                 | RNA             | 300–400 nmol/L      | Schürer et al. (2001)                              |
| Tetracycline                 | RNA             | 1 $\mu$ mol/L       | Berens et al. (2001)                               |
| Chloramphenicol              | RNA             | 25–65 $\mu$ mol/L   | Burke et al. (1997)                                |
| Peptides and proteins        |                 |                     |                                                    |
| T4 DNA polymerase            | RNA             | 5–30 nmol/L         | Tuerk and Gold (1990)                              |

# RNA Mimics of Green Fluorescent Protein

Jeremy S. Paige,<sup>1</sup> Karen Y. Wu,<sup>1</sup> Samie R. Jaffrey<sup>1,2\*</sup>

Green fluorescent protein (GFP) and its derivatives have transformed the use and analysis of proteins for diverse applications. Like proteins, RNA has complex roles in cellular function and is increasingly used for various applications, but a comparable approach for fluorescently tagging RNA is lacking. Here, we describe the generation of RNA aptamers that bind fluorophores resembling the fluorophore in GFP. These RNA-fluorophore complexes create a palette that spans the visible spectrum. An RNA-fluorophore complex, termed Spinach, resembles enhanced GFP and emits a green fluorescence comparable in brightness with fluorescent proteins. Spinach is markedly resistant to photobleaching, and Spinach fusion RNAs can be imaged in living cells. These RNA mimics of GFP provide an approach for genetic encoding of fluorescent RNAs.

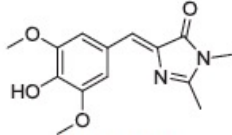
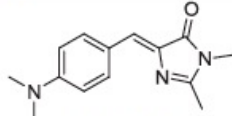
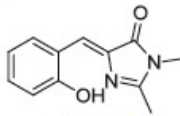
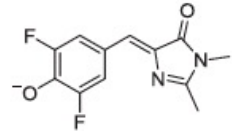
**Fig. 1.** RNA aptamers switch on the fluorescence of GFP-like fluorophores. **(A)** Structures of HBI (green), in the context of GFP, and DMHBI. **(B)** 13-2 enhances the fluorescence of DMHBI. Solutions containing DMHBI, 13-2 RNA, DMHBI with 13-2 RNA, or DMHBI with total HeLa cell RNA were photographed under illumination with 365 nm of light. The image is a montage obtained under identical image-acquisition conditions.



Scheme 1

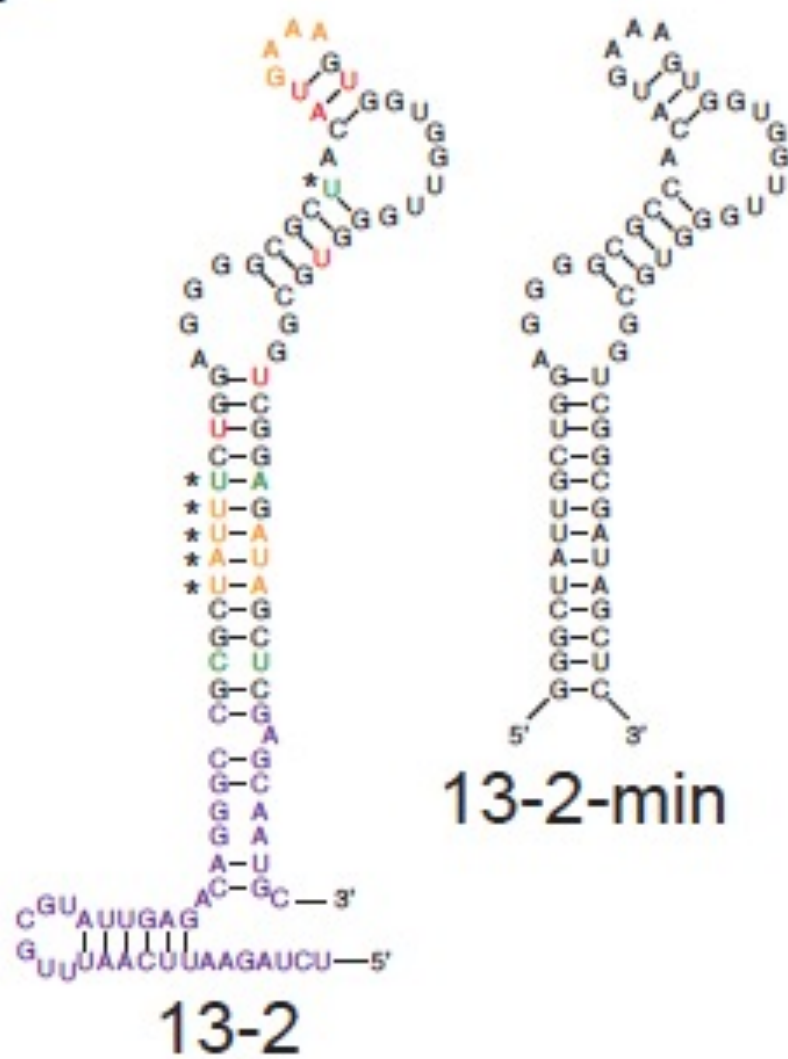


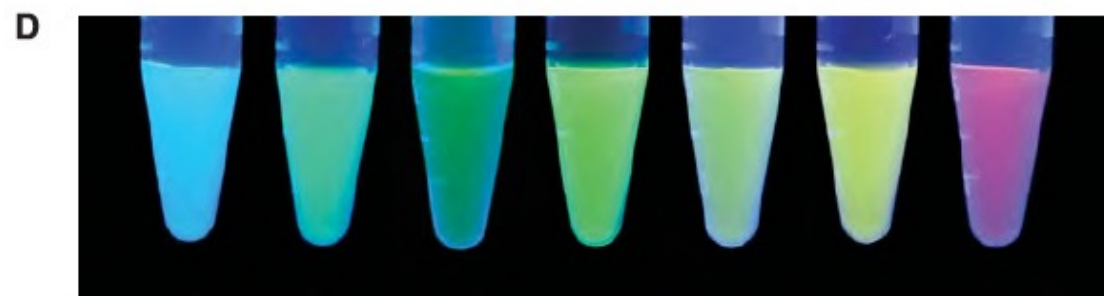
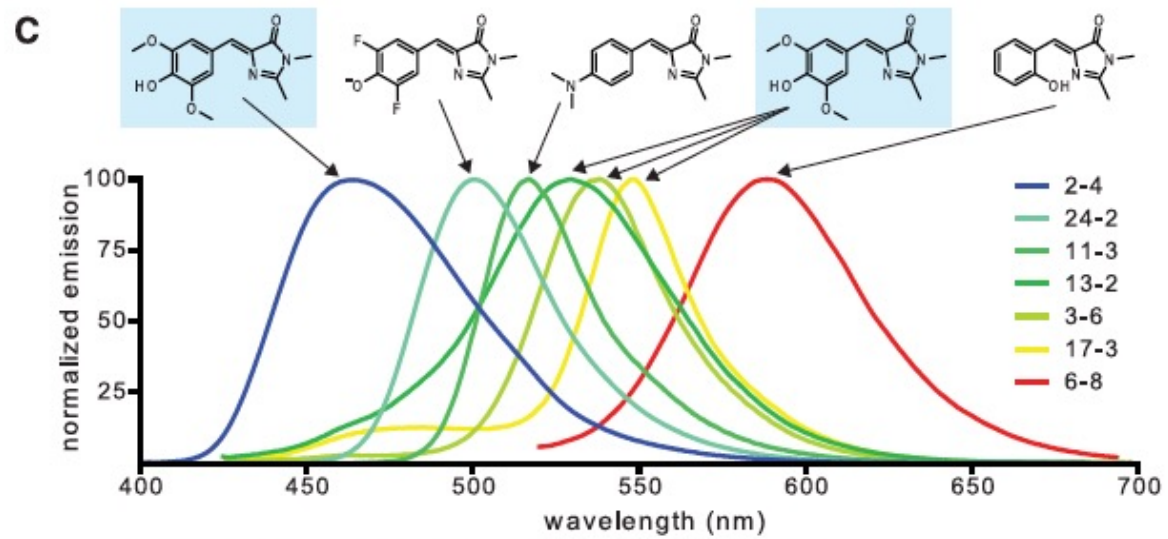
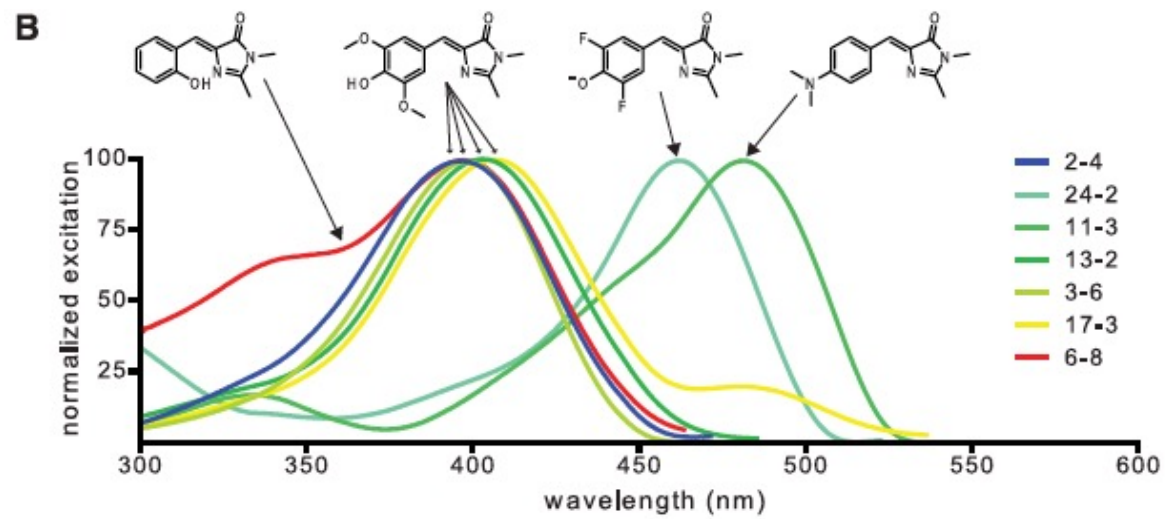
Table S1 Photophysical and binding properties of RNA-fluorophore complexes

