

**New generation of
sensors to map pH
dynamics and Chloride
transport in organelles**

TJC

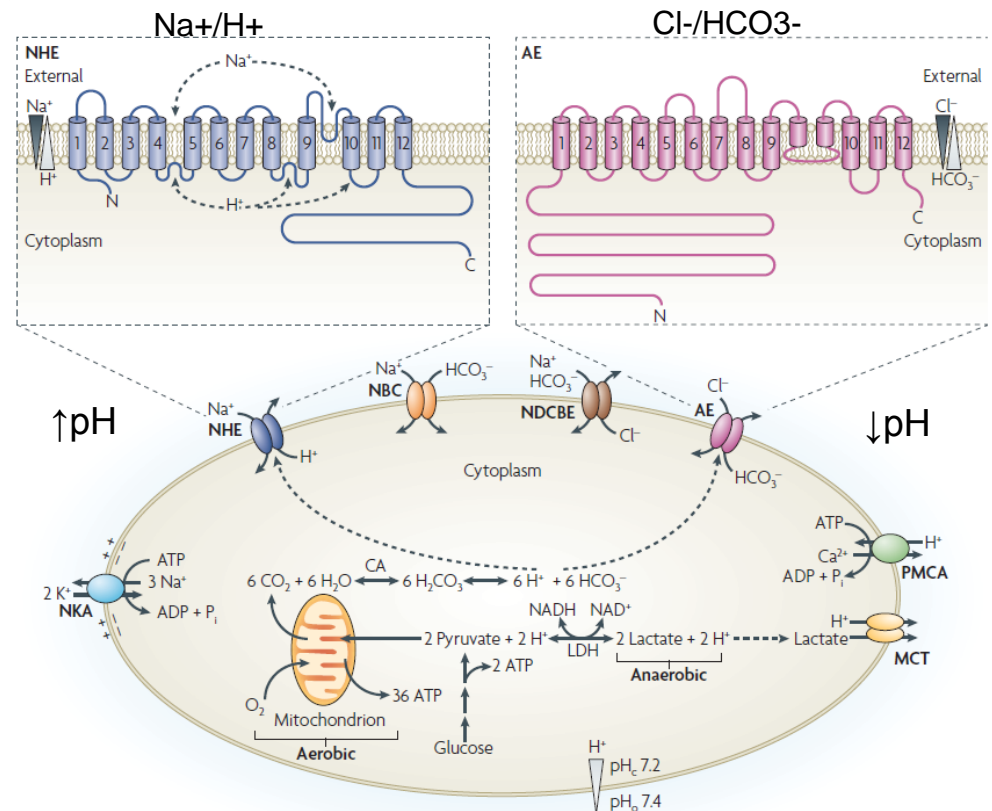
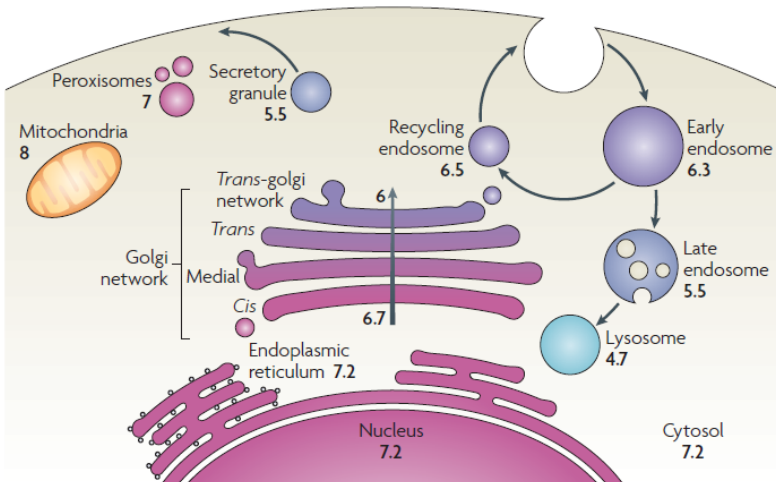
11.08.2015

Rochat Mary-aude

Subcellular pH

- > Cellular Compartmentalization: Segregation of specific function in organelles
- > Tendency of cytosol acidification
- > Multiple pH regulation mechanisms
- > Polarization and cell migration

7.3-7.4



pH of the endocytic pathway

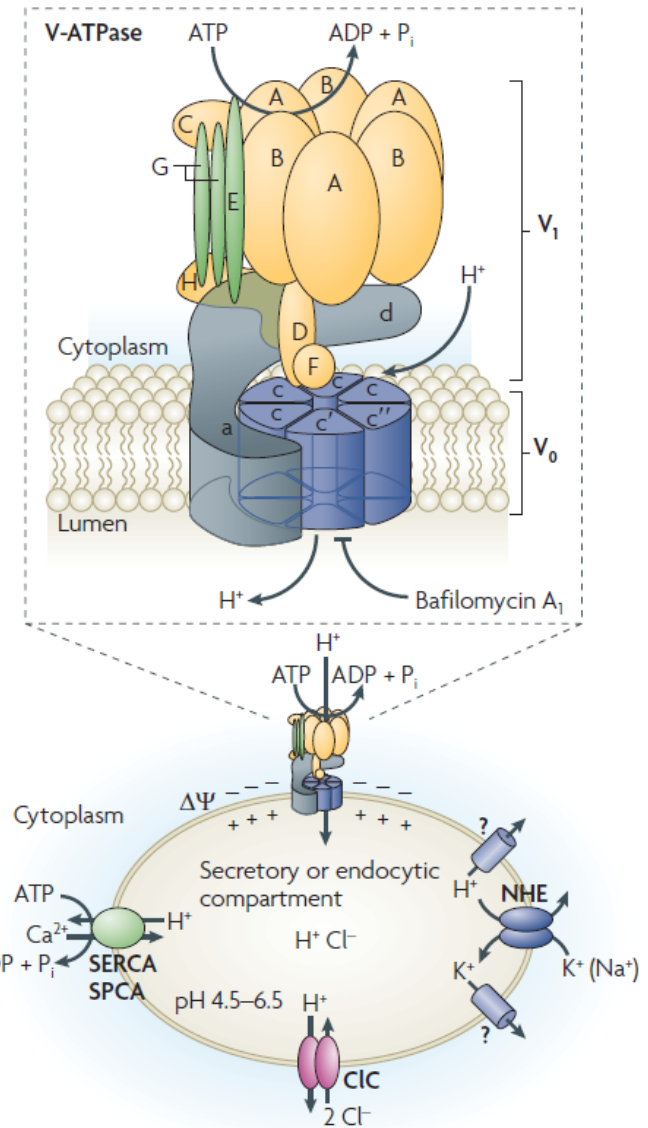
>Function:

- *Ligand-receptor uncoupling
- *Activation protease
- *Protonation of microbicidal factors

>V-ATPase

>ClC exchangers:

- Electrogenic
- Voltage dissipation
- Implicated in disease



Two DNA nanomachines map pH changes along intersecting endocytic pathways inside the same cell

Souvik Modi¹, Clément Nizak², Sunaina Surana¹, Saheli Halder¹ and Yamuna Krishnan^{1*}

Aim: Simultaneous use of different DNA nanodevices to map the pH changes within the same living cells

Principle: Nanodevice programmed to enter via different pathway
Early endosome and trans-golgi network

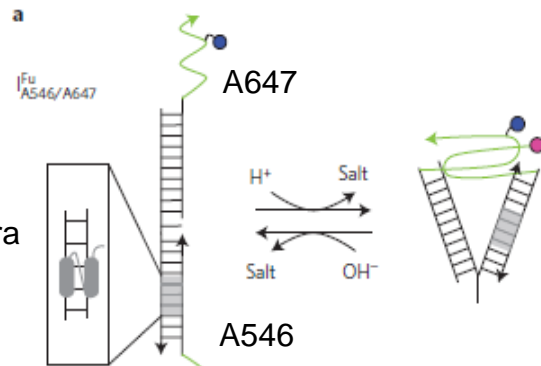
SympHony: Simultaneous pH mapping technology