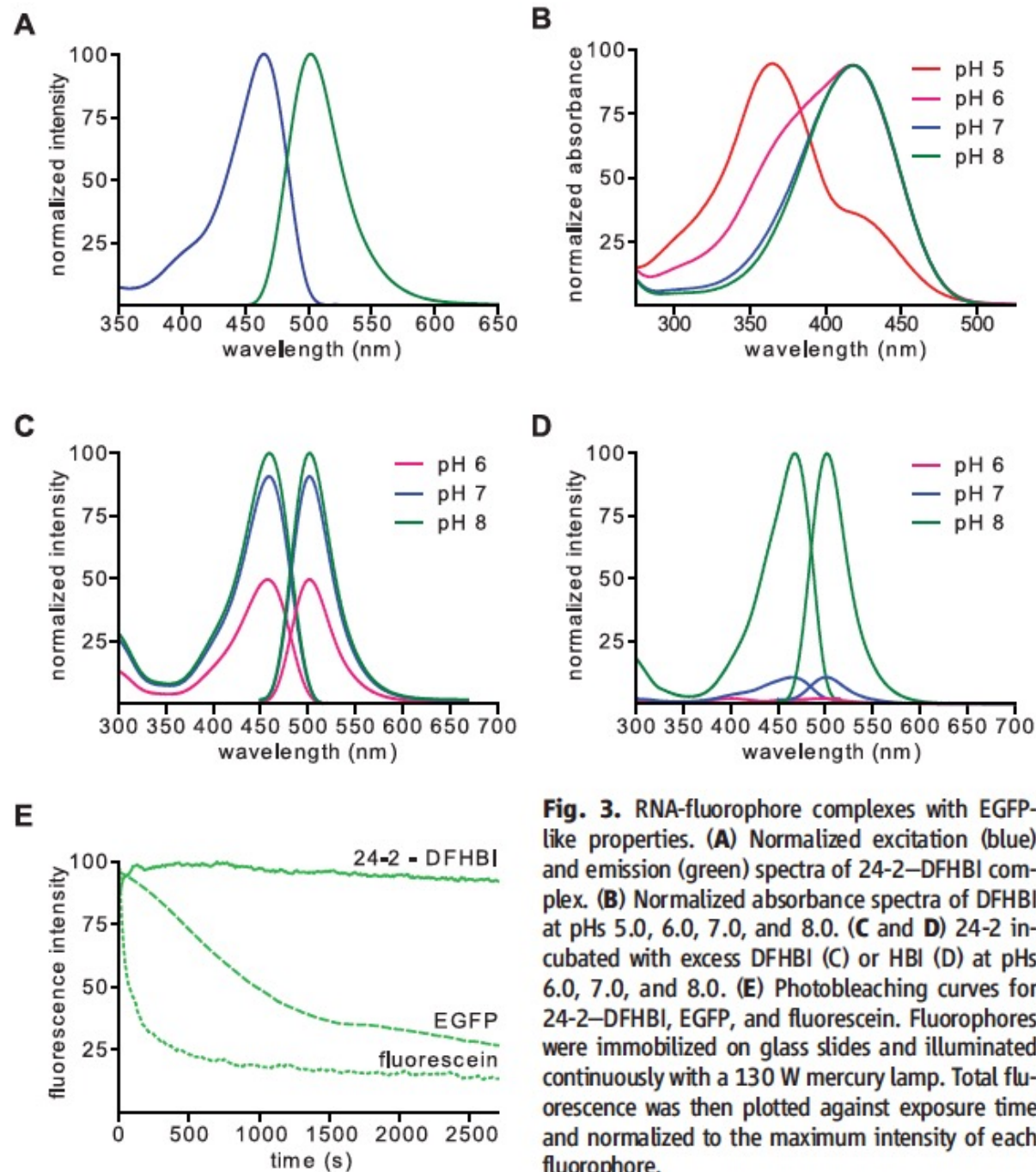
| Fluorophore                                                                                  | Excitation maximum (nm) | Emission maximum (nm) | Extinction coefficient ( $M^{-1}cm^{-1}$ ) <sup>a</sup> | Fluorescence quantum yield | $K_D$             | Brightness <sup>b</sup> |
|----------------------------------------------------------------------------------------------|-------------------------|-----------------------|---------------------------------------------------------|----------------------------|-------------------|-------------------------|
| <i>Aequorea</i> GFP <sup>c</sup>                                                             | 395                     | 508                   | 27,600                                                  | 0.79                       | -                 | 100                     |
| EGFP <sup>c</sup>                                                                            | 489                     | 508                   | 55,000                                                  | 0.60                       | -                 | 151                     |
| <br>DMHBI   | 394                     | 487                   | 23,336                                                  | 0.0005                     | -                 | 0.05                    |
| 2-4                                                                                          | 397                     | 464                   | 21,536                                                  | 0.10                       | N.D. <sup>d</sup> | 6                       |
| 13-2                                                                                         | 398                     | 529                   | 23,391                                                  | 0.06                       | 464 nM            | 12                      |
| 13-2-min                                                                                     | 398                     | 529                   | 23,391                                                  | 0.11                       | N.D.              | 12                      |
| 3-6                                                                                          | 398                     | 537                   | 25,300                                                  | 0.05                       | 406 nM            | 6                       |
| 17-3                                                                                         | 405                     | 547                   | 18,127                                                  | 0.09                       | N.D.              | 7                       |
| <br>DMABI   | 447                     | 519                   | 14,329                                                  | 0.003                      | -                 | 0.2                     |
| 11-3                                                                                         | 489                     | 512                   | 18,743                                                  | 0.05                       | N.D.              | 4                       |
| <br>2-HBI  | 399                     | 461                   | 10,943                                                  | 0.0005                     | -                 | 0.02                    |
| 6-8                                                                                          | 396                     | 588                   | 10,714                                                  | 0.004                      | N.D.              | 0.2                     |
| <br>DFHBI | 405                     | 498                   | 11,864                                                  | 0.0007                     | -                 | 0.04                    |
| 24-2 (Spinach)                                                                               | 469                     | 501                   | 24,271                                                  | 0.72                       | 537 nM            | 80                      |

<sup>a</sup>Extinction coefficients for fluorophores alone were measured at a pH in which all species were in the phenolic form except for DFHBI which was measured in its phenolate form. RNA-fluorophore complex extinction coefficients were all measured at pH 7.4. <sup>b</sup>Brightness (extinction coefficient  $\times$  quantum yield) is reported relative to *Aequorea* GFP. <sup>c</sup>See reference 40. <sup>d</sup>N.D. = not determined.

E

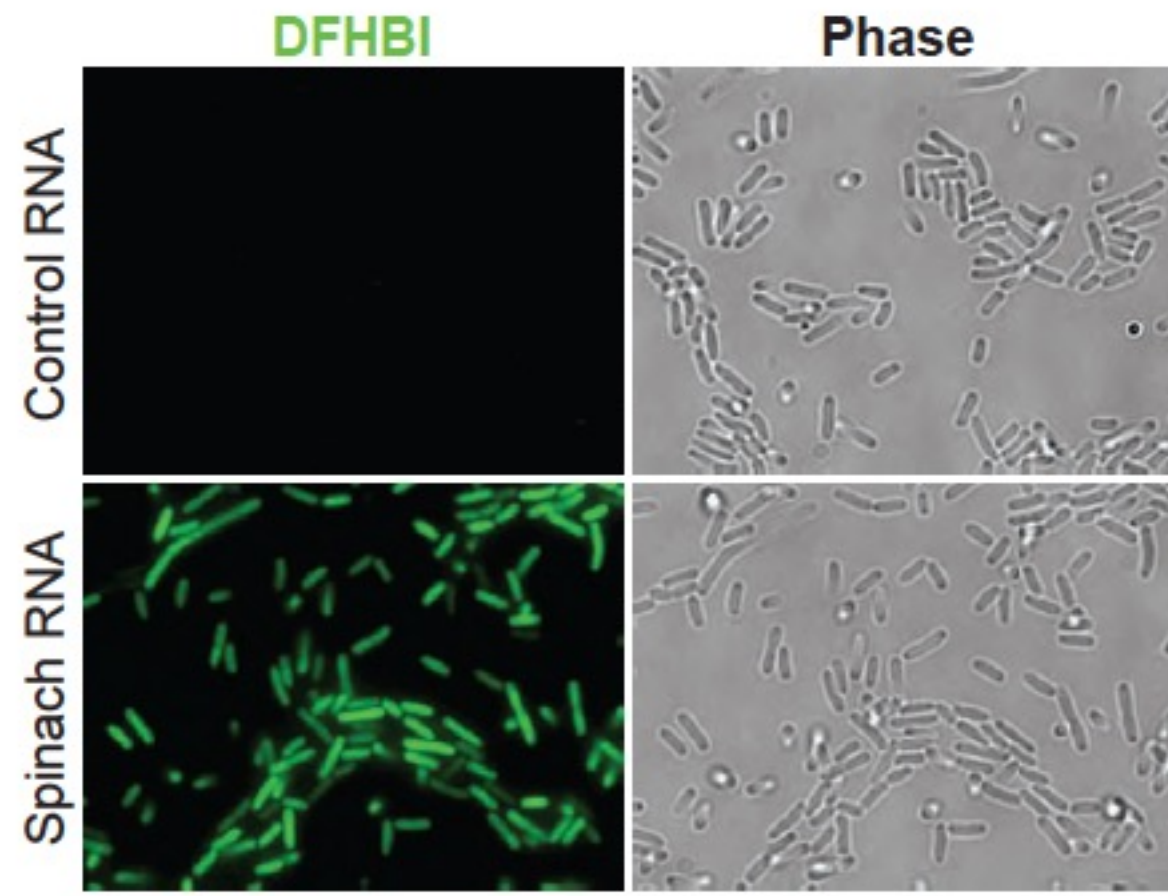




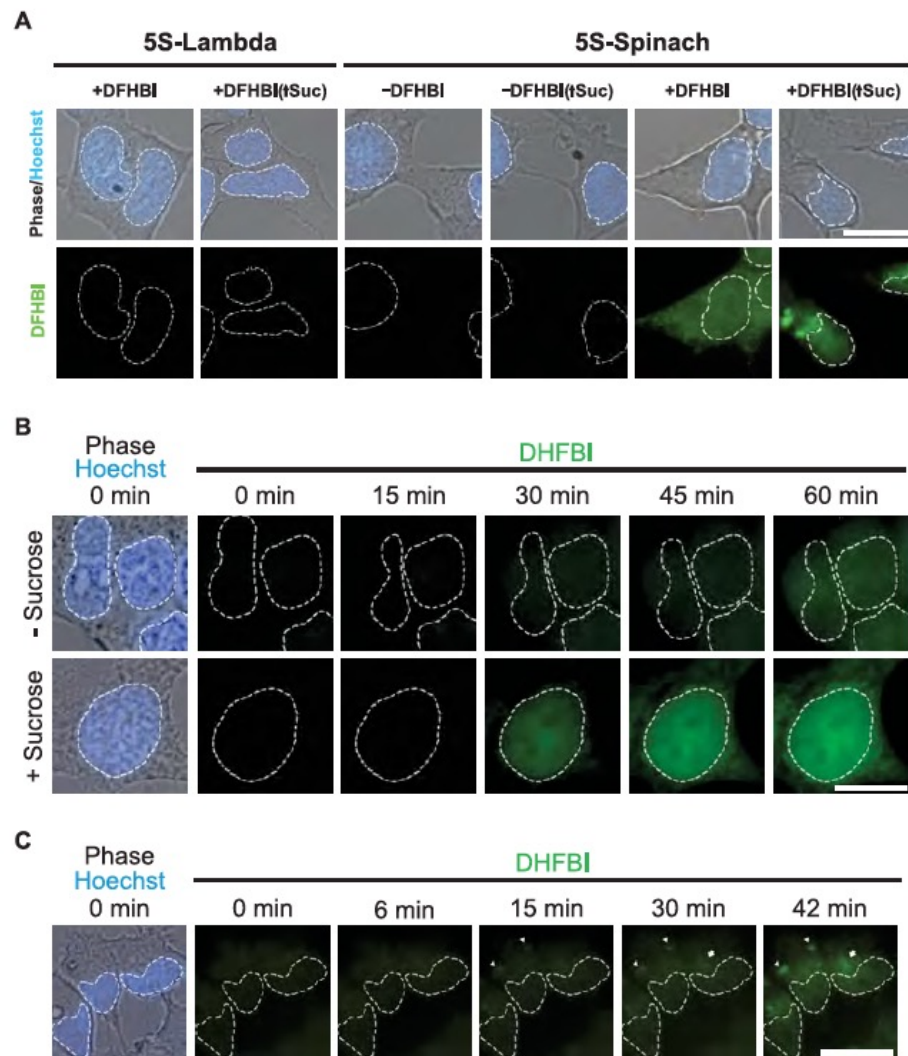


**Fig. 3.** RNA-fluorophore complexes with EGFP-like properties. **(A)** Normalized excitation (blue) and emission (green) spectra of 24-2-DFHBI complex. **(B)** Normalized absorbance spectra of DFHBI at pHs 5.0, 6.0, 7.0, and 8.0. **(C and D)** 24-2 incubated with excess DFHBI (C) or HBI (D) at pHs 6.0, 7.0, and 8.0. **(E)** Photobleaching curves for 24-2-DFHBI, EGFP, and fluorescein. Fluorophores were immobilized on glass slides and illuminated continuously with a 130 W mercury lamp. Total fluorescence was then plotted against exposure time and normalized to the maximum intensity of each fluorophore.



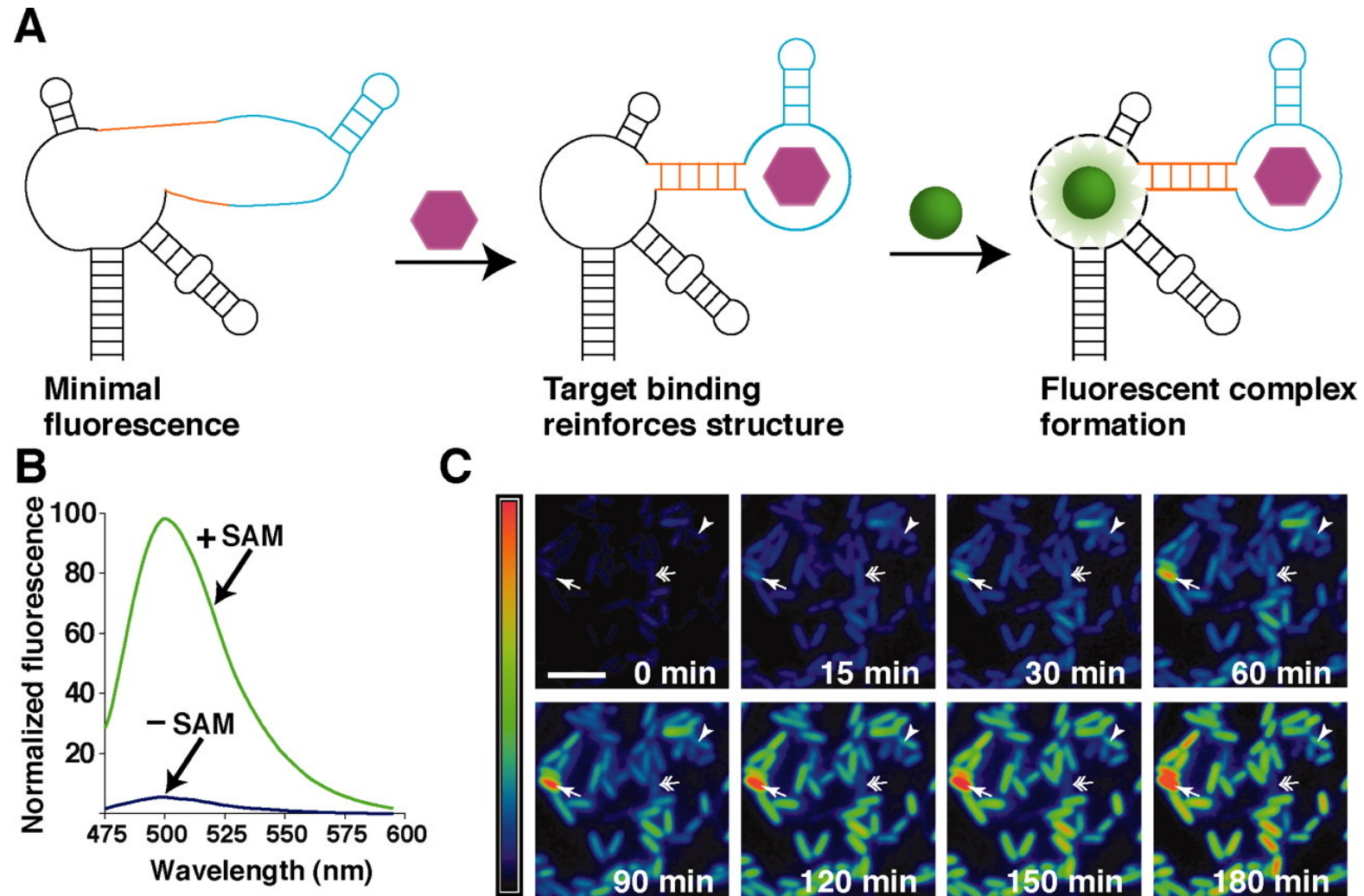






**Fig. 4.** Live-cell imaging of Spinach fusion RNAs. **(A)** Live-cell imaging of Spinach-tagged 5S RNA. Fluorescence and phase images of HEK293 T cells expressing 5S tagged with either Spinach or Lambda, a control RNA. Fluorescence is detected in 5S-Spinach-expressing cells in the presence of 20  $\mu$ M DFHBI, with granule formation present in cells treated with 600 mM sucrose for 30 min ( $\uparrow$ Suc). White dashed lines indicate nuclear borders assessed by means of Hoescht 33342 staining. **(B)** 5S-Spinach RNA induction in response to stress. 5S-Spinach-expressing HEK293 T cells were pretreated with 30 nM ML-60128 for 16 hours and then treated with vehicle or 600 mM sucrose for 60 min. Treatment of cells with sucrose resulted in a rapid induction of 5S-Spinach RNA and an increase in total 5S-Spinach levels compared with control cells. **(C)** 5S-Spinach RNA localization into granules. 5S-Spinach-expressing HEK293 T cells were stimulated with 600 mM sucrose in order to monitor the rate of formation of 5S-Spinach-containing granules. Arrowheads indicate granules that formed earliest, and arrows indicate granules that developed later during the time course of treatment. Scale bar, 10  $\mu$ m.

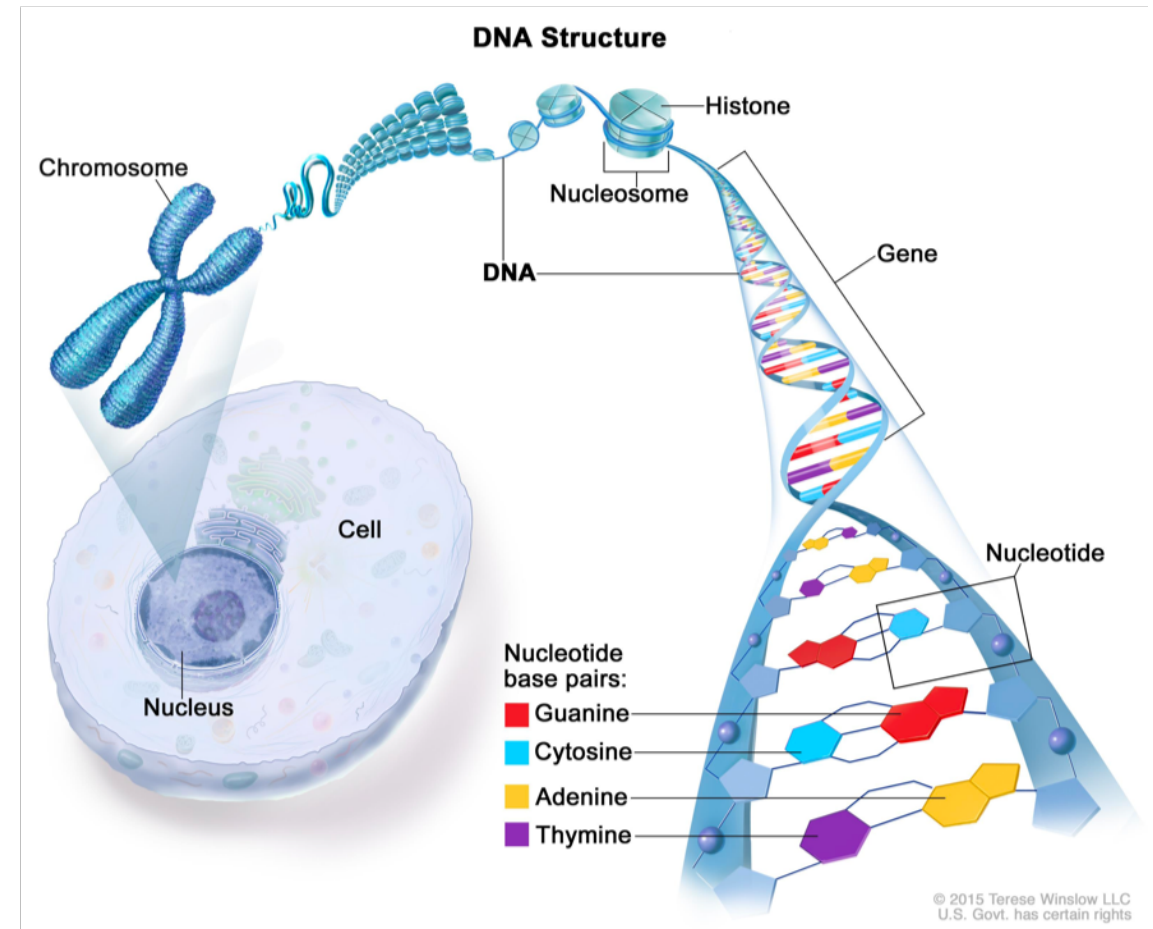
**Fig. 1 Imaging SAM in living cells with RNA. (A) The sensor RNA comprises Spinach (black), a transducer (orange), and a target-binding aptamer (blue).**



Jeremy S. Paige et al. Science 2012;335:1194

## Goal of DNA imaging:

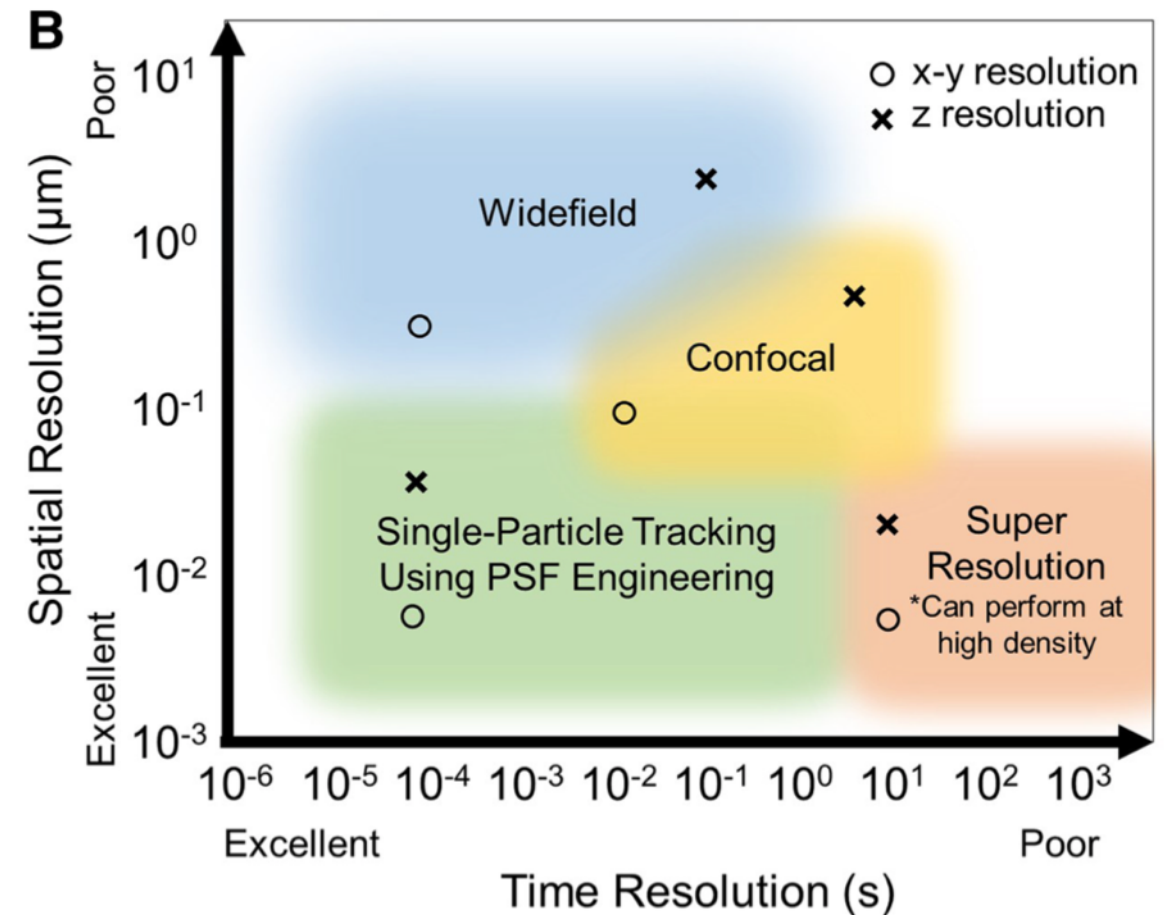
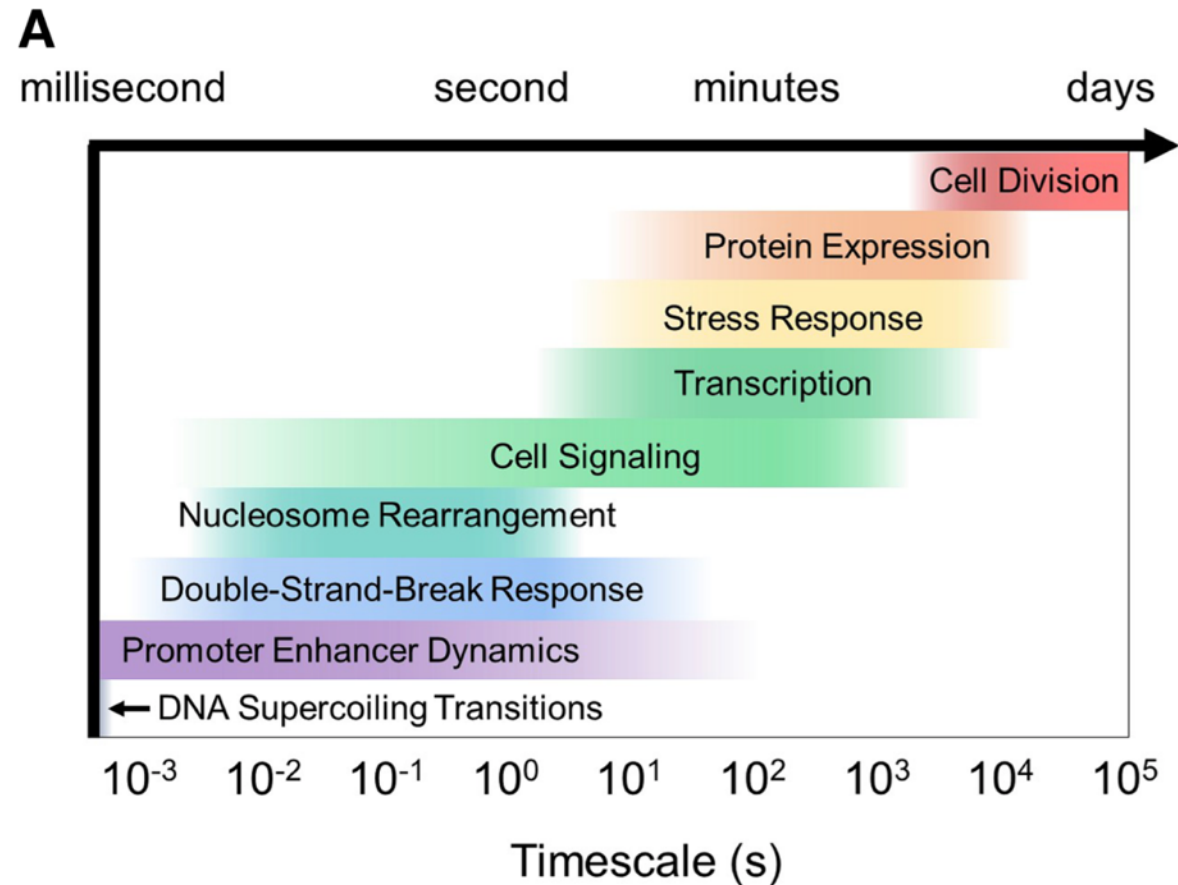
Time-resolved, 3D positions of the entire genome, with information on sequence and epigenetic modification.