SimpHony

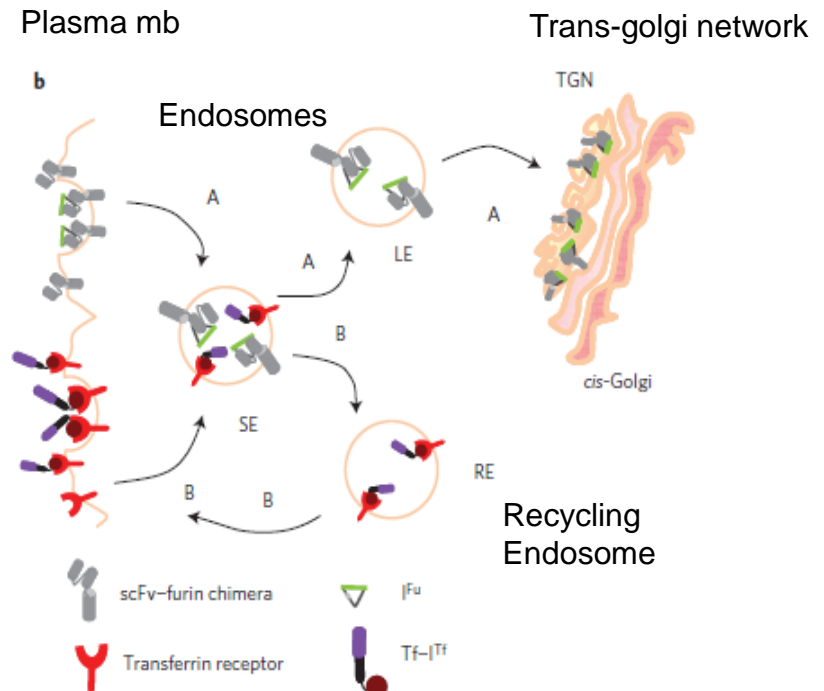
- Simultaneous pH mapping technology
 - >FRET-based DNA nanodevice
 - *Subcellular targeting : -Furin retrograde endocytic pathway
 - Transferrin endocytic/recycling pathway
 - *FRET-pairs: Alexa 546/647
 - Alexa 488/647

Furin

scFv-furin chimera
Binding element



Transferrin



Characterization of scFv

> Phage display screen of scFv recombinant antibodies against dsDNA of Furin nanodevice

* Sequence specific binding capacity

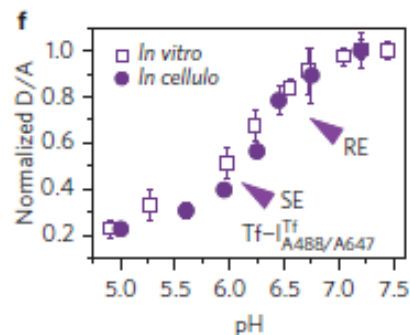
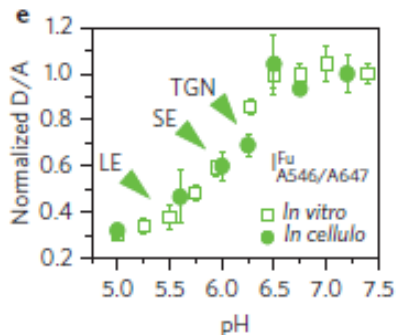
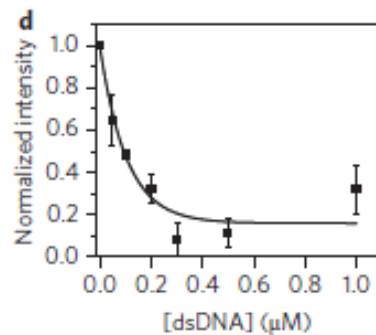
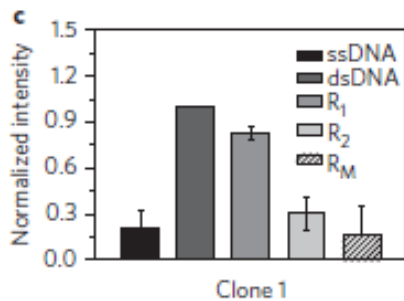
* K_d : 80nM

> Expression of scFv-furin

> Artificial receptor for nanodevice

* pH dependant FRET

* Relevant pH sensitivity for the targeted organelle



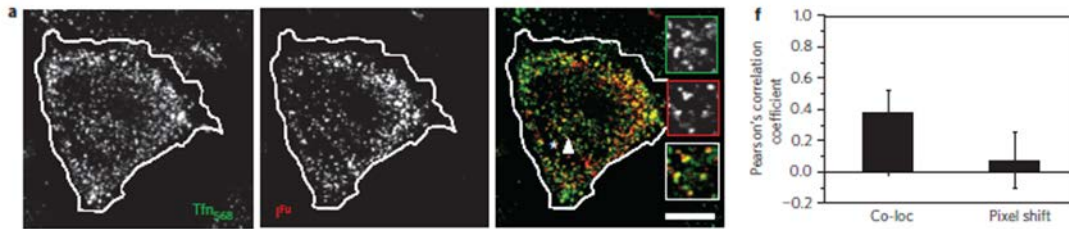
LE: late endosome

SE: sorting endosome

RE: recycling endosome

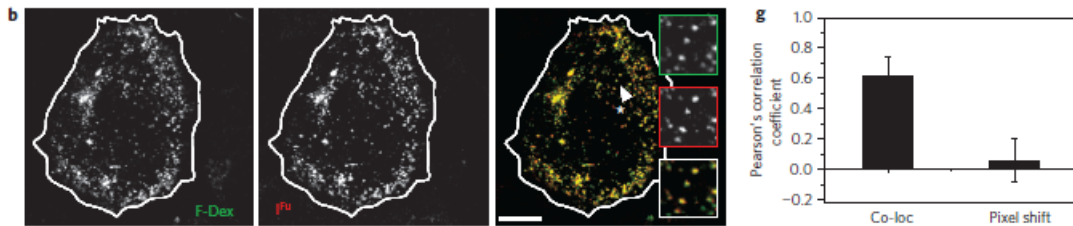
Trafficking specificities

Furin-nanodevice

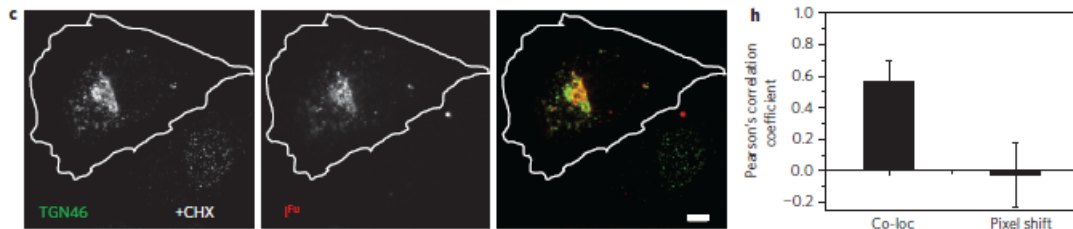


+ A568-transferrin:
EE/SE co-localization

+Fic-Dextran:
Chased 60min
Late endosome



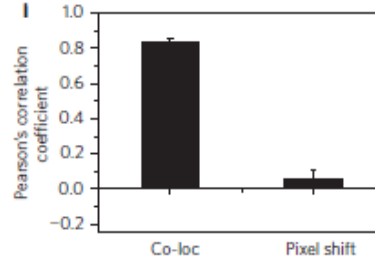
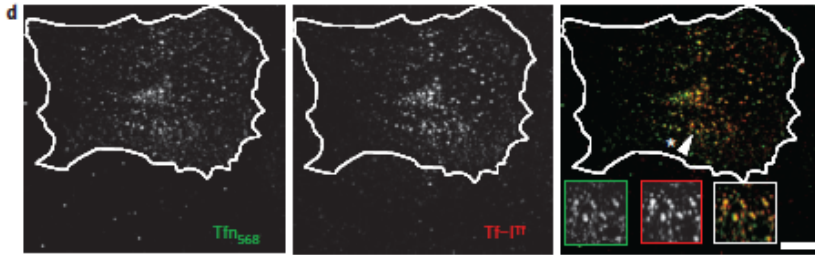
+TGN46 + Cycloheximid
Chased 90min
TGN network



>> Specificities of Furin
nanodevice to the furin
retrograde pathway

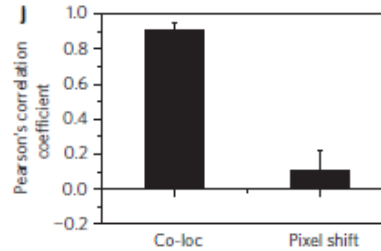
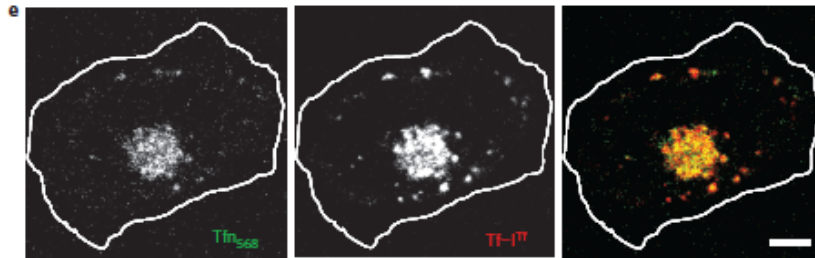
Trafficking specificities