<https://www.cancer.gov/about-nci/organization/ccg/research/structural-genomics/tcga>, 29/5/2020

1. Why should we study DNA in cells?
2. DNA labeling methods:
  - a) DNA dye staining
  - b) Unnatural nucleosides and pro-nucleotides
  - c) CASFISH – a CRISPR-dCas9 method
3. Paper
  - Rieder et al. – Alkene-Tetrazine Ligation for Imaging Cellular DNA

# 1. DNA is involved in many cellular processes that occur on a time scale that can be observed with common imaging techniques





# 1. Questions that could be addressed with DNA imaging

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- To what extent does the structuring of chromatin in topological regions (e.g., chromatin domains) contribute to nuclear functioning? On what length scales does this structuring occur?
- How is it dynamically organized e.g., during the cell cycle or during DNA replication or transcription?
- How are the dynamical processes themselves responsible for the possibly induced structural changes of chromatin?

Genes **2019**, *10*, 493.

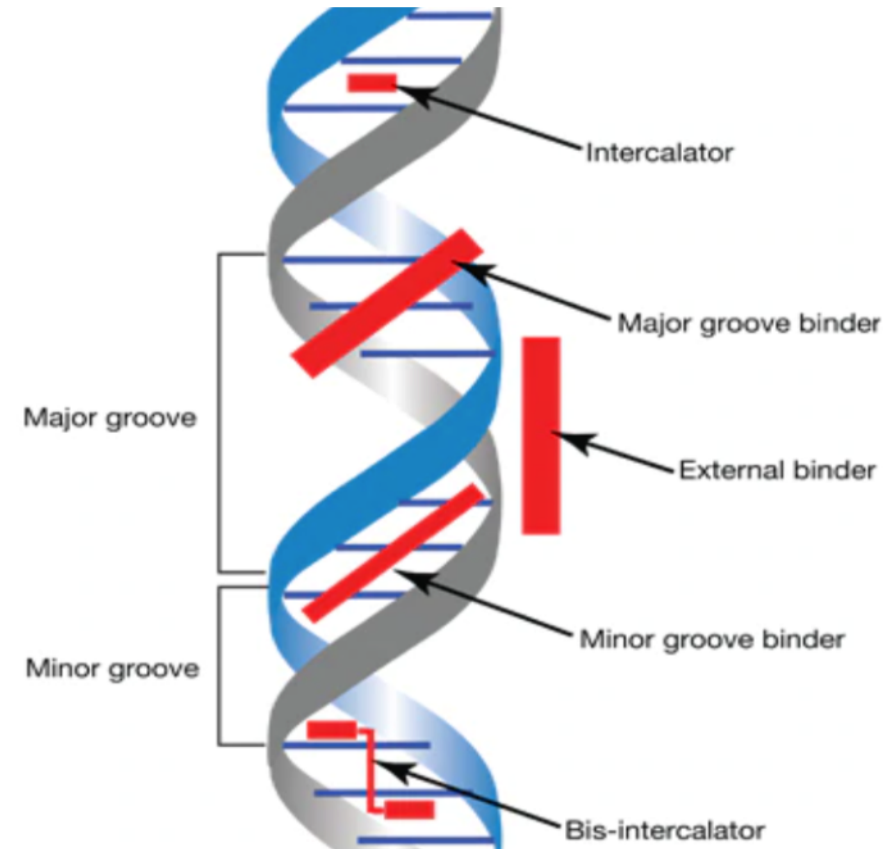


# 1. Key applications of DNA imaging

| Application                                     | Labeling method                                     |
|-------------------------------------------------|-----------------------------------------------------|
| Cell viability test                             | Cell impermeable DNA stain                          |
| Imaging chromatin                               | Cell permeable DNA stain                            |
| Cell cycle analysis, DNA replication and repair | DNA stain, Unnatural nucleosides or pro-nucleotides |
| Tracking repetitive loci                        | DNA binding proteins (Zinc finger proteins)         |
| Tracking specific, nonrepetitive loci           | DNA binding proteins (CRISPR-dCas9 method)          |

## 2. DNA labeling methods: a) DNA dye staining

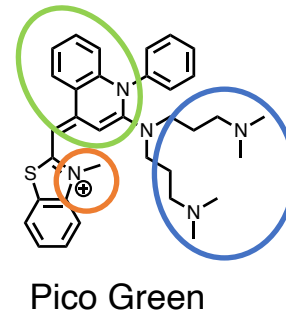
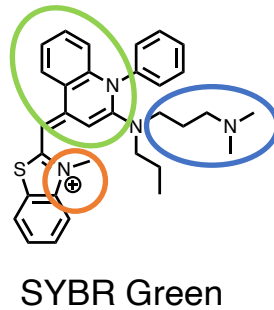
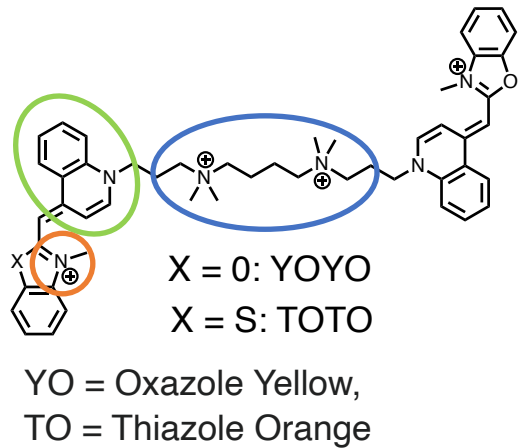
- Intercalating between base pairs
  - Cyanine monomers and dimers (Mono- and bisintercalator)
  - Phenanthridines and acridines
- Minor groove binder
  - Indoles and imidazoles



<https://www.thermofisher.com/de/de/home/references/molecular-probes-the-handbook/nucleic-acid-detection-and-genomics-technology/nucleic-acid-stains.html>, 29/5/20

## 2. DNA labeling methods: a) DNA dye staining

### Cyanine Dyes:



- Intercalator
- Interaction with phosphate backbone
- Interaction with DNA minor groove

### Properties:

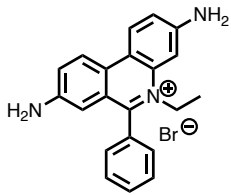
- High turn-on upon DNA binding
- Mostly cell impermeable

### Applications:

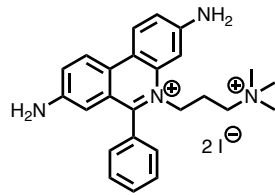
- Ultrasensitive nucleic acid detection and quantification in solution or gels
- Dead-live stain to determine cell cytotoxicity

## 2. DNA labeling methods: a) DNA dye staining

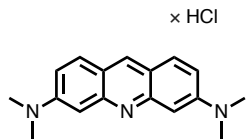
### Classical Intercalators: phenanthrines and acridines



Ethidium bromide (EtBr)



Propidium iodide (PI)



Acridine orange

#### *Properties:*

- EtBr and PI are potent mutagens.
- Acridine orange has an emission maximum at 525 nm when bound to DNA that is shifted to 650 nm when bound to RNA.

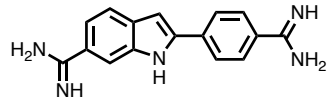
#### *Applications:*

- PI is commonly used as chromosome and nuclear counterstain *e. g.* in flow cytometry.
- Acridine Orange is a dual-fluorescence nucleic acid stain that can be used for RNA/DNA discrimination measurements.

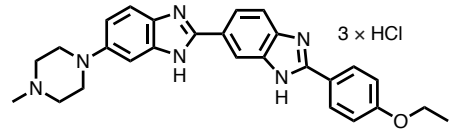


## 2. DNA labeling methods: a) DNA dye staining

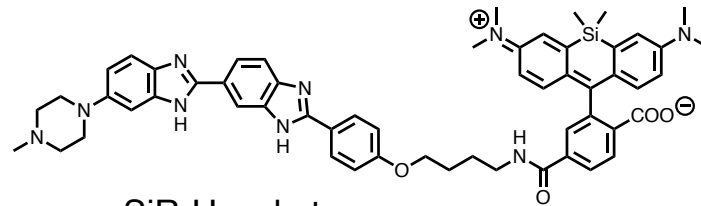
### Minor groove binders: Indoles and Imidazoles



DAPI



Hoechst33342



SiR-Hoechst

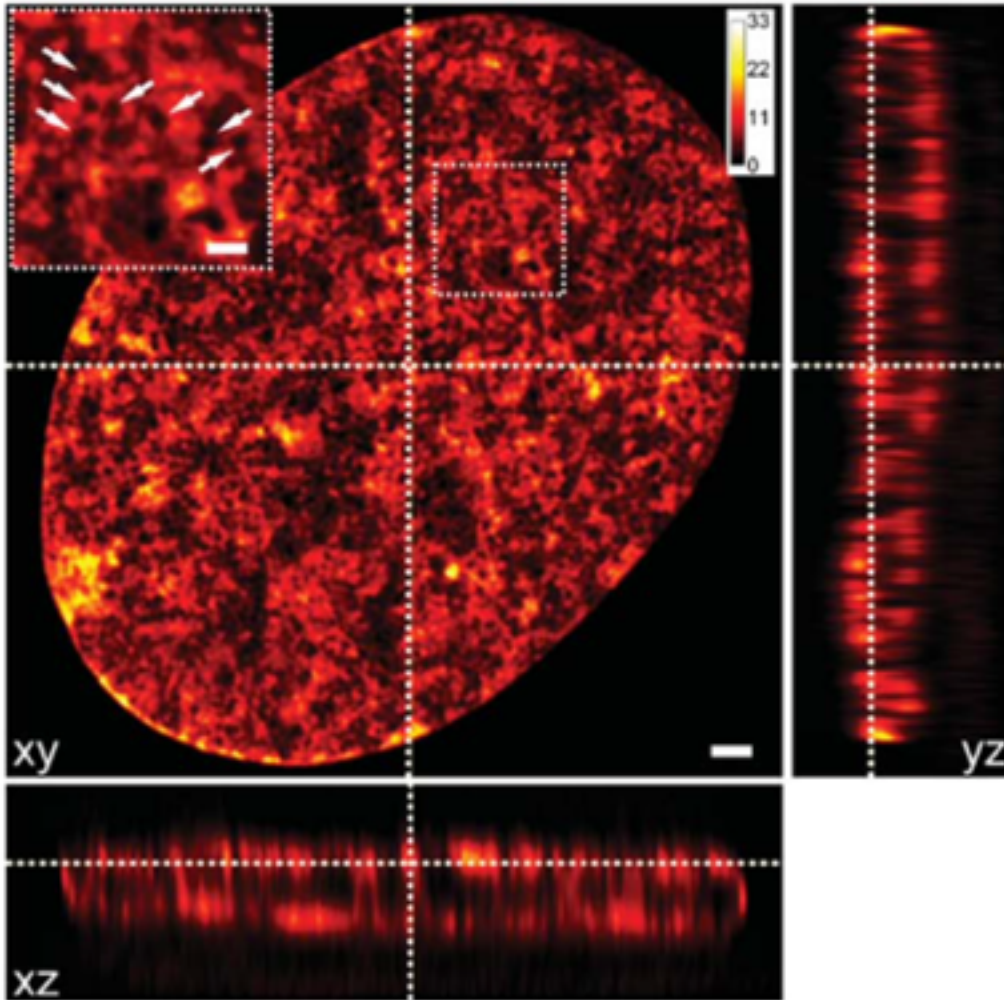
### *Properties:*

- AT-selective.
- DAPI is cell-impermeable whereas Hoechst33342 and SiR-Hoechst are cell permeable.

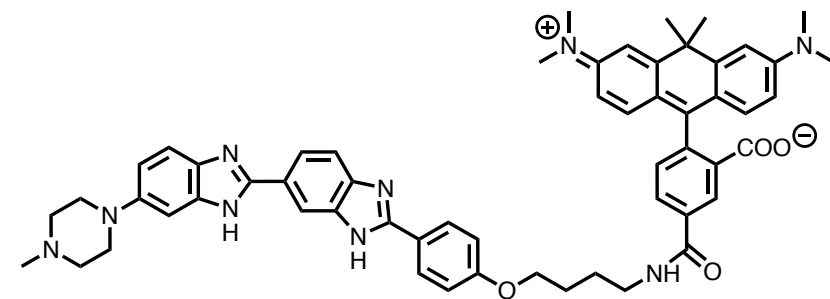
### *Applications:*

- Nuclear counterstains for fluorescence microscopy.
- Cell cycle analysis for flow cytometry or fluorescence microscopy.

## 2. DNA labeling methods: a) DNA dye staining can give super-resolved images of chromatin



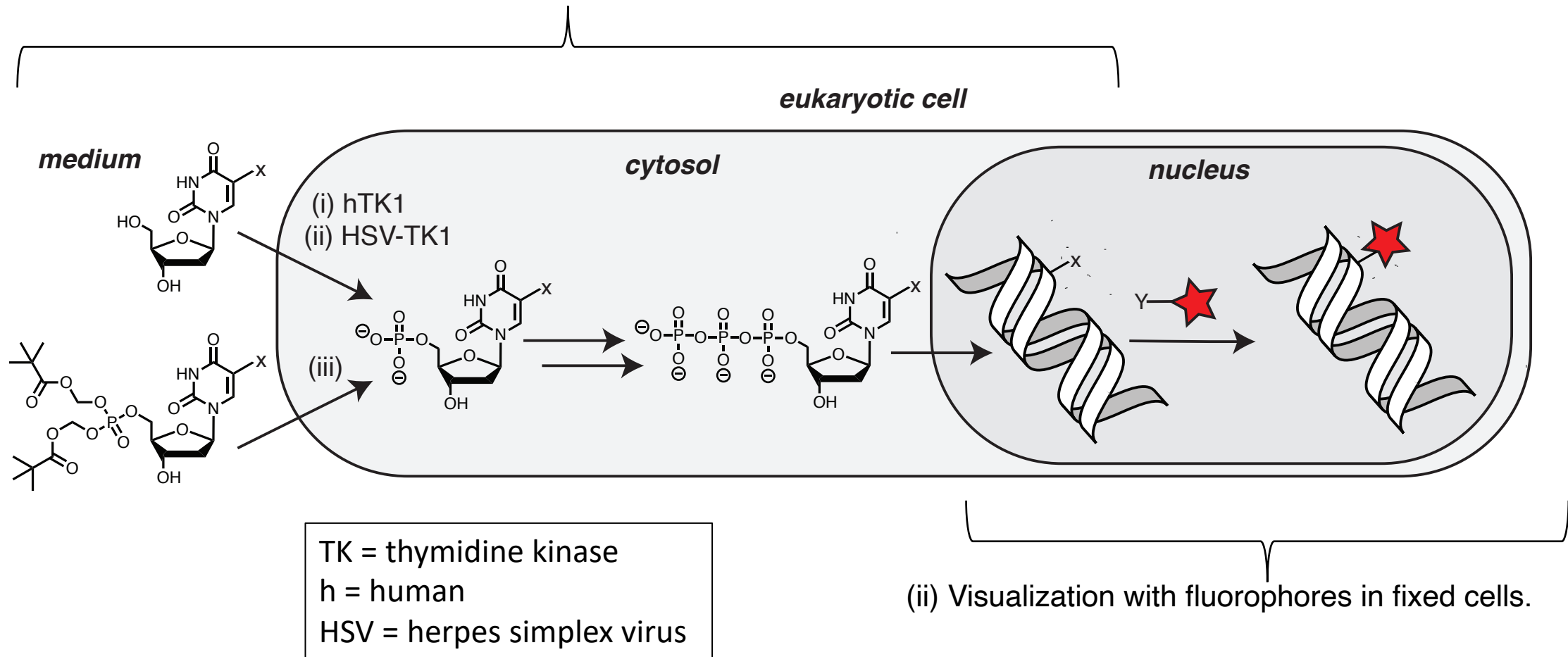
STED nanoscopy images of nuclei in living fibroblast stained 5-610CP-Hoechst. Arrows indicate heterochromatin exclusion zones (HEZs). Scale bar: 1 mm in the large image and 0.5 mm in the inset.



Chem. Sci. **2019**, *10*, 1962–1970.

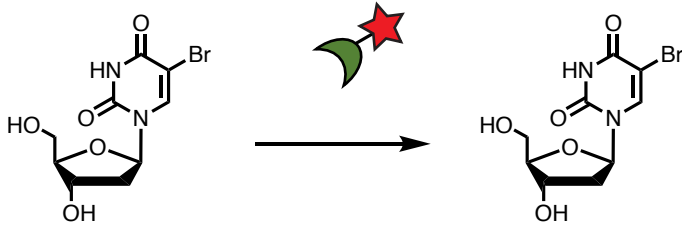
## 2. DNA labeling methods: b) Labeling newly synthesized DNA

(i) Metabolic processing of nucleoside or pro-nucleotide analogues in living cells.



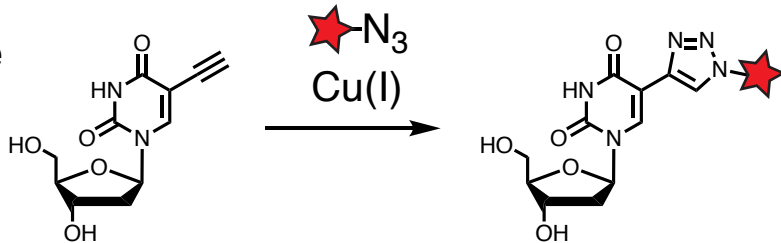
## 2. DNA labeling methods: b) Labeling newly synthesized DNA: Unnatural nucleosides

**5-Bromo-  
2'-deoxyuridine  
(BrdU)**



Immunostaining with antibody

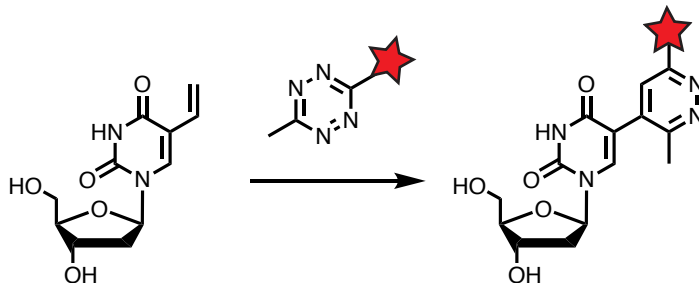
**5-Ethynyl-  
2'-deoxyuridine  
(EdU)**



Copper-catalyzed azide-alkyne cycloaddition (CuAAC)

$$k \sim 10 - 200 \text{ M}^{-1} \text{ s}^{-1}$$

**5-Vinyl-  
2'-deoxyuridine  
(VdU)**



Inverse electron-demand Diels Alder reaction (iEDDA)

$$k_{\text{VdU}} \sim 0.021 \text{ M}^{-1} \text{ s}^{-1}, \text{ (other dienophiles: } k_{\text{tetrazines}} \sim 1 - 10^4 \text{ M}^{-1} \text{ s}^{-1})$$

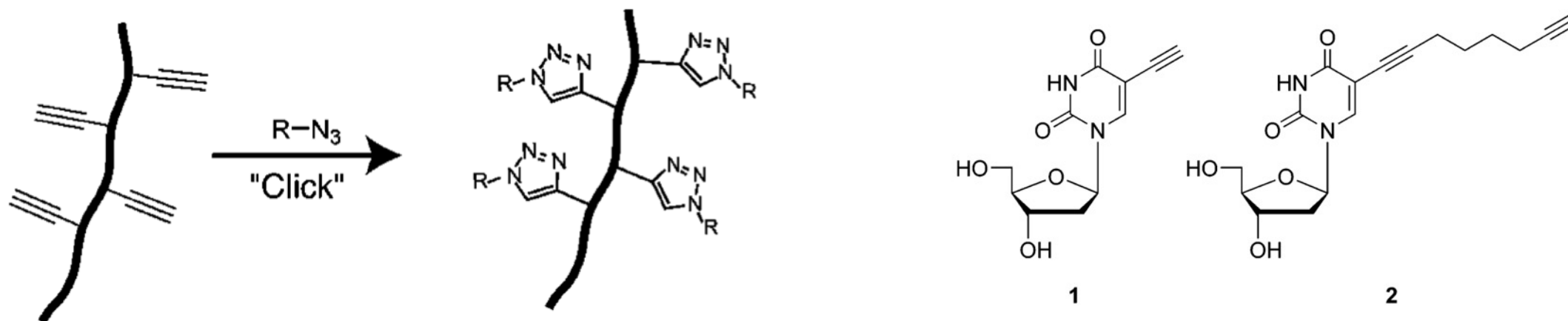
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Angew Chem Int Ed Engl **2014**, 53, 9168-72.

ACS Chem. Biol. **2014**, 9, 16-20.

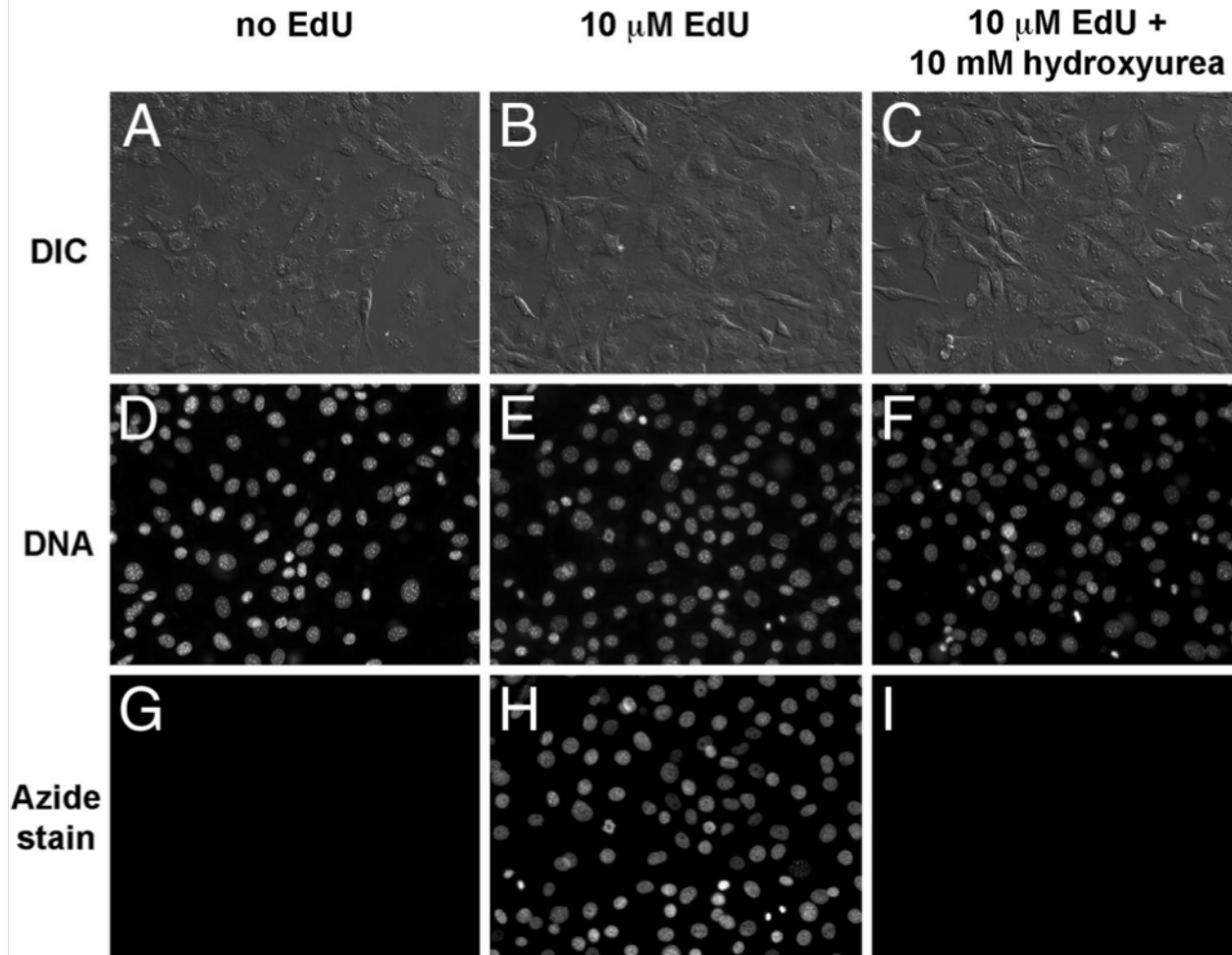
## 2. DNA labeling methods: b) Labeling newly synthesized DNA: EdU



- Postsynthetic modification of oligonucleotides using Cu(I) catalyzed azide-alkyne cycloaddition was first demonstrated by Carell in 2006.
- Applications in mammalian cells were published two years later in 2008.

Org. Lett. 2006, 8, 17, 3639-42.