Transferrin-nanodevice



>A568-transferrin +
Tf-Irf

10min Pulse:
Sorting endosomes

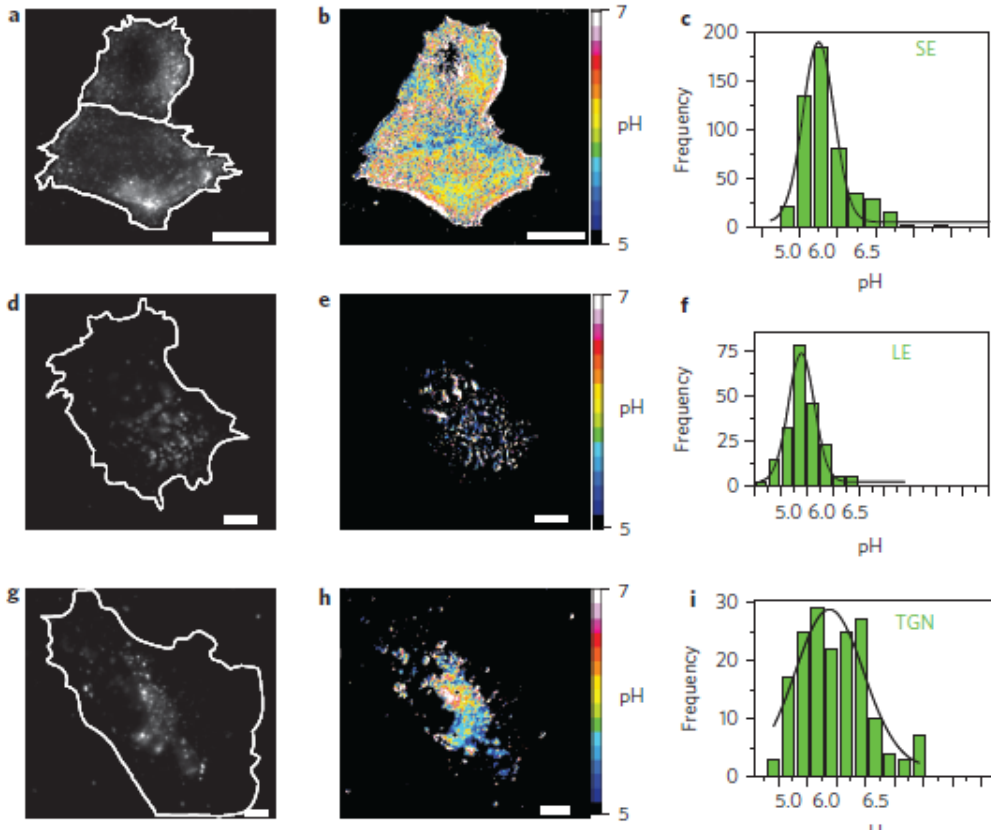


10min Pulse/12min
Chase:
Perinuclear Recycling
endosomes

DNA nanodevice are trafficking down of their specific
endocytic pathways
Without perturbation of the ligand

Spatiotemporal pH mapping

- Furin nanodevice



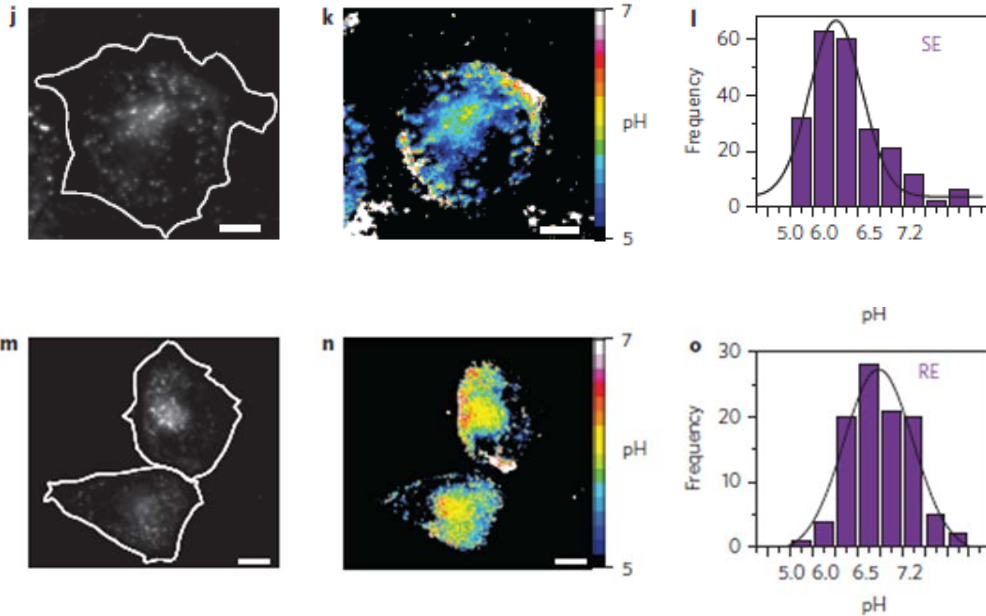
10min pulse
>EE/SE: pH of 6

30min pulse/45min chased
>LE: pH of 5.5

2h pulse/90min chased
Cycloheximid
>TGN: broader range of pH
From 5.5-6.5

Spatiotemporal pH mapping

- Transferrin nanodevice



10min pulse
>SE: pH of 6

10 min pulse/12min chase
>RE: pH of 6.5

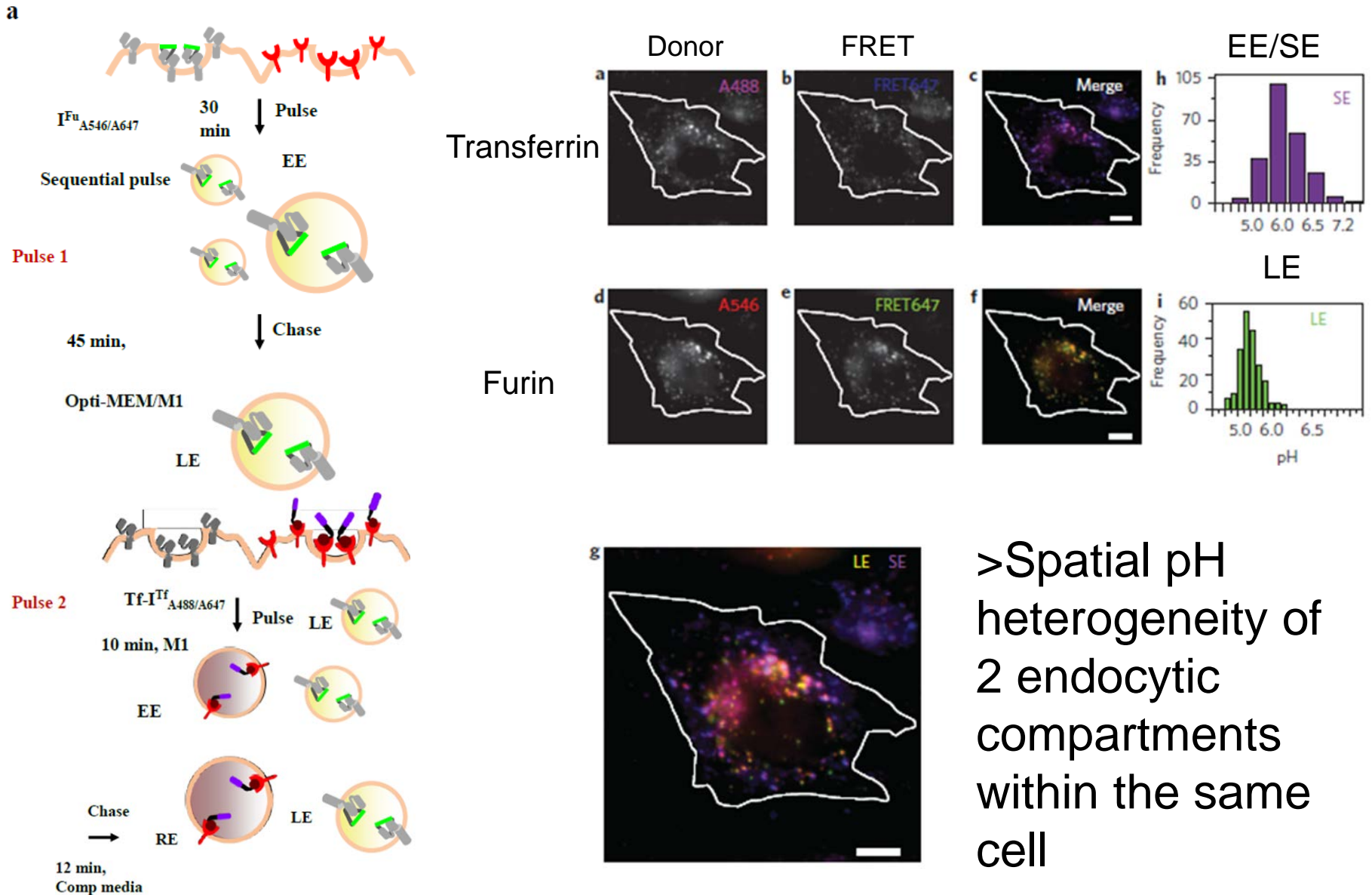
Each DNA nanomachine:

- *Specific endocytic pathway

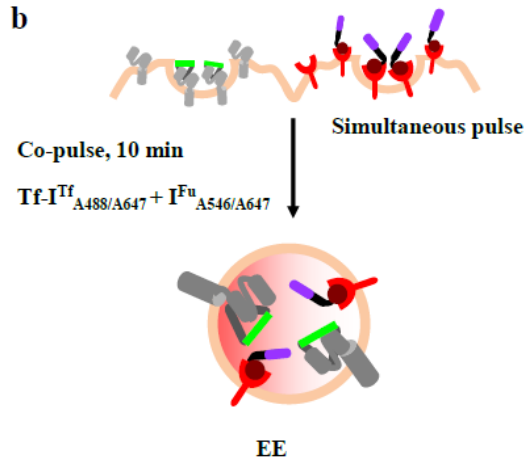
- *Quantitative pH sensing

- *High resolution spatiotemporal maps of subcellular organelles

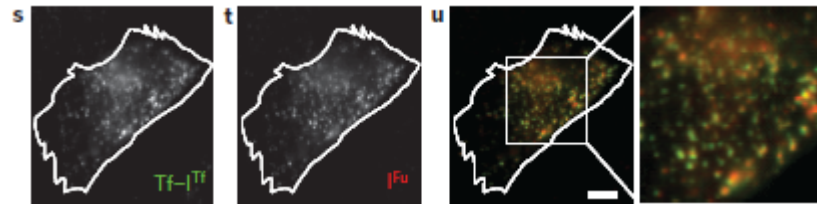
Sequential pH mapping



Simultaneous pH mapping



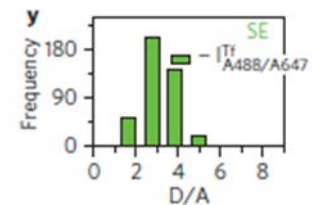
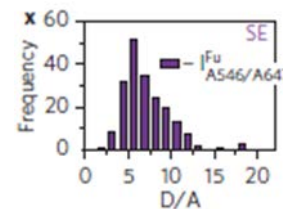
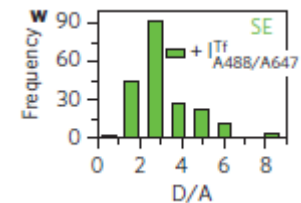
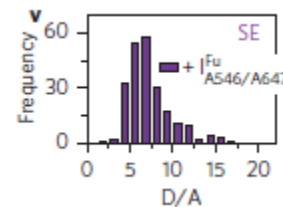
Co-localization in the early endosomes



Similar D/A ratio distribution in simultaneous Vs nanodevice alone

DNA nanodevices either alone or simultaneously, along with literature reported.

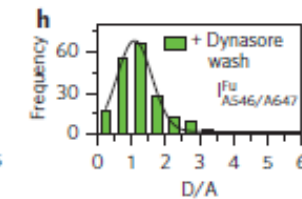
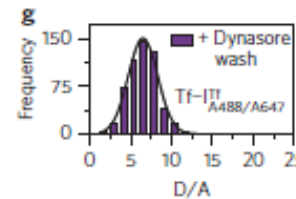
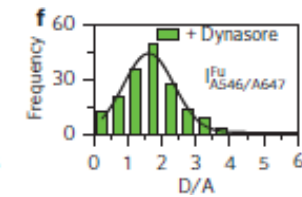
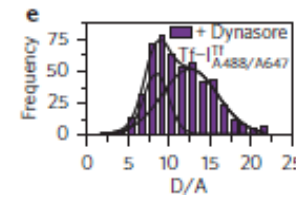
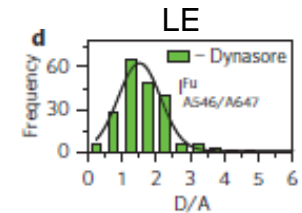
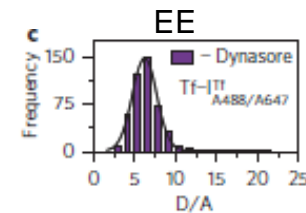
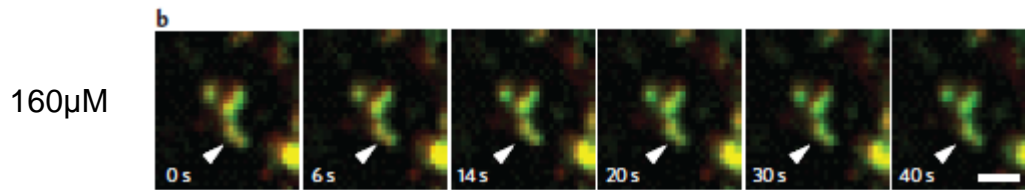
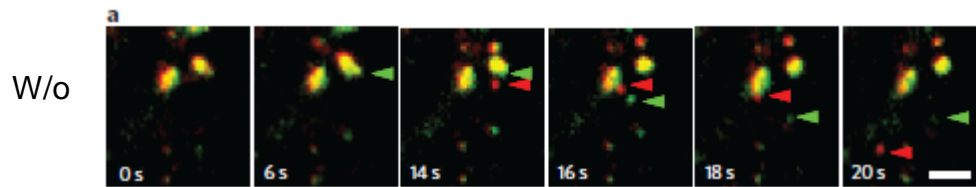
Compartments	I ^{Fu} _{A546/A647}		Tf-I ^{Tf} _{A488/A647}		Reported ^a	Reported ^b
	Single	SimpHony	Single	SimpHony		
SE	5.98 ± 0.2	6.0 ± 0.05*	6.09 ± 0.01	6.09 ± 0.09	6.1 (2)	6.2 ± 0.1 (3)
LE	5.72 ± 0.8	5.43 ± 0.19	-----	-----	5.8 ± 0.1 (4)	5.2-5.8 (5)
RE	-----	-----	6.35 ± 0.04	6.56 ± 0.11	6.5 (2)	6.43 ± 0.03 (6)
TGN	6.18 ± 0.01	6.16 ± 0.09	-----	-----	5.95 ± 0.03 (7)	6.19 (8)



Generation of non-interfering and autonomous DNA-based pH-sensitive nanodevice

Organelles morphology

- Dynasore: *LE: furin (red)
*EE: transferrin (green)



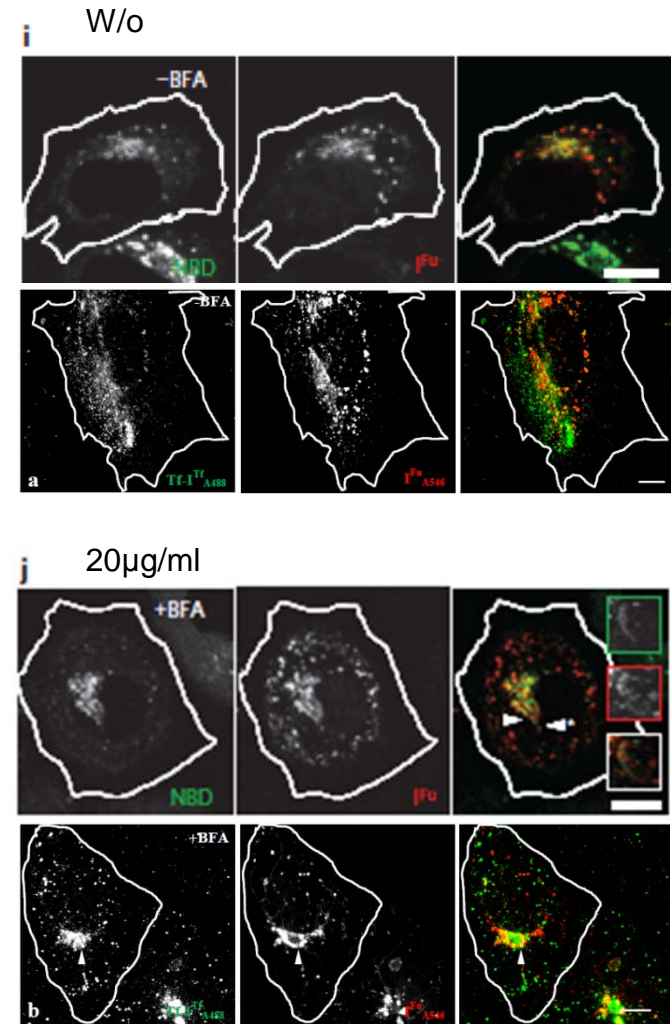
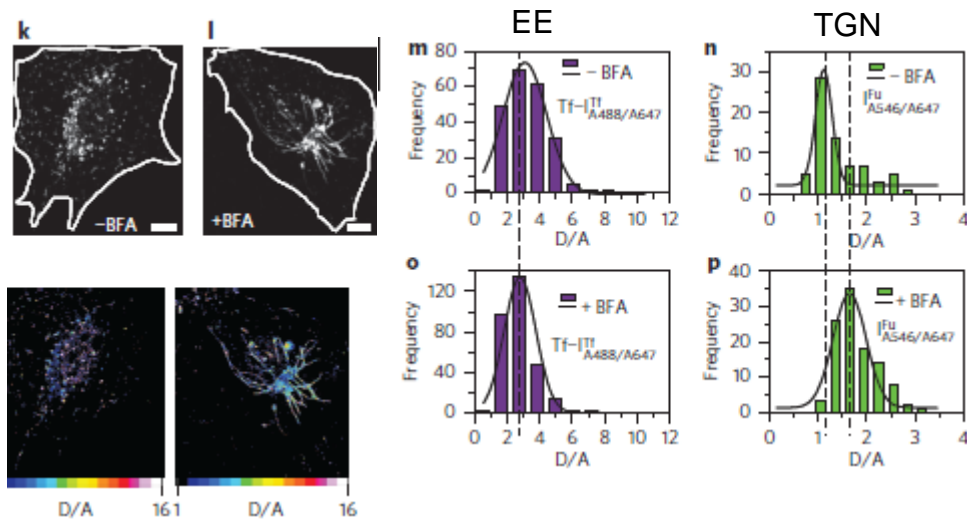
- >Inhibit EE fission
- >EE tubulation characteristics
- >Elevation of luminal pH in the EE
- >LE is unaffected
- >Restoration of endosomal dynamics and pH after wash-out