## 2. DNA labeling methods: b) Labeling newly synthesized DNA: EdU

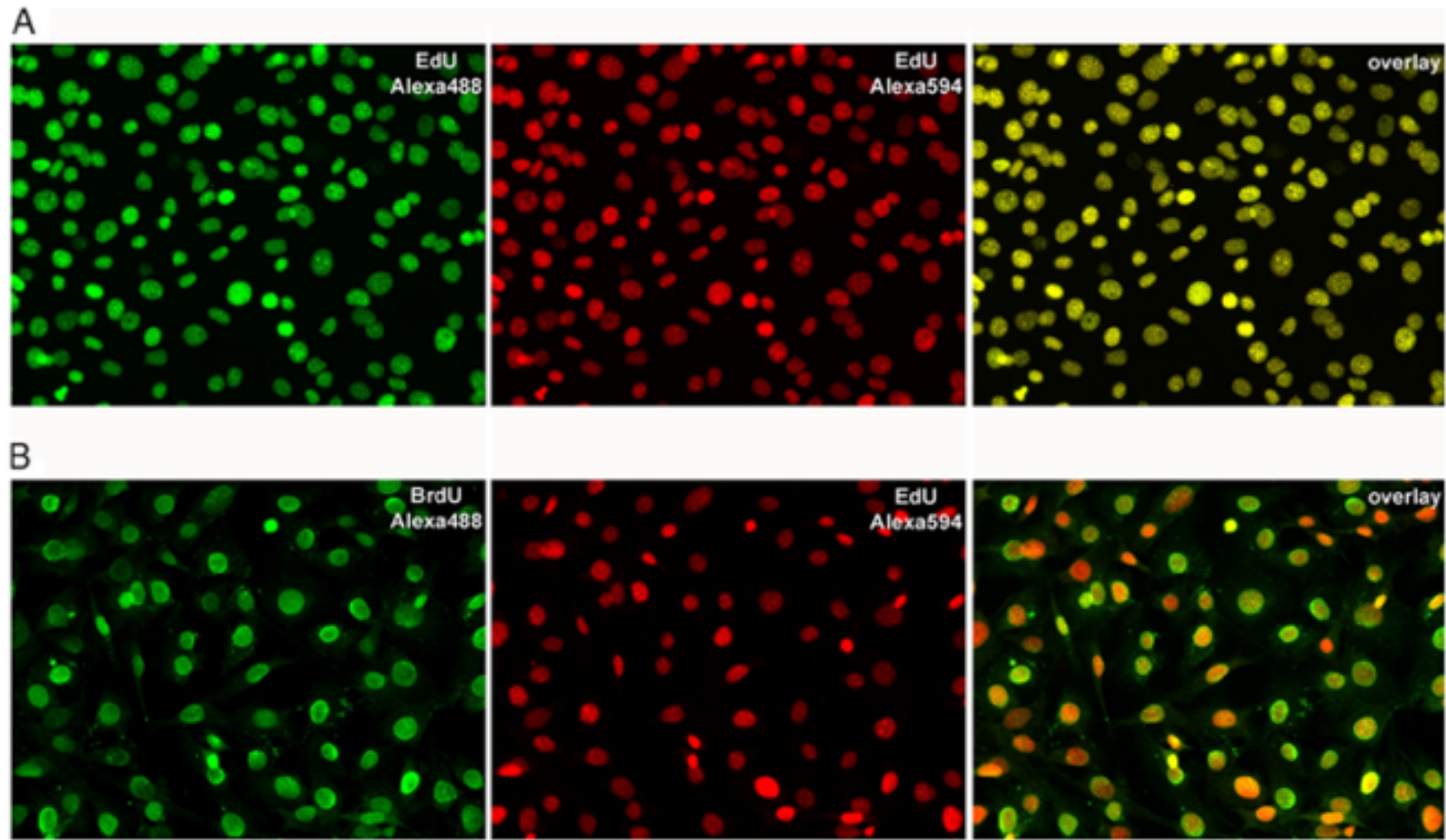


**Fig. 2.** Detection of EdU incorporated into the DNA of cultured NIH 3T3 cells (A–M) and HeLa cells (N–P) by fluorescence microscopy. NIH 3T3 cells were incubated in media without EdU (A, D, G, J, and L), media supplemented with 10  $\mu$ M EdU (B, E, and H) or media with 10  $\mu$ M EdU and 10 mM hydroxyurea to block DNA synthesis (C, F, I, K, and M). In A–M, the cells were fixed and then reacted with 10  $\mu$ M Alexa568-azide (Fig. 1C, compound 2) for 10 min. The cells were then counterstained with Hoechst to reveal cellular DNA, washed, and imaged by fluorescence microscopy and differential interference contrast (DIC). Note the strong specific and low nonspecific azide stain in the presence (H) and absence (G) of EdU, respectively. Not all nuclei in H are labeled after overnight incubation with EdU, suggesting that only cells that went through S phase became labeled. Blocking DNA replication with hydroxyurea abolishes EdU incorporation almost completely (I).

- EdU incorporation can be diminished with the DNA synthesis inhibitor hydroxyurea.



## 2. DNA labeling methods: b) Labeling newly synthesized DNA: EdU

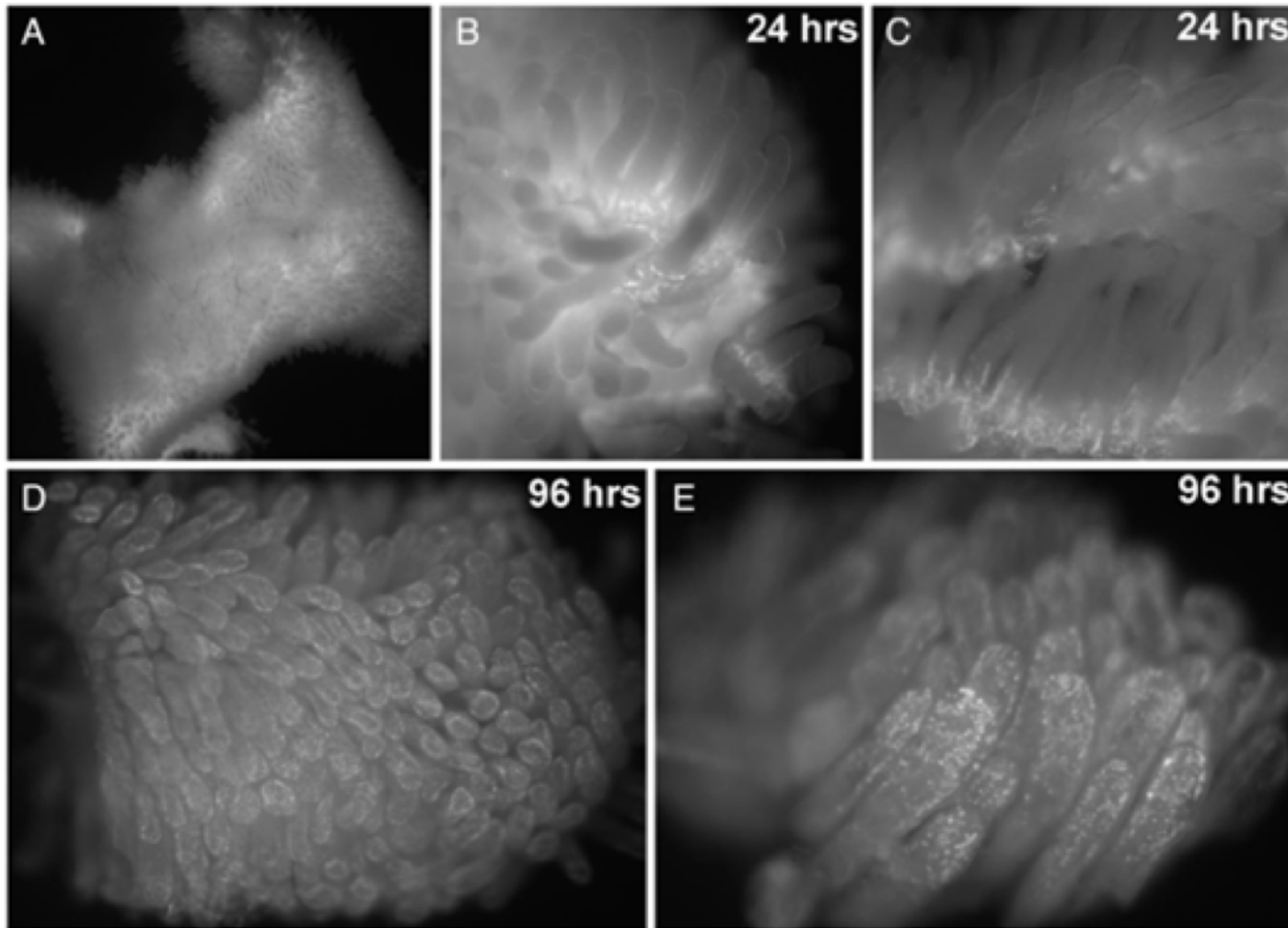


- EdU signals colocalize with BrdU signals.

**Fig. 3.** Reproducibility of EdU labeling (A) and comparison with BrdU (B)

PNAS 2008, 105, 2415-20.

## 2. DNA labeling methods: b) Labeling newly synthesized DNA: EdU

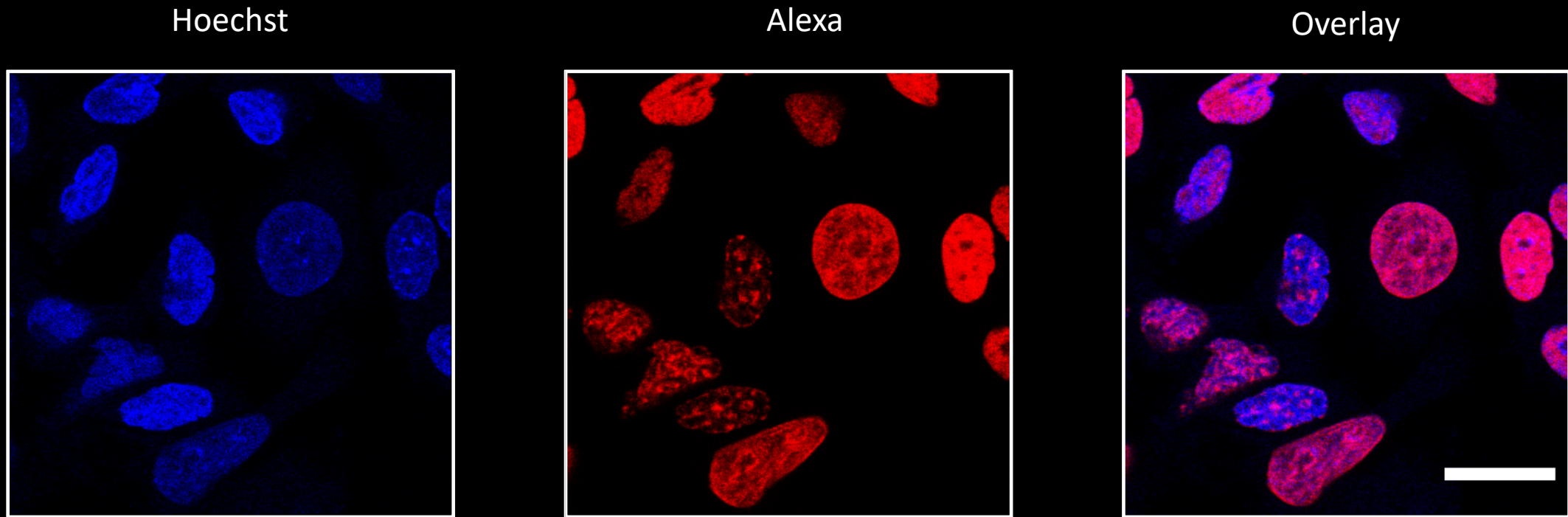


- Whole-mount fluorescent images of mouse small intestine, stained to detect EdU incorporation.

**Fig. 5.** Exploring cell proliferation and tissue dynamics in animals using EdU.

PNAS 2008, 105, 2415-20.

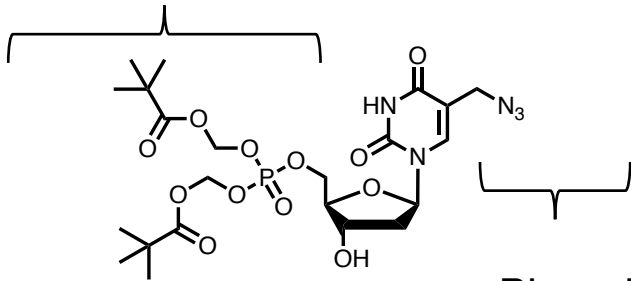
## 2. DNA labeling methods: b) Labeling newly synthesized DNA: EdU



Confocal images of HeLa cells treated with EdU (10  $\mu$ M, 16 h) before PFA fixation and labeling with Alexa647-azide and Hoechst33342, scale 25  $\mu$ m.

## 2. DNA labeling methods: b) Labeling newly synthesized DNA: Unnatural pro-nucleotides

Pivaloyloxymethyl  
phosphotriester (=POM) is  
cleaved by endogenous  
esterases



Bio-orthogonal handle for strain-promoted  
azide-alkyne cycloaddition (SPAAC)

$$k \sim 0.01 - 1 \text{ M}^{-1} \text{ s}^{-1}$$

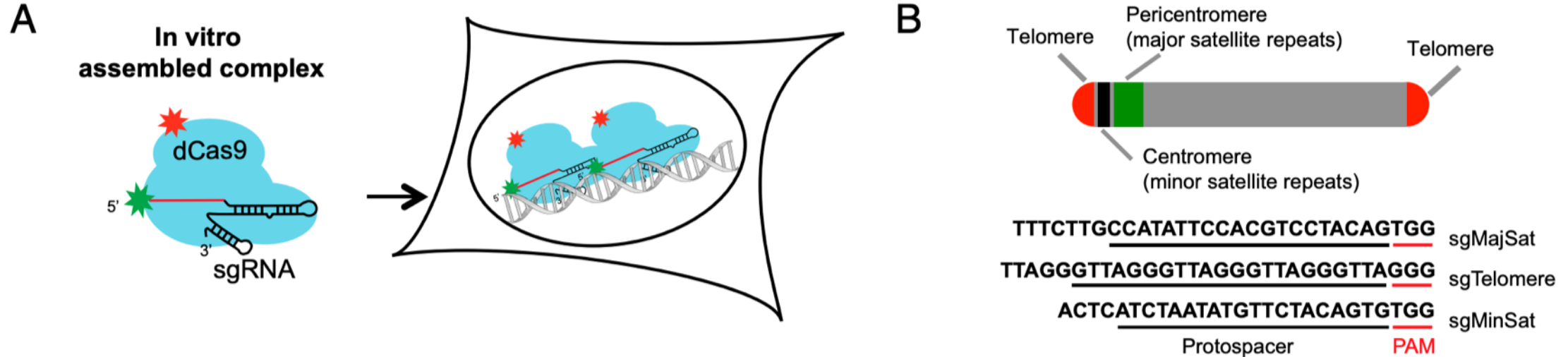
→ The larger the handle, the less likely to  
be a substrate of endogenous enzymes.

More efficient incorporation into DNA through  
protected monophosphate.

→ First phosphorylation is critical for incorporation  
into DNA.

## 2. DNA labeling methods: c) CASFISH – a CRISPR-dCas9 method for fixed cells

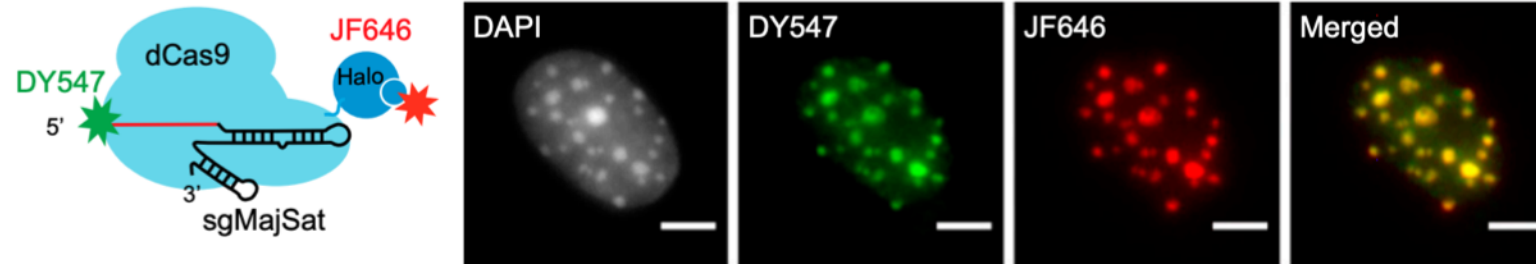
- *In vitro* assembled dCas9/sgRNA fluorescently labels genomic DNA in cells
  - System is derived from Fluorescence-in situ hybridization (FISH), but no DNA denaturation needed.
    - (i) CRISPR (clustered regularly interspaced short palindromic repeats) -dCas9 (nuclease-deficient caspase 9) mutant fused HaloTag labelled with JF646 (A)
    - (ii) sgRNA that is targeting the desired sequences in the genome (B)



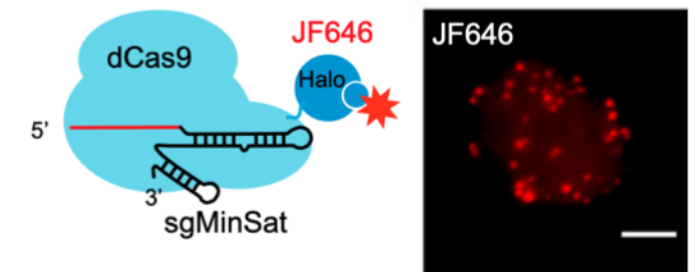


## 2. DNA labeling methods: c) CASFISH – a CRISPR-dCas9 method for fixed cells

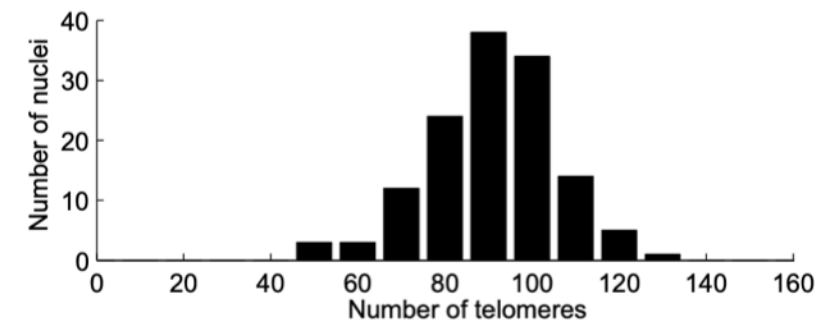
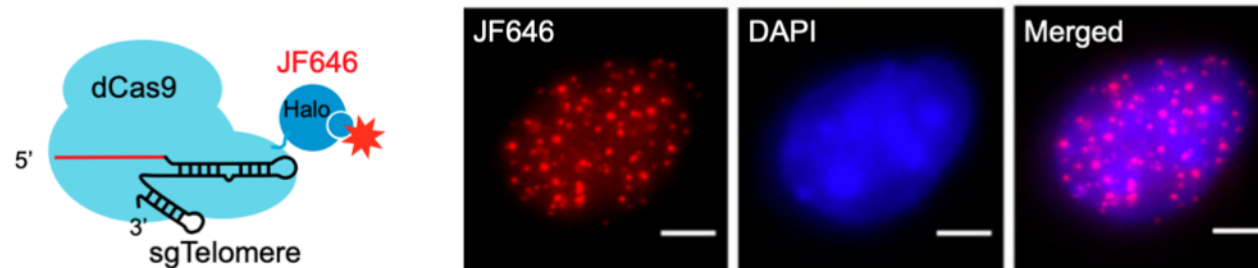
### C CASFISH of pericentromeres



### E CASFISH of centromeres



### D CASFISH of telomeres



PNAS 2015, 112, 38, 11870-1187.



### 3. Paper: a) VdU: Rieder *et al.* - Alkene-Tetrazine Ligation for Imaging Cellular DNA

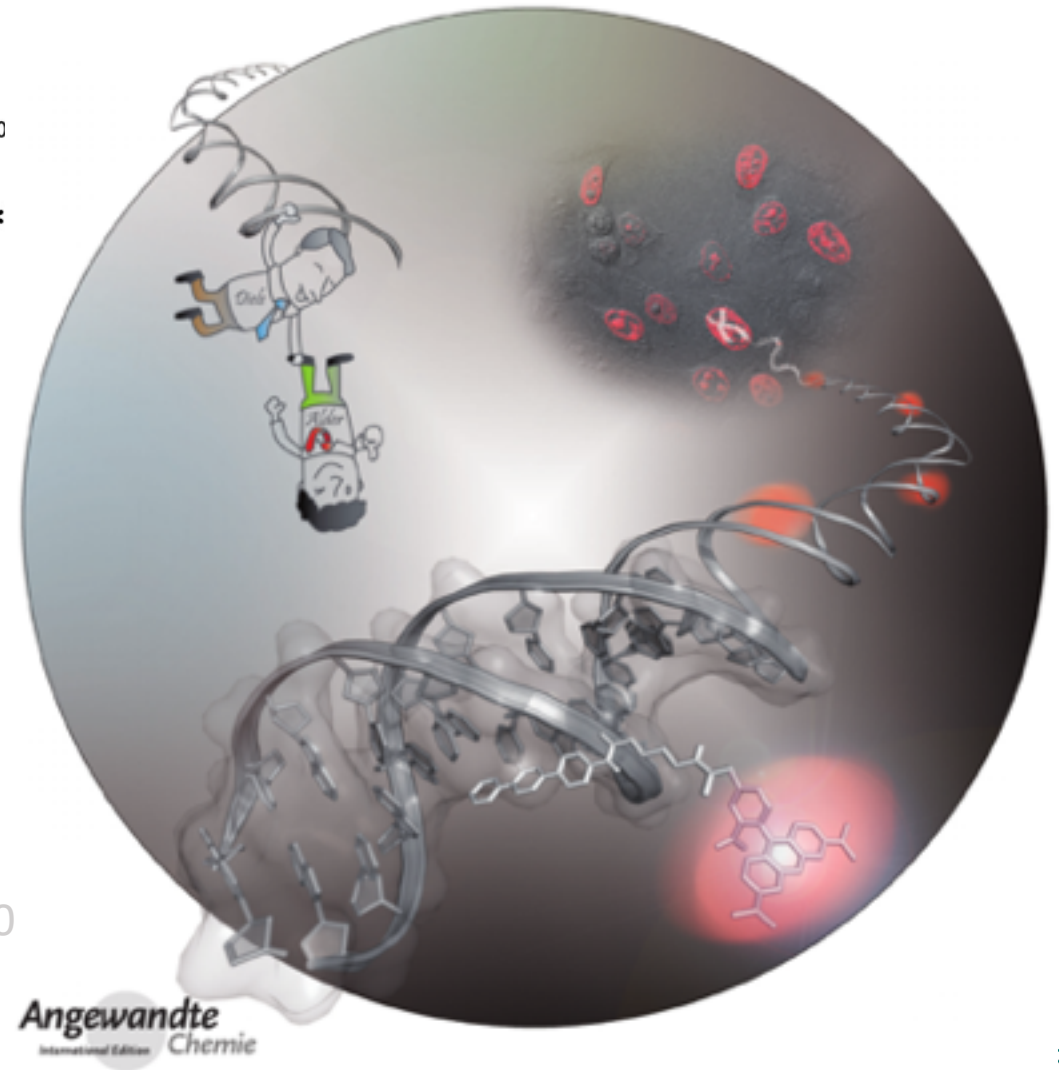
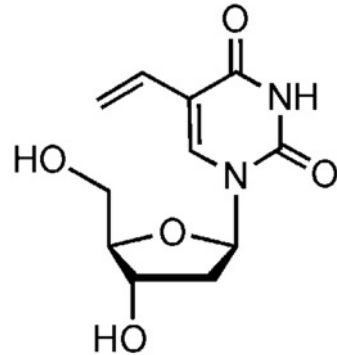
Angewandte  
Communications

VIP DNA Labeling Very Important Paper

DOI: 10.1002/anie.201403580

#### Alkene–Tetrazine Ligation for Imaging Cellular DNA\*\*

Ulrike Rieder and Nathan W. Luedtke\*

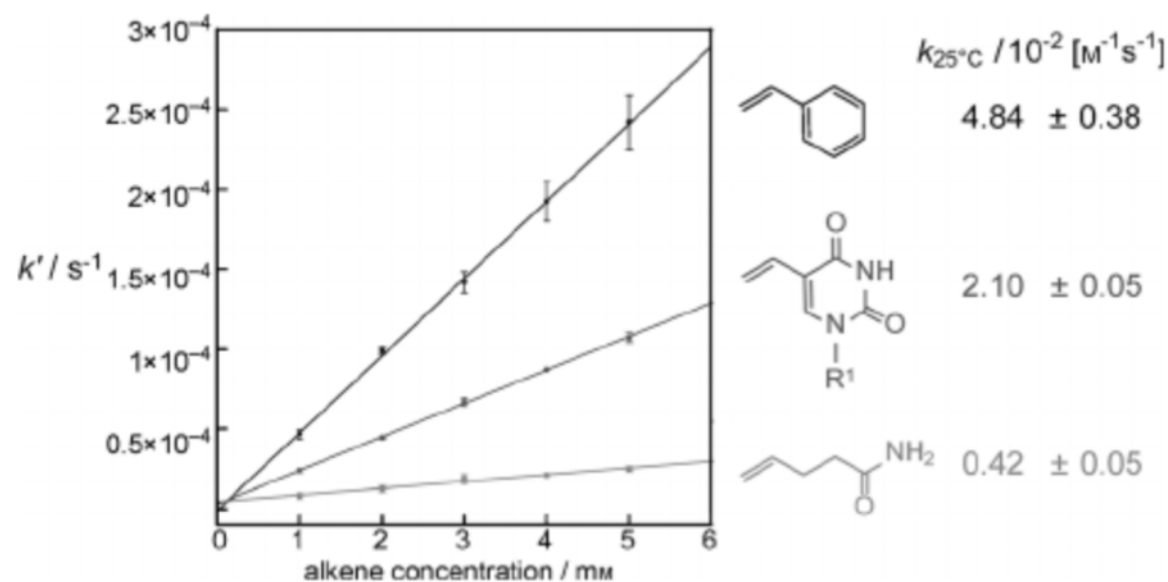


<https://onlinelibrary.wiley.com/doi/epdf/10.1002/anie.201403580>

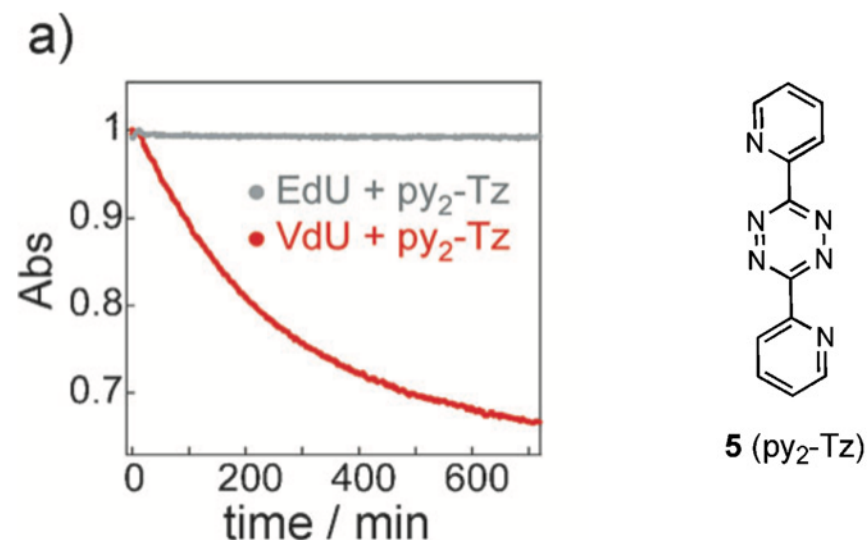
Angewandte  
International Edition  
Chemie