- Hypoacidification of the lumen in the tubular EE due to dynasore

Organelles morphology

- **Brefeldin A:**

- >Alteration distribution nanodevice
- >Tubular extension of the TGN
- >Higher pH in the Brefeldin A treated TGN
- >pH in EE is unaffected



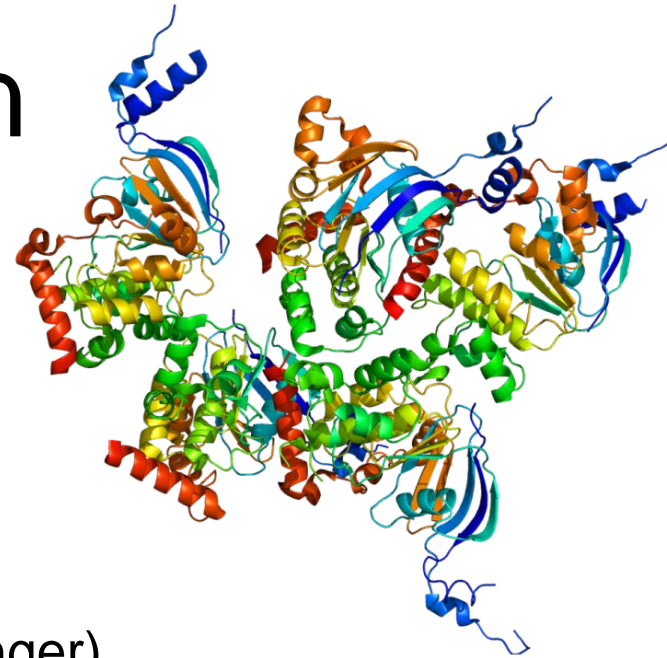
- Morphology alteration due to Brefeldin A is associated with defective acidification of the TGN

Conclusions

- Specific targeting of subcellular compartment using trafficking proteins
- Modular DNA nanodevice containing analyte sensing domains and fluorophores
- Simultaneous pH measurement in different compartments in living cells
- Pos
 - >Labeling in living cell
 - >Greater resolution than small molecule sensors
 - >iMotif stability is tunable (pH range)
 - >Generalizable to other chemical molecules
 - >Application for pH determination and compartments dynamics
- Neg
 - >Unprecise characterization of the subcellular compartments for Furin
 - >Different sensitivity/precision between nanodevice
 - >Fluorescence overlap
 - >Trafficking pathway and pH estimates needed
 - >Loss of signal

Chloride ion

- Broad range (5-130mM)
- Tightly regulated :
 - >Chloride channels
 - >Cl⁻/H⁺ exchanger
- Functions:
 - >pH regulation (chloride and bicarbonate exchanger)
 - >Cell excitability/secretion (cell mb)
 - >Mb voltage dissipation (endo-lysosome)
 - >Volume homeostasis (ClC2)
 - >Phagosome (HOCl)
- Disease:
 - >Cystic fibrosis: lung, pancreas, intestine..(CFTR)
 - >Epilepsy/congenital myotonia (ClC1)
 - >Bartter's syndrome: kidney (ClCkb)
 - >Dent's disease: kidney



Cl ion transporter sensing

Cl-sensitive small-molecule dye:

*SPQ

*MQAE

*Lucigenin

+ Bicarbonate and pH insensitive

- Not ratiometric

- Excitation in the UV range

- Cannot be specifically localized within organelles

- Unstable loading and retention over 30°C

- Chemical conjugation: *loss sensitivity (massive quenching)

*variable sensor characteristics

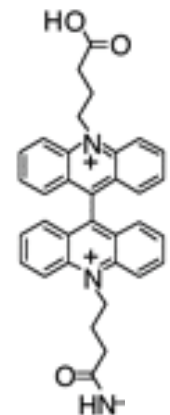
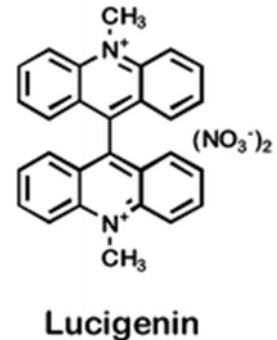
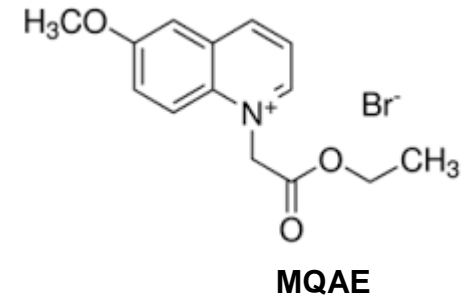
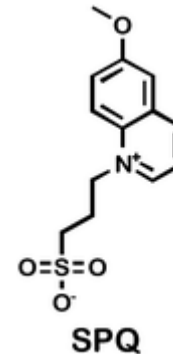
>Not suitable for live cell imaging

*BAC:

+ Bioconjugatable

+ Can be targeted to specific intracellular location

+ Wide range of Chloride concentration (0-200mM)



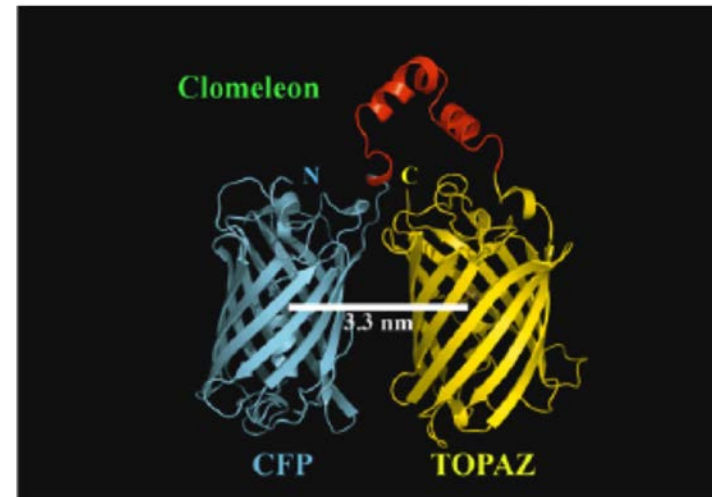
Cl ion transporter sensing

>Cl sensitive proteins reporters: YFP mutant/FRET

*Clomeleon

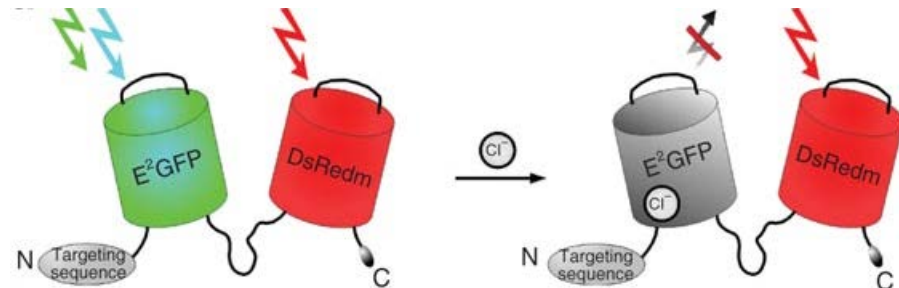
*Cl-sensor

- + Targetable
- + Photostable
- + Live imaging
- + Ratiometric
- pH sensitive and dependancy
- Low chloride affinity



*ClopHensor:

- + pH reporter module/correction factor
- Reduction of time resolution
- Complicated analysis



Need for

- >Modular device
- >Entire physiological range
- >Uniform Sensor characteristics
- >Ratiometric: ratio of 2 optical signals
- >pH independant
- >Live imaging
- >Targeted to specific pathway

A pH-independent DNA nanodevice for quantifying chloride transport in organelles of living cells

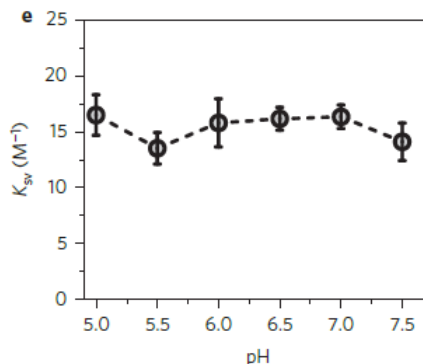
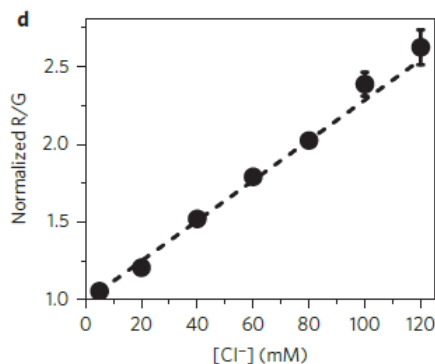
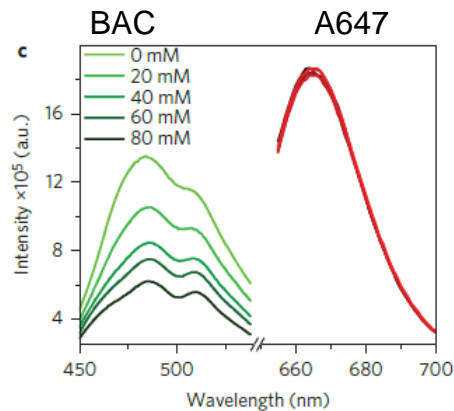
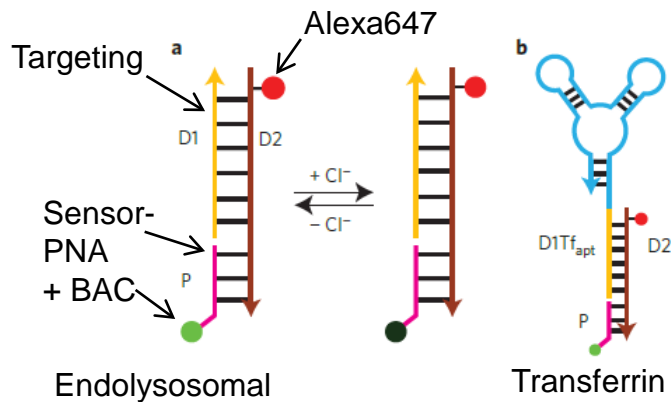
Sonali Saha¹, Ved Prakash², Saheli Halder¹, Kasturi Chakraborty² and Yamuna Krishnan^{1,2*}

Aim: precisely measure the activity and the subcellular location of chloride transporter in living cells using a nanodevice

Principle: specific targeting to the endolysosomal pathway

Clensor

- Fluorescent (BAC/A647) and ratiometric DNA-based nanodevice
- Coupled to endolysosomal/transferrin pathway
- pH independant



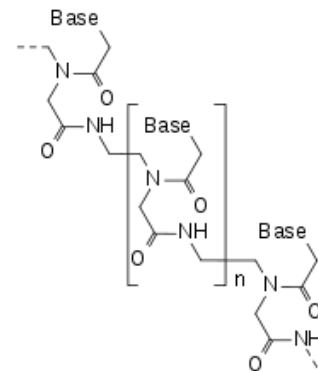
PNA: Peptide nucleic acid

Pos

- >Binding strength
- >High specificity
- >Short sequence
- >Resistant
- >Broad pH

Neg

- >Hydrophobic
- >High melting T^o

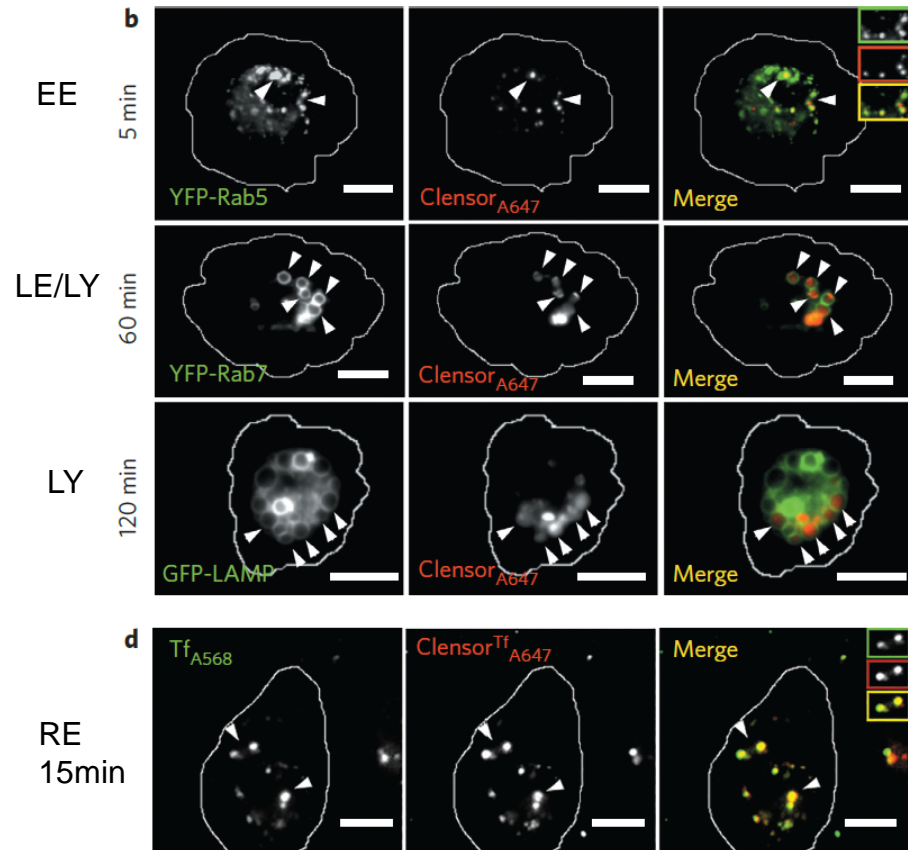
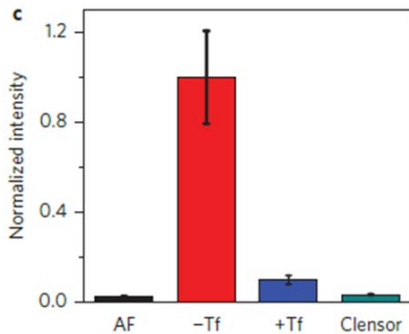
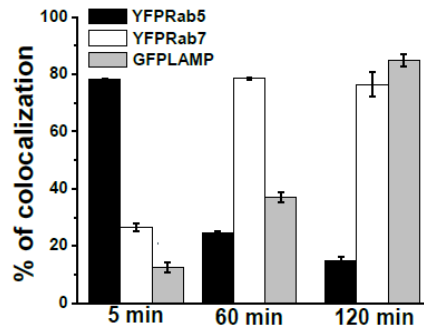
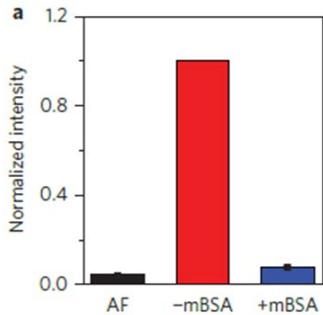


G (BAC)= 435/505nm
>Collisional quenching
R (A647)=650/670nm

Cl- measured in lysosome

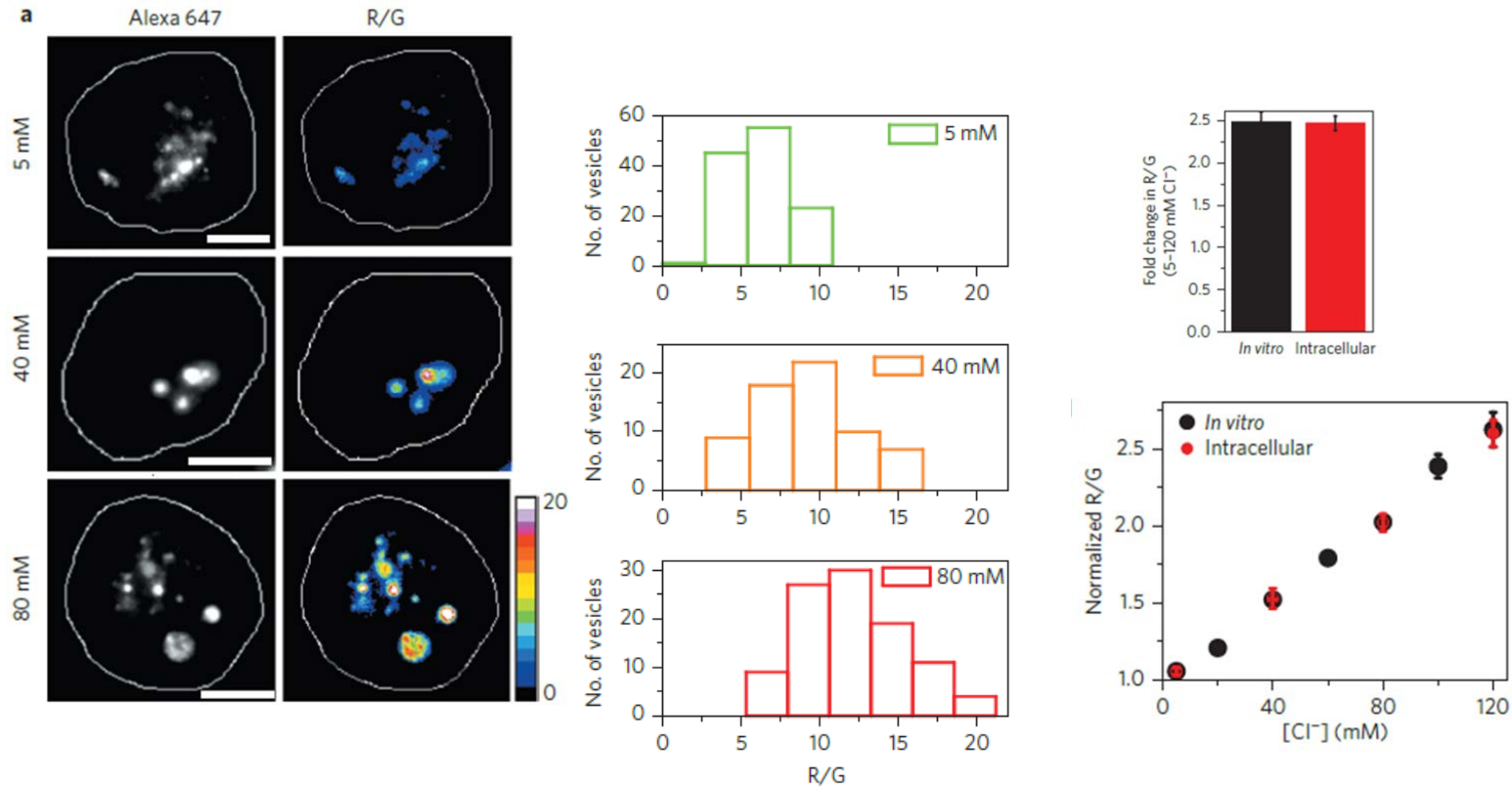
- *Drosophila* hemocytes:
 - >Endocytosis through ALBR
 - >Endolysosomal pathway migration
 - >Specific targeting Transferrin-pathway

Competition assay



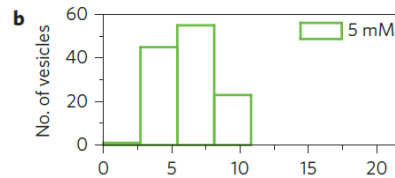
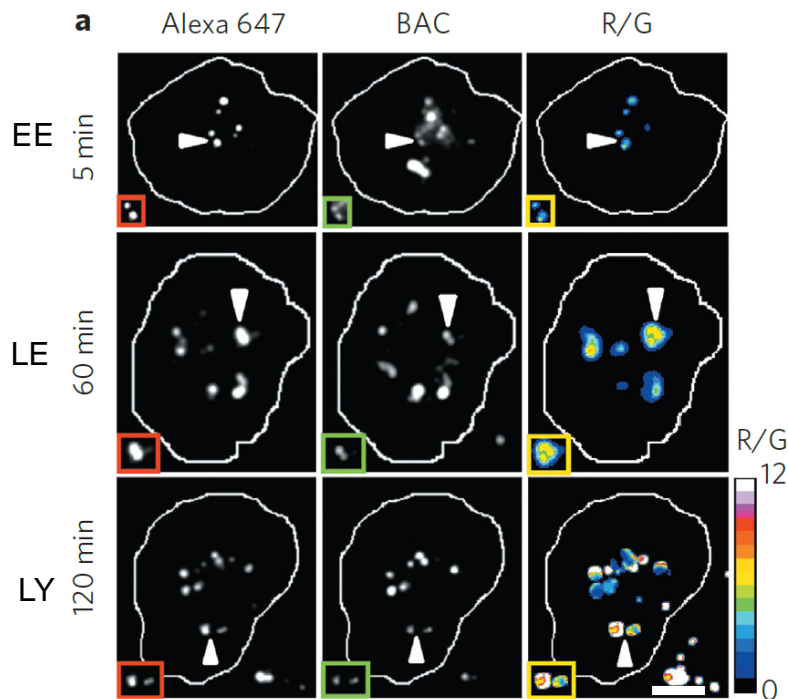
Intracellular functionality of Clensor

- Quantitative, linear, $[Cl^-]$ dependant R/G ratio
- 2.5fold between 5-120mM
- Endocytosis does not affect sensing properties of Clensor

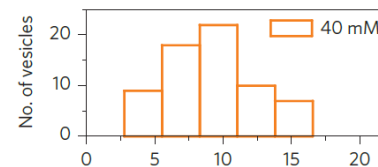


Spatiotemporal change of [Cl⁻]

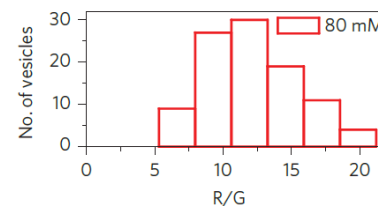
- Progressive accumulation of Cl⁻ along the endolysosomal pathway
- Reliable quantification
- Cl⁻ alteration detection after chemical perturbation of chloride conductance



37mM



60mM



108mM

Endosomes (min)	Mean [Cl ⁻] ± s.e.m (mM)	
	-NPPB	+NPPB
EE (5)	37.0 ± 1.6	9.3 ± 1.5
LE (60)	60.4 ± 2	33.8 ± 2.5
LY (120)	108.5 ± 1.4	86.5 ± 3.5

Localization and activity CIC

- Specific Knock-down of DmCIC-b and DmCIC-c

- DmCIC-c: EE

RE: moderately

>Inter-relation $[Cl^-]$ and pH

- DmCIC-b: LE

LY

>No role on pH, similar to CIC6,7

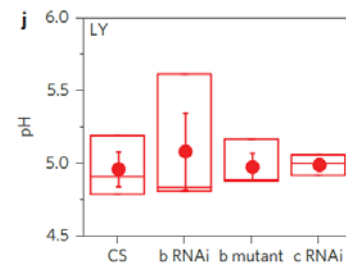
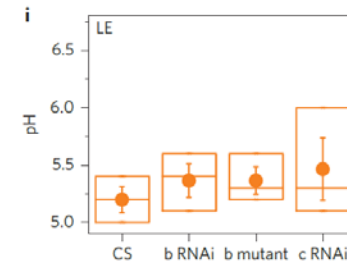
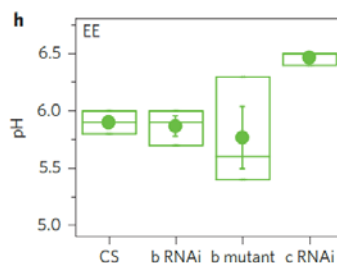
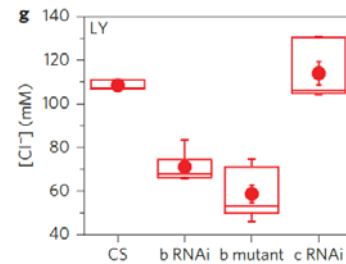
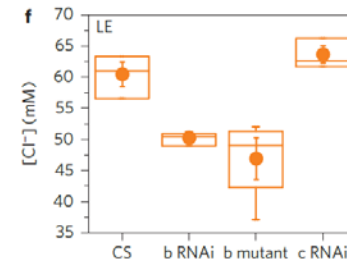
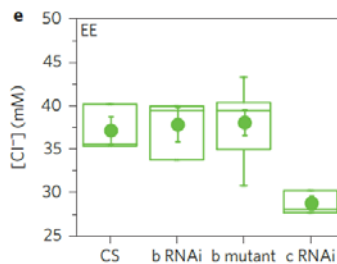
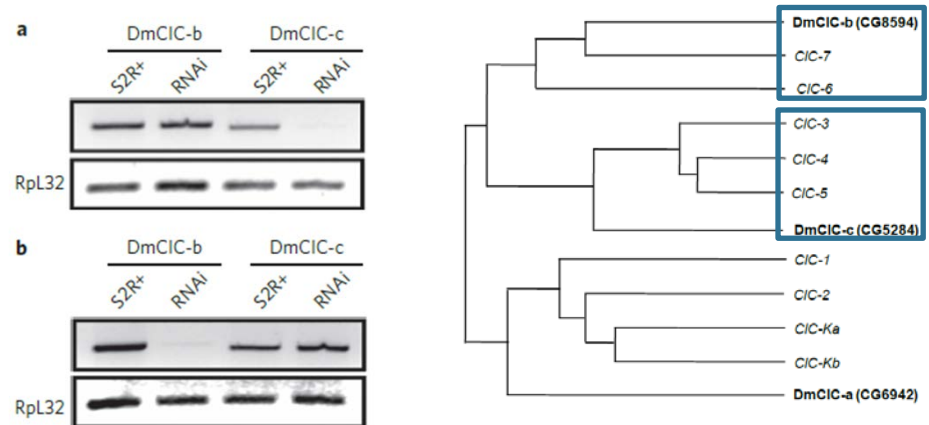


Table 1 | Luminal $[Cl^-]$ within recycling endosomes.

	Mean $[Cl^-]_{RE} \pm$ s.e.m. (mM)	Mean $pH_{RE} \pm$ s.e.m.
S2R+	39.9 ± 1.2	6.3 ± 0.09
DmCIC-c RNAi	33.1 ± 1.5	6.4 ± 0.03
DmCIC-b RNAi	39.1 ± 0.7	6.3 ± 0.09

Conclusions

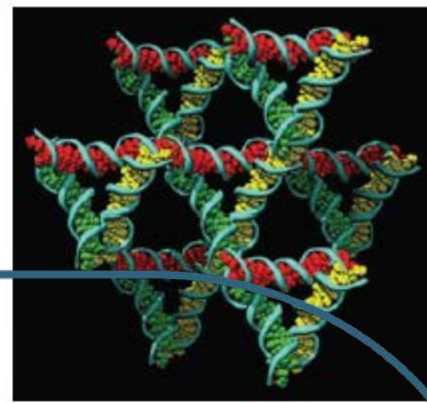
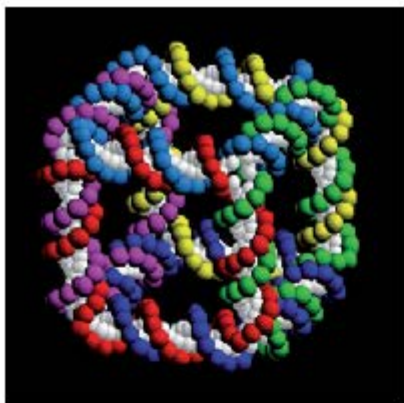
- Generation of modular nanodevice with identical sensor characteristics
 - >Sensing module: P-BAC
 - >Normalizing module: D2
 - >**Targeting module**: D1 (+/- aptamer)
- Quantitative and functional studies possible
 - >Physiology and Disease: *Lysosomal disorder
 - *Cystic fibrosis (CFTR)
 - *Golgi pH regulator (GPHR)
 - >Role of Chloride homeostasis : *Secretory pathway
 - *Synaptic vesicles

Pos

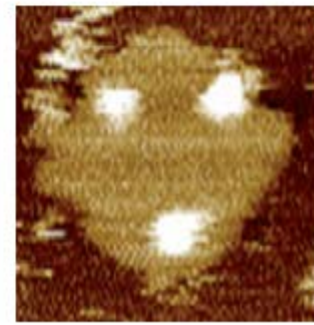
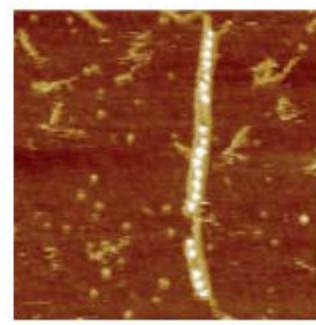
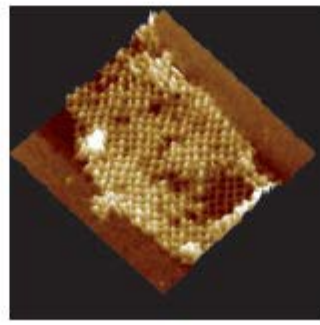
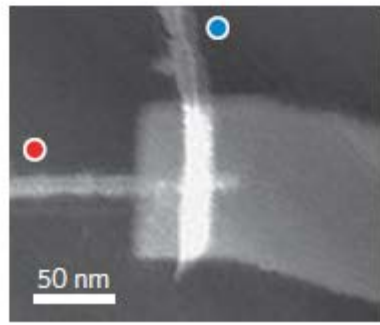
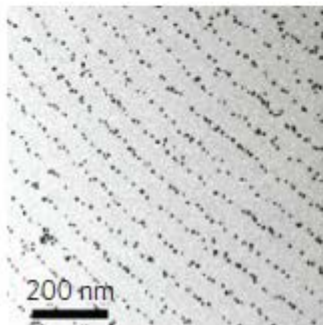
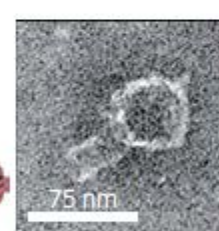
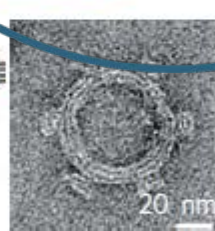
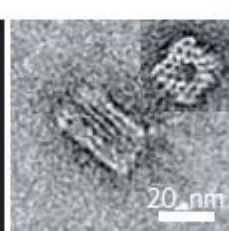
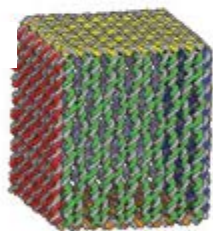
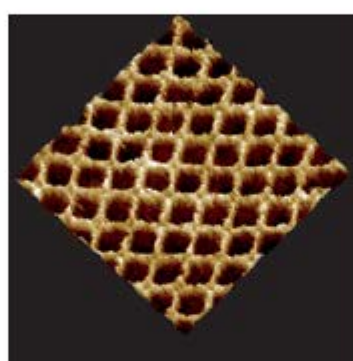
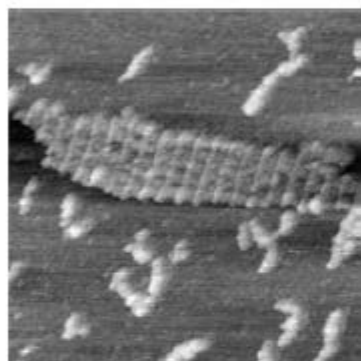
- >Labeling in living cell
- >Functional assay

Neg

- >Sensitivity?

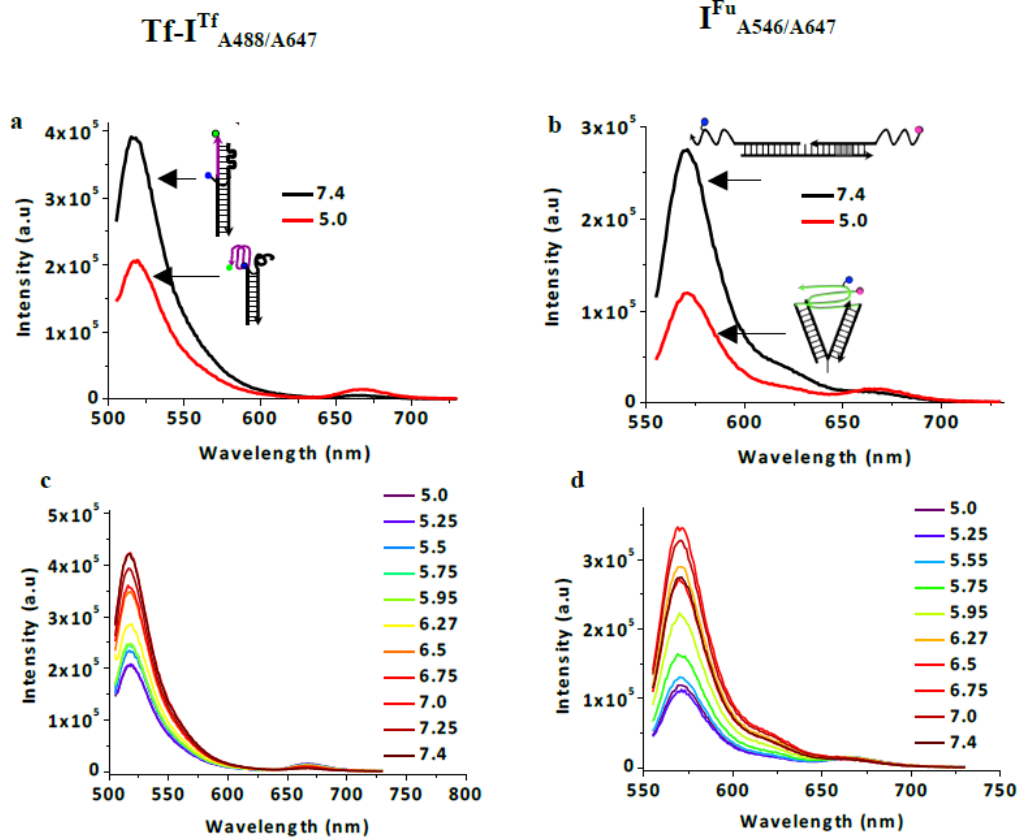


THANK YOU!



Back-up1

Steady state fluorescent spectra of DNA nanomachines at clamped pH



Back-up2

Clensor and *Clensor^{Tf}* sequences

Name	Sequence	Comment
P	BAC-NH _ε -Lys-ATC AAC ACT GCA-Lys-COOH	PNA strand: Sensing module
D2	5' TATA T ATA GGATCTTGCTGTCTGGTG TGC AGT GTT GAT 3'	DNA strand: Normalizing module; internal Alexa 647 modification on the T shown in bold
D1	5' CACCAGACAGCAAGATCC TATATATA 3'	DNA strand: Targeting module
D1Tf _{apt}	5' CACCAGACAGCAAGATCCTATATATAGGGGGA UC _{AA} UCC _{AA} AGGGA CCC _{GGAAA} C _G CUCCCUU _{ACA} CCCC 3'	DNA RNA hybrid strand : Targeting module with RNA aptamer against human transferrin receptor.

Back-up2

- Gel mobility shift assay