### 3. Paper: a) VdU reacts with pyridyl-tetrazine but not EdU



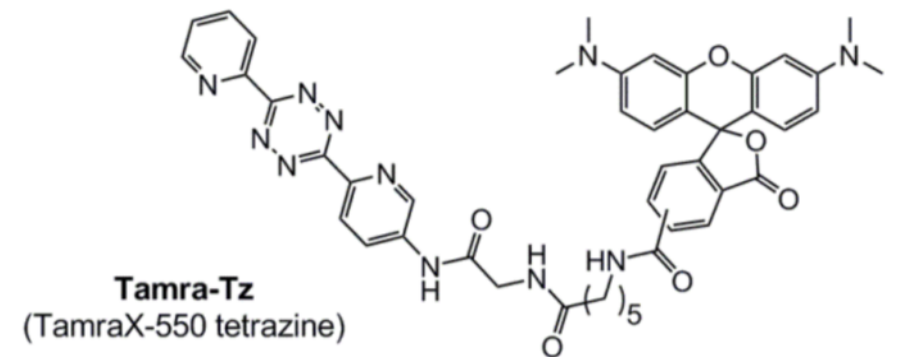
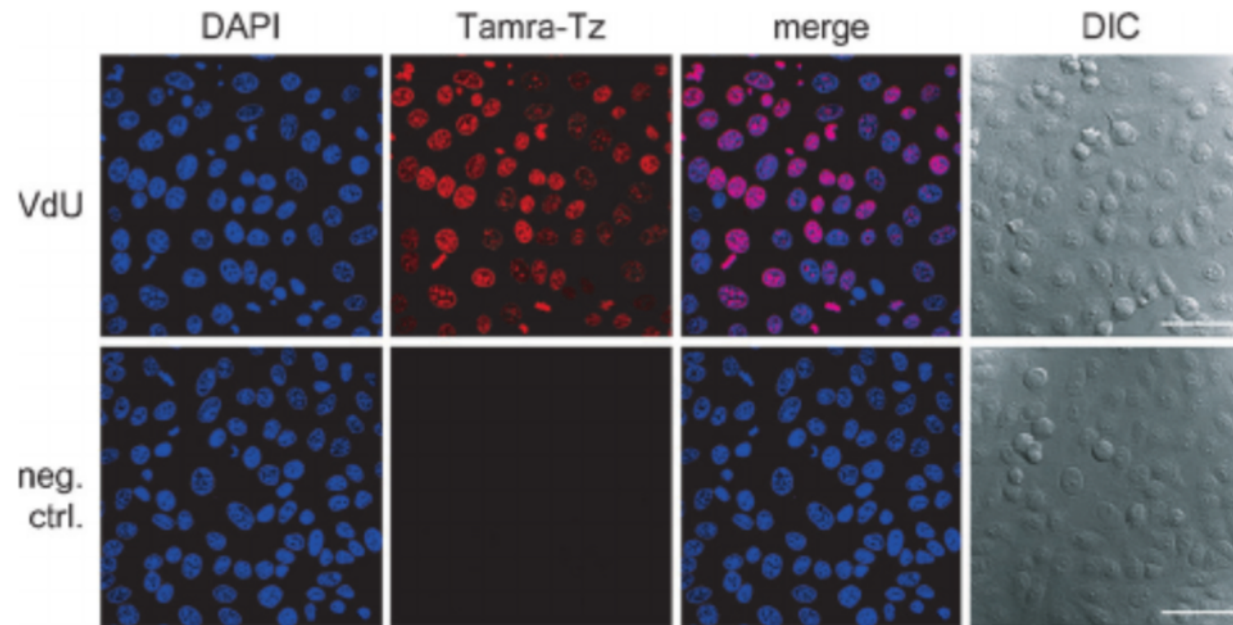
**Figure 1.** Pseudo-first-order reaction rates ( $k'$ ) versus alkene concentration for the consumption of tetrazine **5** (0.1 mM) in a 1:1 (v/v) mixture of methanol and water at 25 °C. The slope of each linear regression provides the rate constant  $k$ .  $\text{R}^1 = 2'$ -deoxyribose.



**Figure 4.** a) The consumption of tetrazine **5** (0.1 mM) according to normalized absorbance changes at 530 nm in the presence of 5 mM of EdU or VdU in a 1:1 (v/v) mixture of methanol and water at 25 °C.

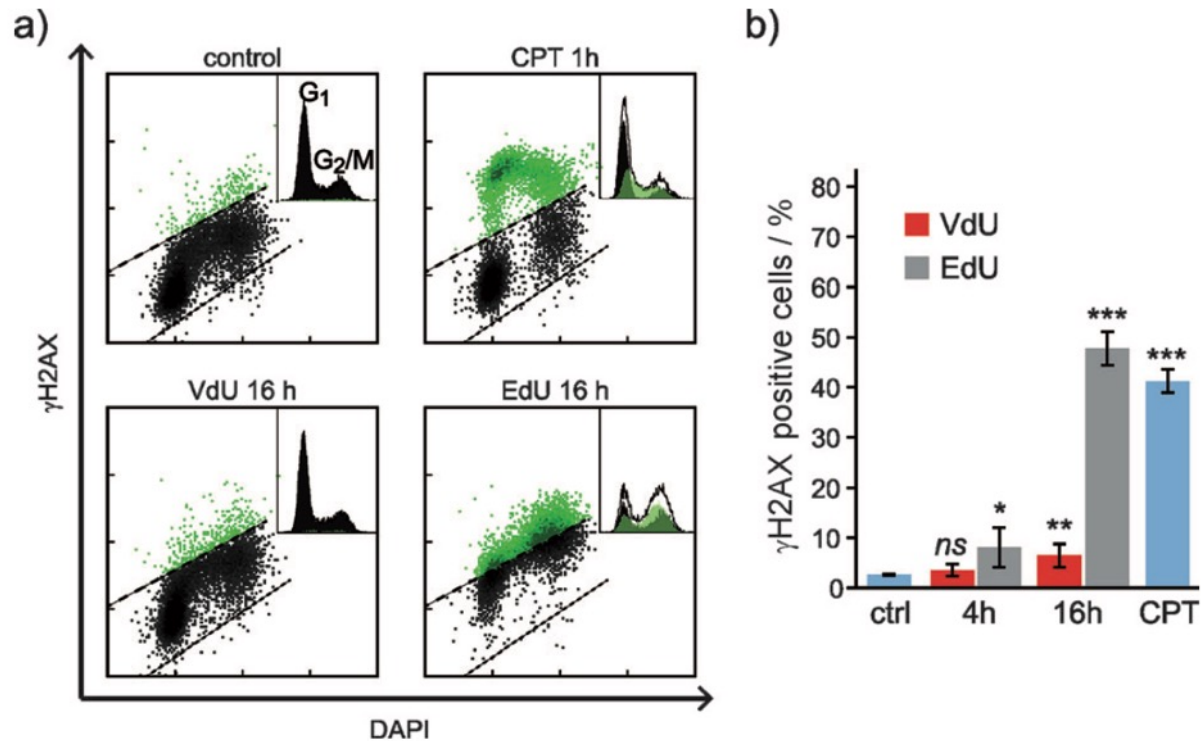
→ The VdU ligation is bio-orthogonal to EdU.

### 3. Paper: a) VdU is incorporated into DNA and can be visualized with Tamra-Tz



**Figure 2.** Bioorthogonal invDA ligation of VdU in newly synthesized DNA upon the addition of the fluorescent tetrazine Tamra-Tz. HeLa cells were treated with 30  $\mu\text{M}$  VdU for 16 h, followed by fixation, DNA denaturation, and incubation with 5  $\mu\text{M}$  Tamra-Tz for 4 h. Total cellular DNA was stained with DAPI. The negative control (neg. ctrl.) samples received identical treatment but were not incubated with VdU prior to staining. Scale bars: 50  $\mu\text{m}$ . DIC = differential interference contrast image, merge = overlay of Tamra-Tz and DAPI channels.

### 3. Paper: a) VdU is less toxic than EdU



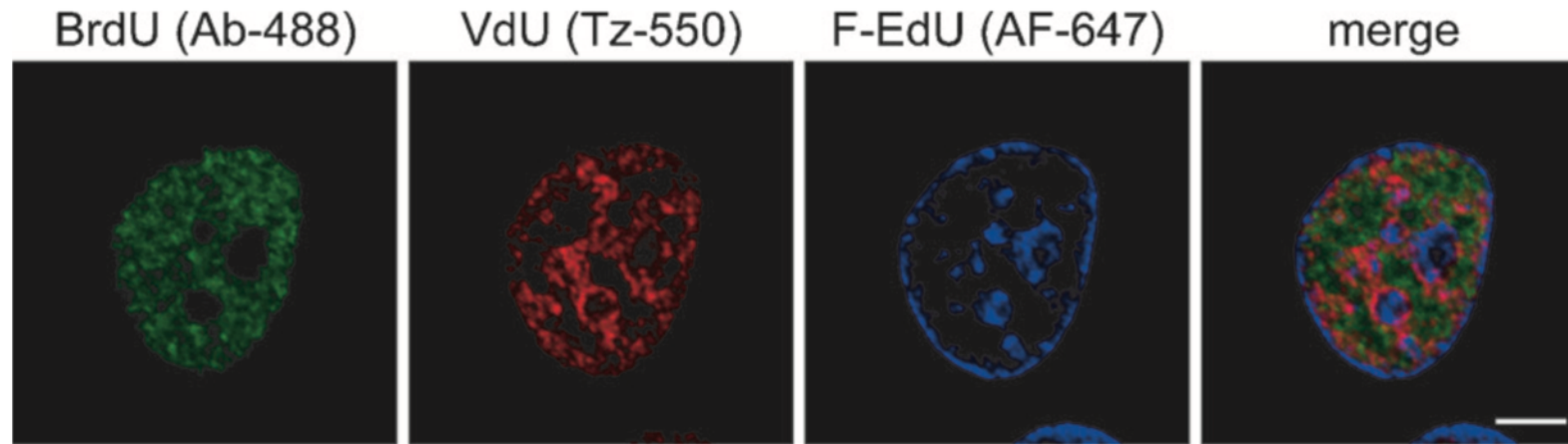
**Figure 3.** Flow cytometry analysis of  $\gamma$ H2AX versus DAPI staining of HeLa cells treated with 30  $\mu$ M VdU or 10  $\mu$ M EdU for 16 h. These concentrations were selected based upon the equivalent staining frequencies and intensities observed for EdU and VdU (Figure 4 b, c, and Figure S15, 16).  $\gamma$ H2AX positive cells were detected by using a phosphospecific antibody. a) The dot plots illustrate  $\gamma$ H2AX abundance versus the total cellular DNA content of each cell. The insets display histograms of the DNA content (from the cell cycle phases G<sub>0</sub>/G<sub>1</sub> to G<sub>2</sub>/M) versus cell count. The dashed lines show the thresholds used for defining  $\gamma$ H2AX positive/negative cells based on the control. For a positive control, cells were treated for 1 h with camptothecin (CPT, 0.5  $\mu$ M), which causes DNA damage.<sup>[21]</sup> b) A graphical representation of results where  $n = 5$ , \* $P < 0.02$ , \*\* $P < 0.002$ , \*\*\* $P < 0.0001$ , ns = not significant, compared to the control. For 4 h time points and U2OS cells see Figure S11.

→ VdU less toxic as phosphorylation of histones H2AX is less pronounced.

→  $\gamma$ H2AX formation is associated with DNA strand breakage.



### 3. Paper: a) VdU can be used for multicolor imaging of newly synthesized DNA



**Figure 5.** Visualizing the progression of the S-phase in a single cell by using BrdU (30  $\mu\text{M}$ , 1<sup>st</sup> pulse, green), VdU (30  $\mu\text{M}$ , 2<sup>nd</sup> pulse, red), and F-*ara*-EdU (10  $\mu\text{M}$ , 3<sup>rd</sup> pulse, blue). HeLa cells were incubated for 2 h 45 min with each nucleoside and washed for 15 min with fresh media in between nucleoside treatments. After fixation and DNA denaturation, the cells were stained with Tamra-Tz (red), BrdU antibody (green), and AF-azide (blue). Scale bars: 5  $\mu\text{m}$ . See Figure S17 for additional examples.

Cell cycle:

