

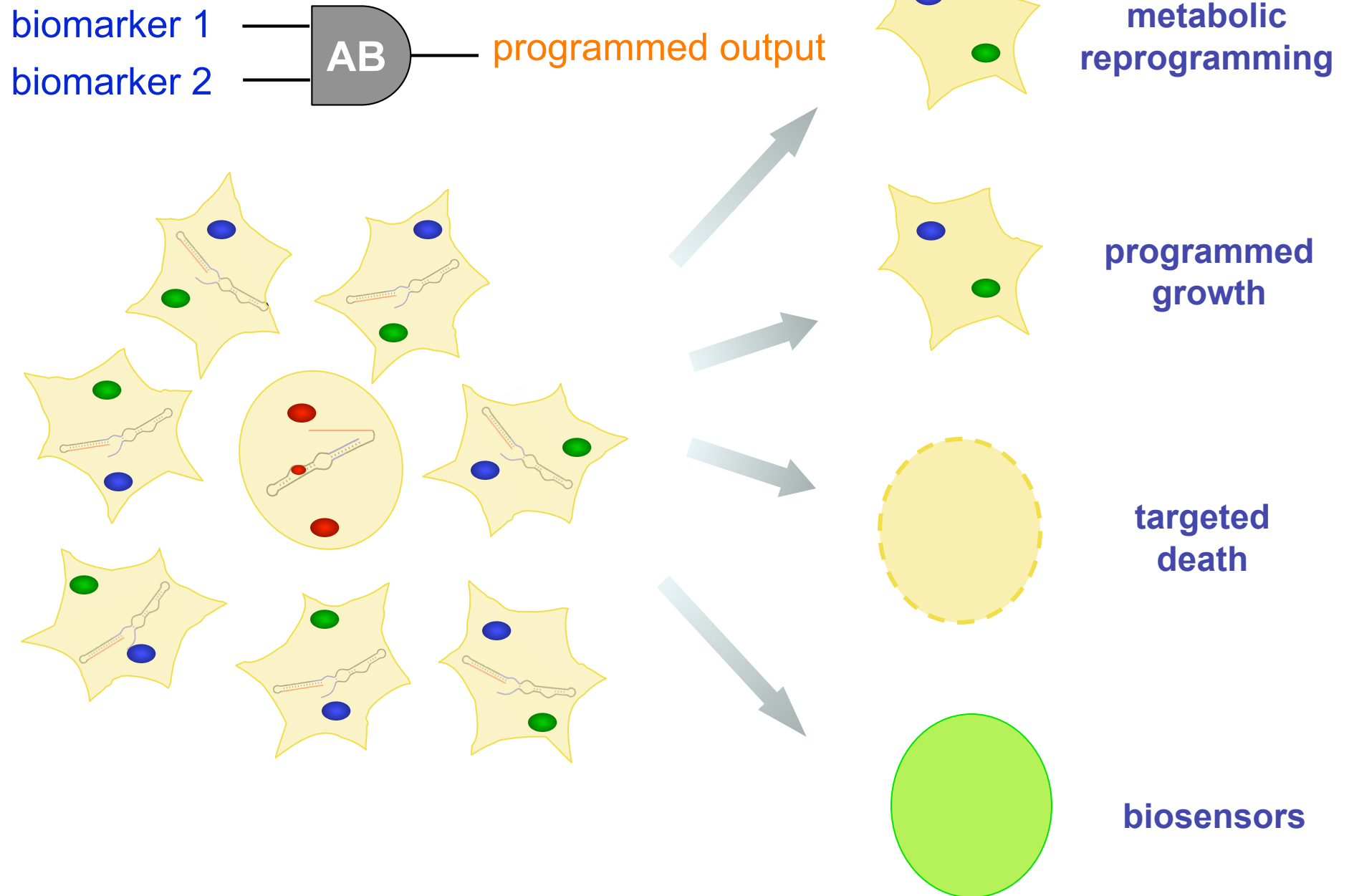
Programming cellular behavior with RNA controllers

Christina D. Smolke
Department of Bioengineering
Stanford University

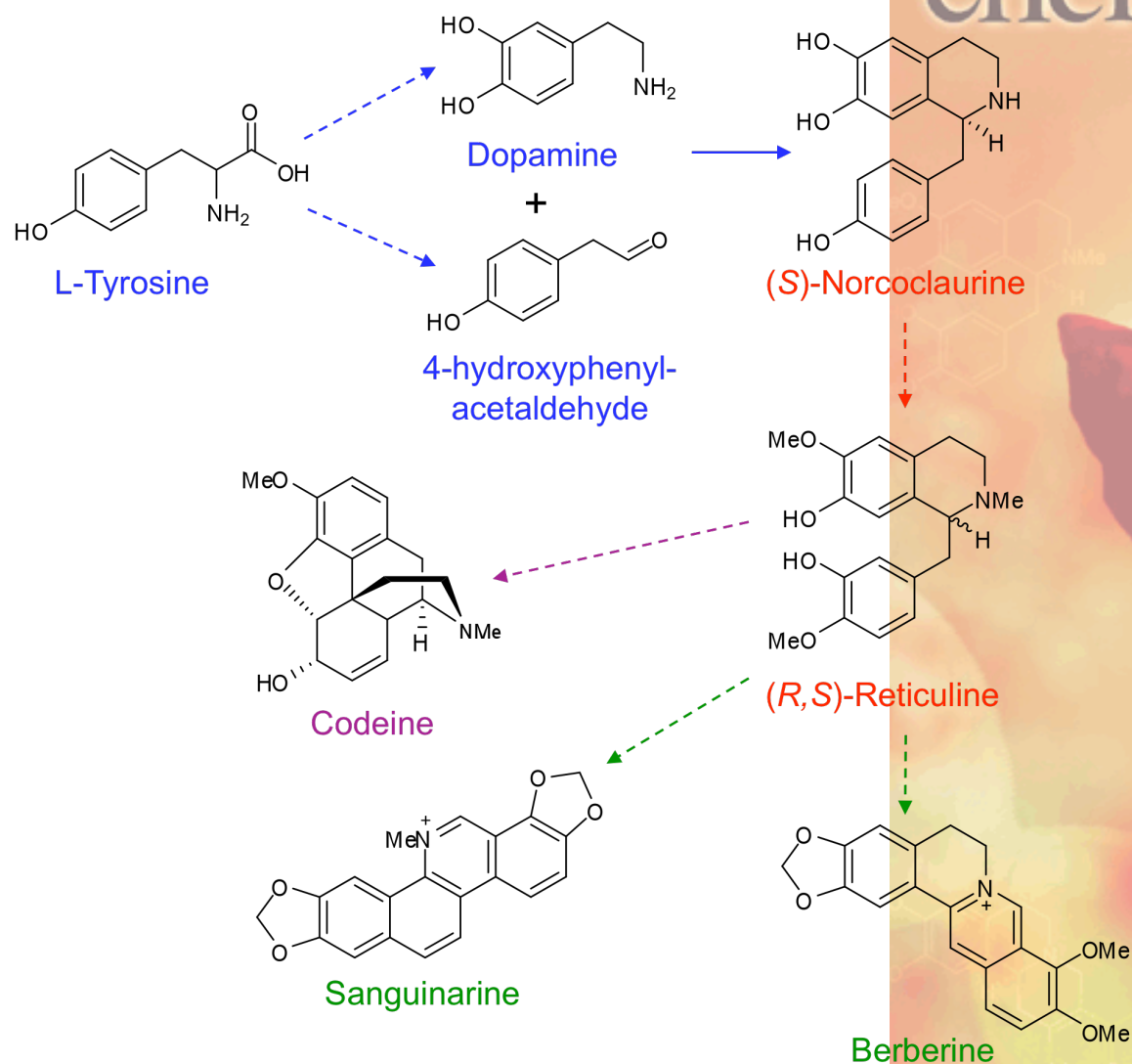
July 13, 2010
RoSBN Net Synthetic Biology Workshop
St. Anne's College, University of Oxford



Targeted therapeutics



Manufacturing and discovery platforms

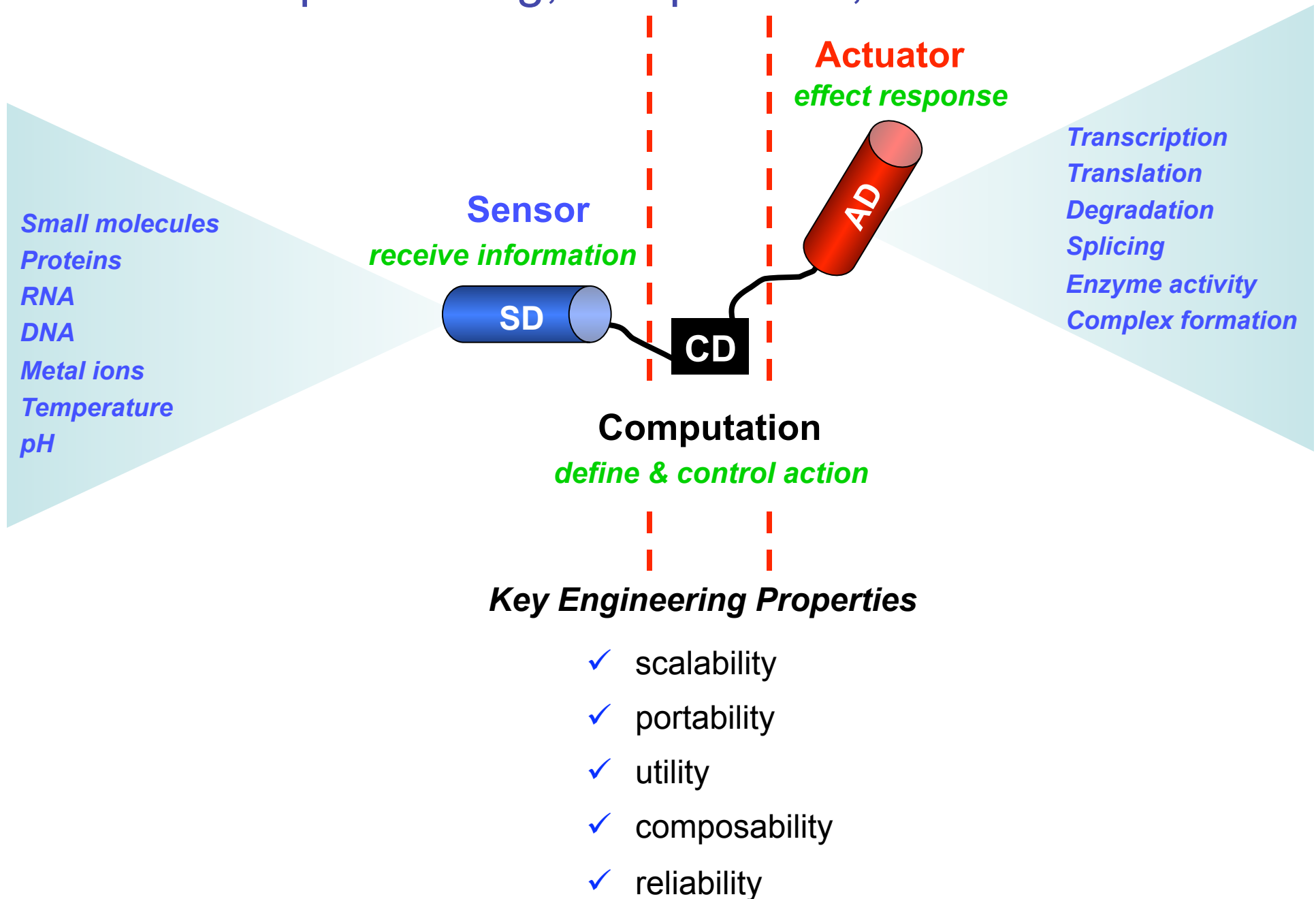


Apps

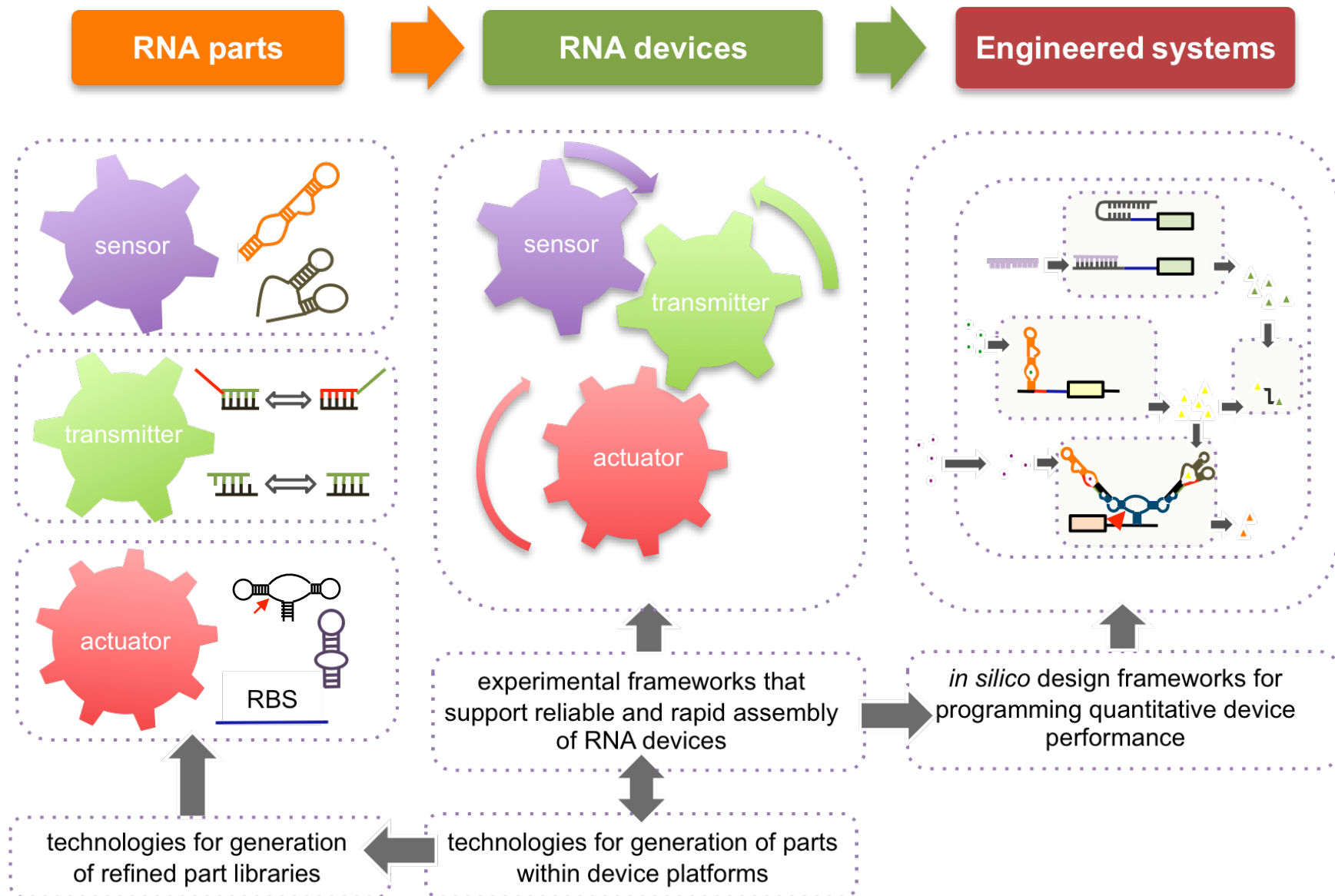
Tools

7. Control & dyn. systems
6. Reverse engineering
5. Fab, CAD & EDA
4. Standards & abstraction
3. Languages & grammars
2. Device design
1. Info. theory & signal proc.

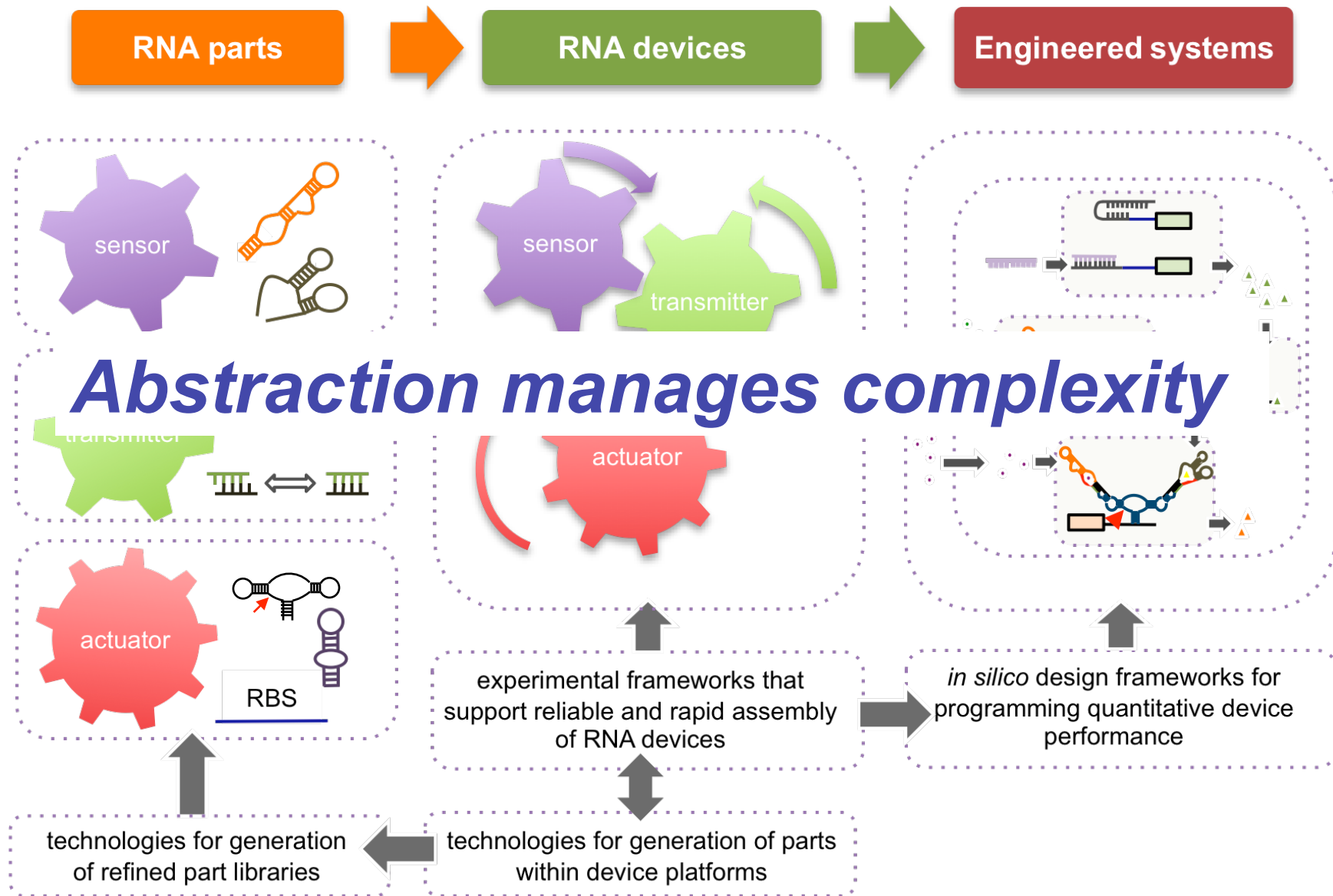
Information processing, computation, and control



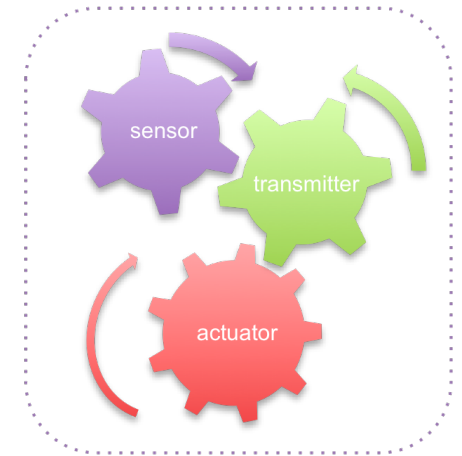
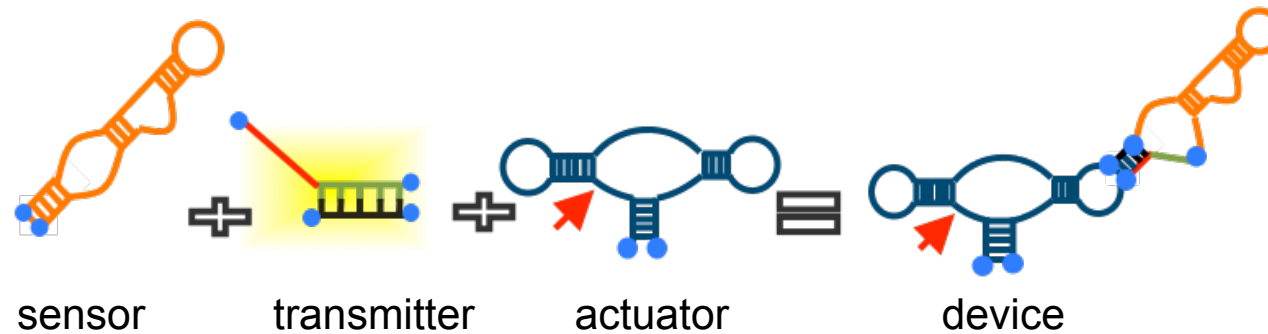
Plan for developing a programmable input/output tool



Plan for developing a programmable input/output tool

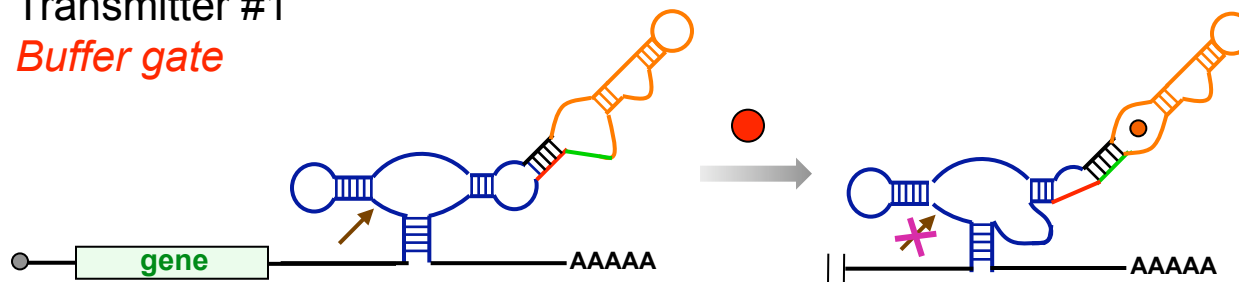


A platform for programming device function

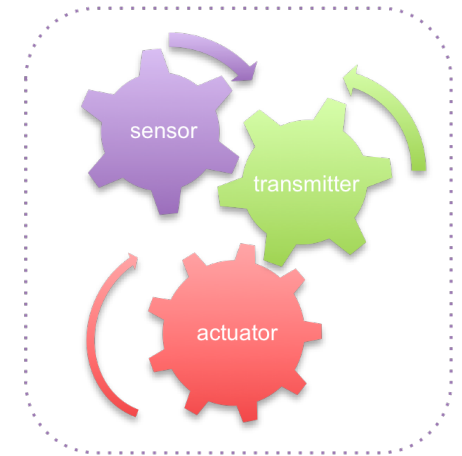
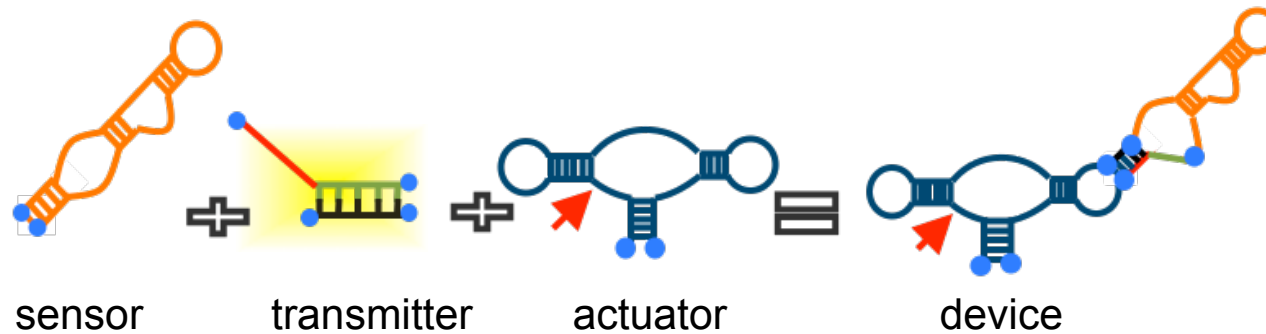


Transmitter #1

Buffer gate

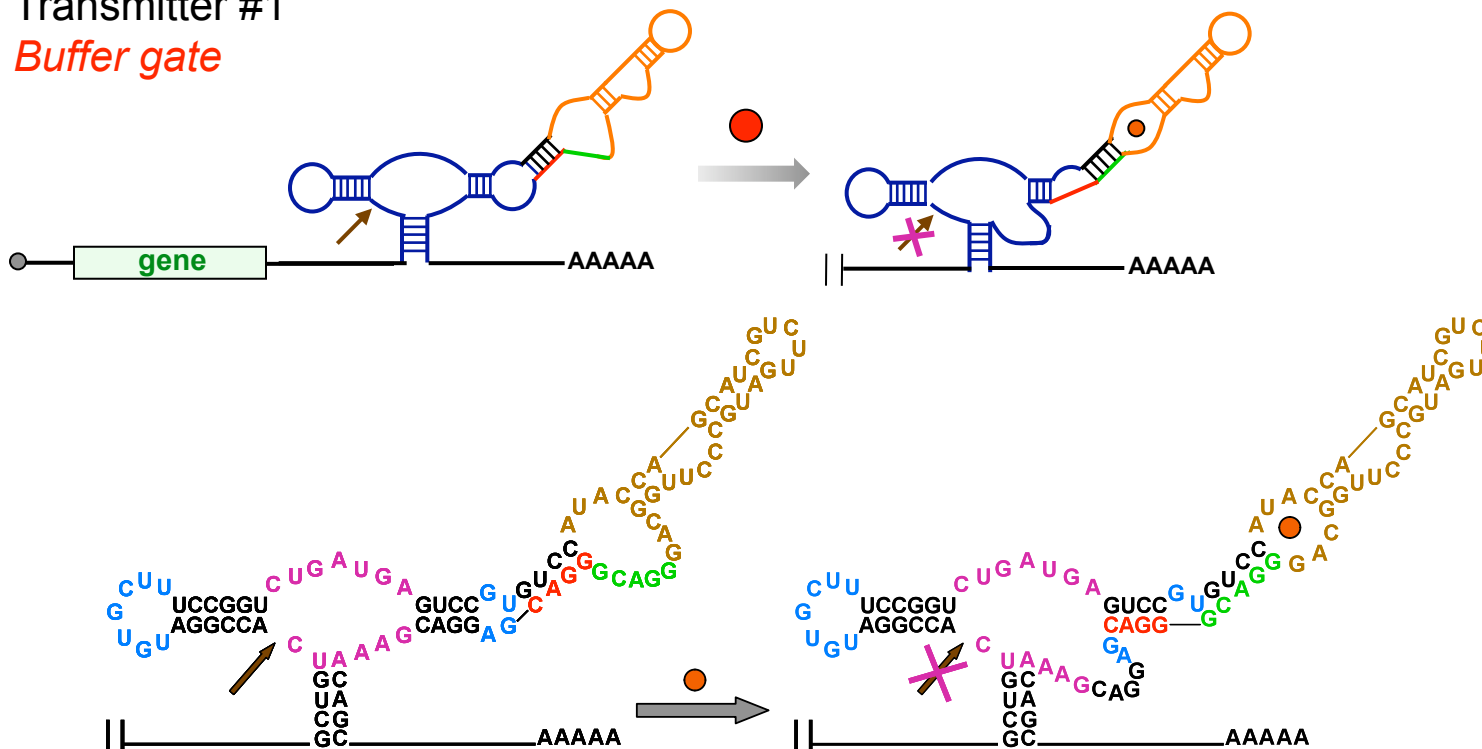


A platform for programming device function

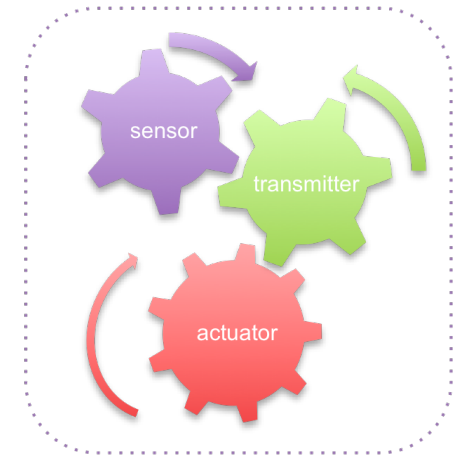
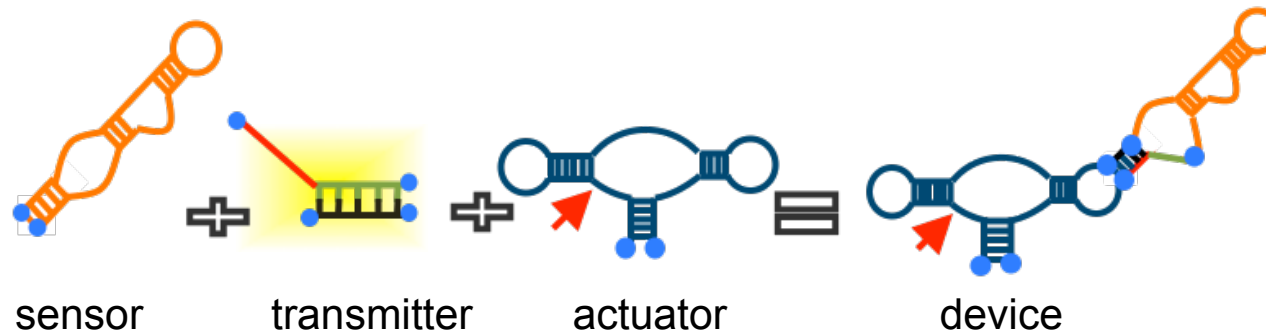


Transmitter #1

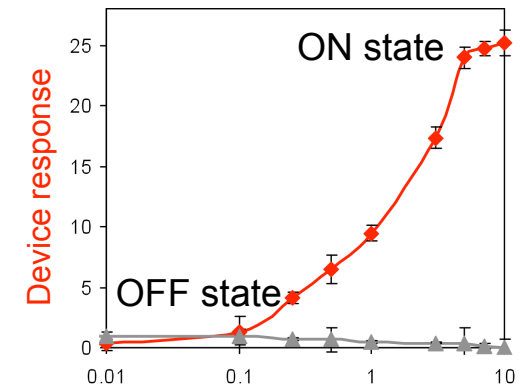
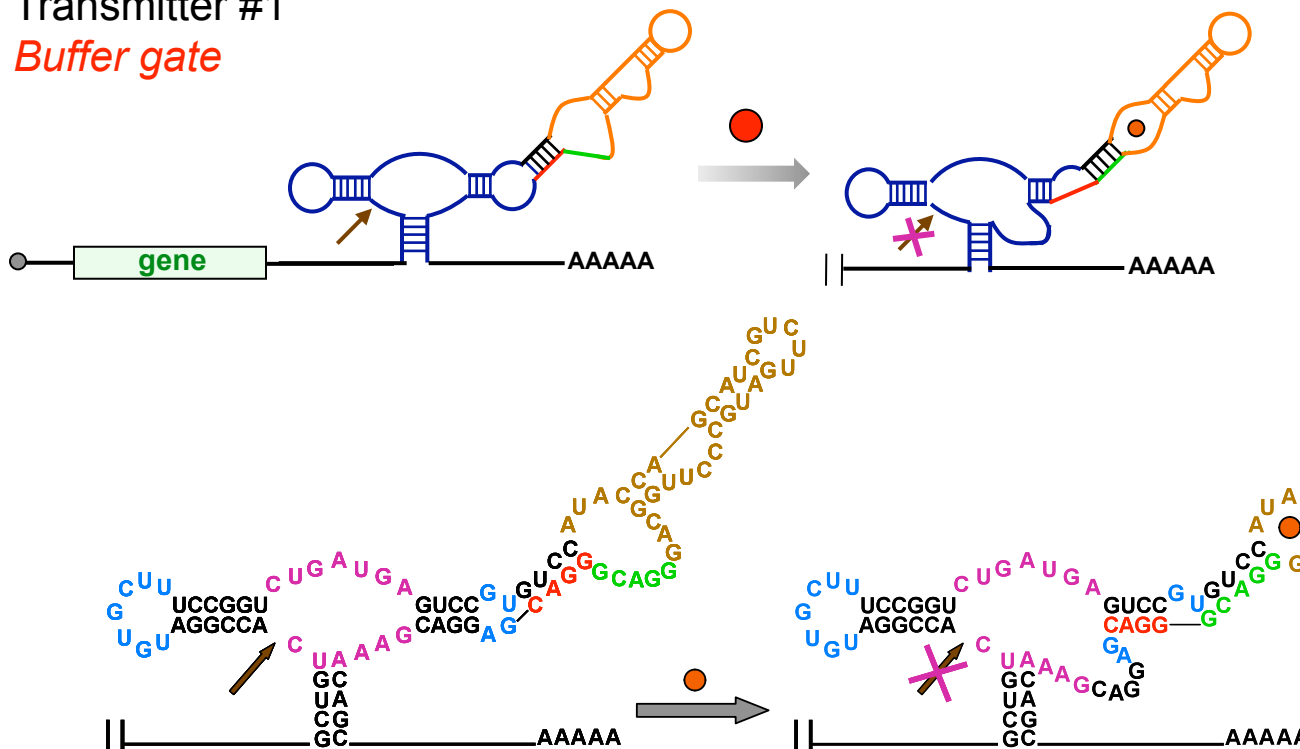
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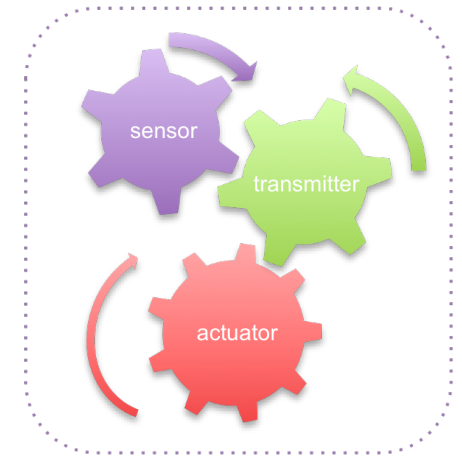
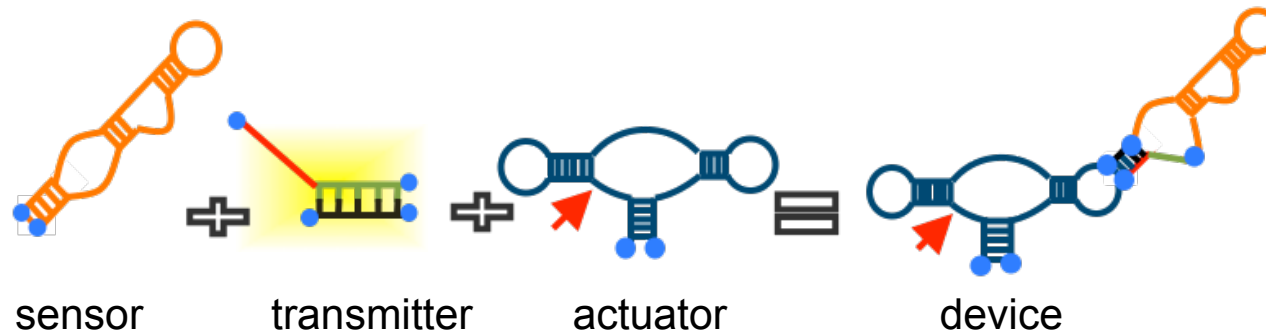
A platform for programming device function



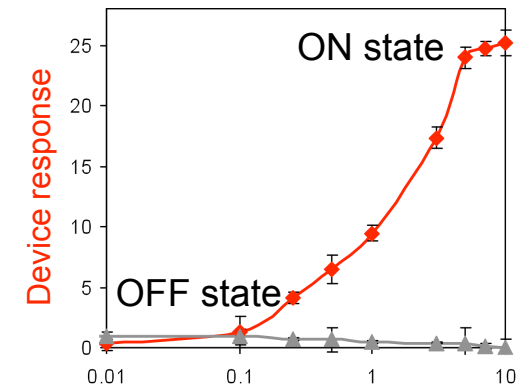
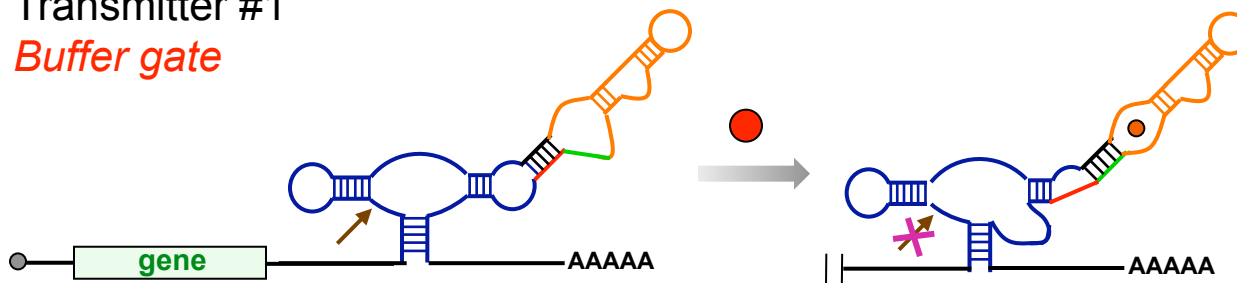
Transmitter #1
Buffer gate



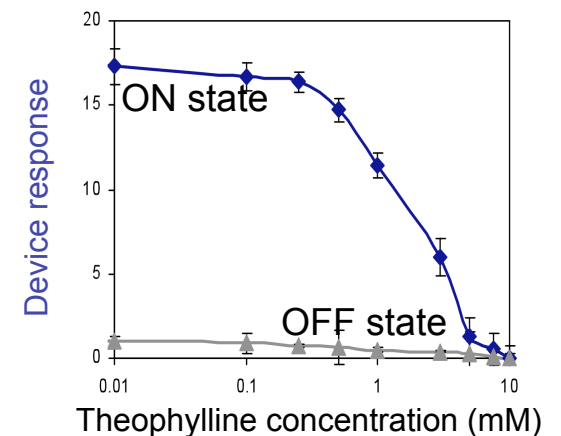
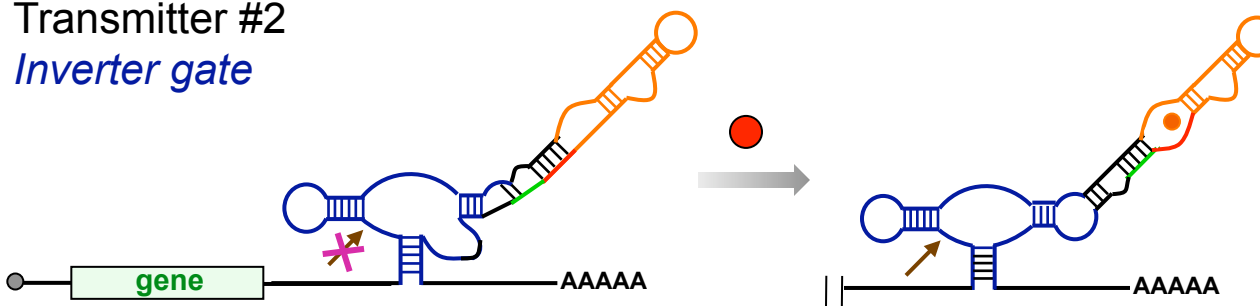
A platform for programming device function



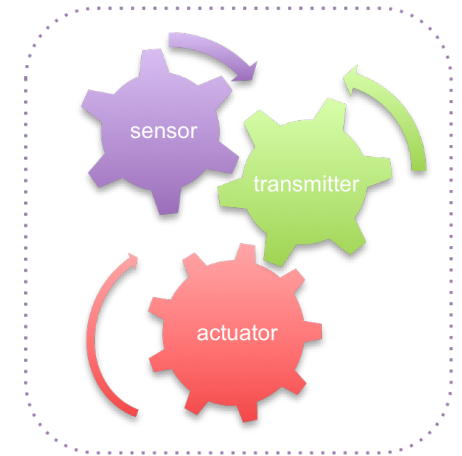
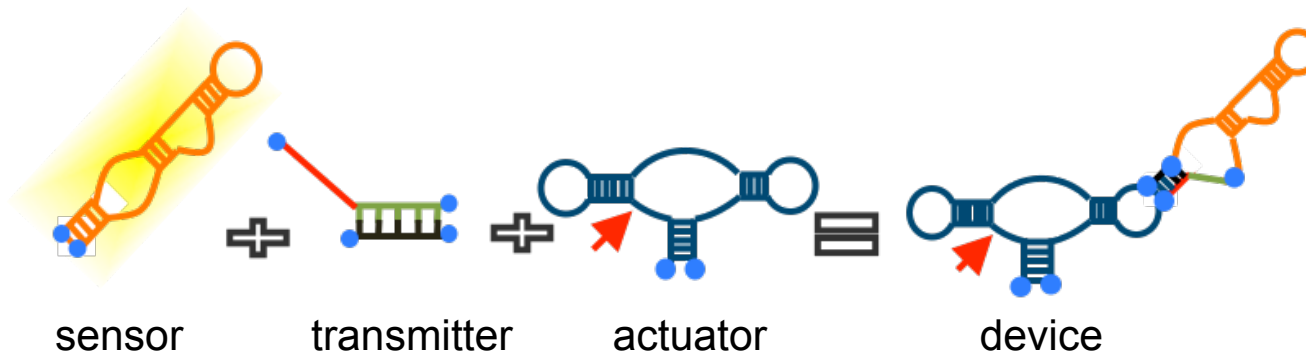
Transmitter #1
Buffer gate



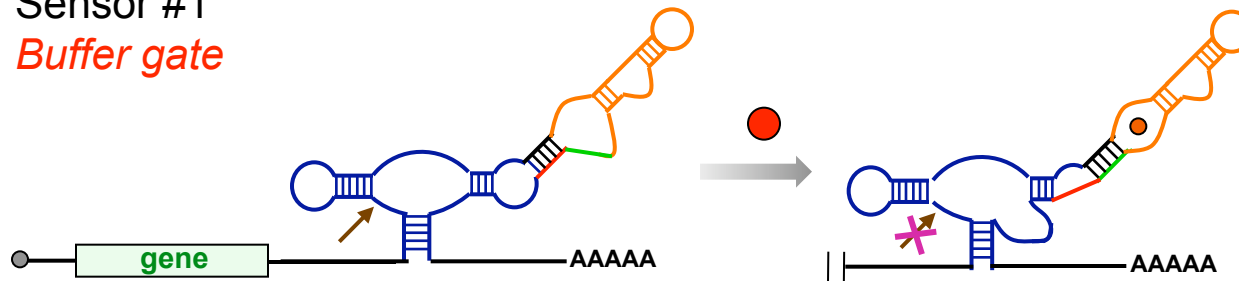
Transmitter #2
Inverter gate



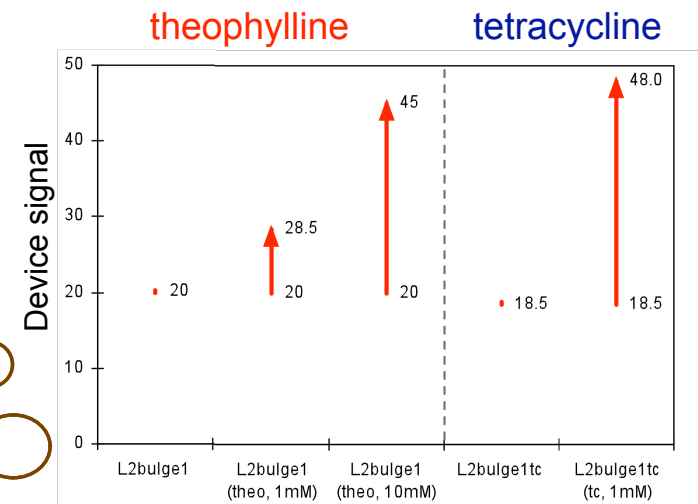
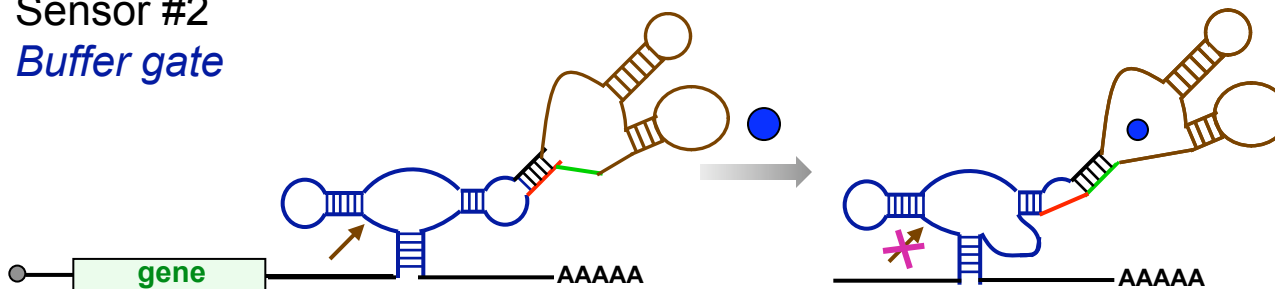
A platform for programming device inputs



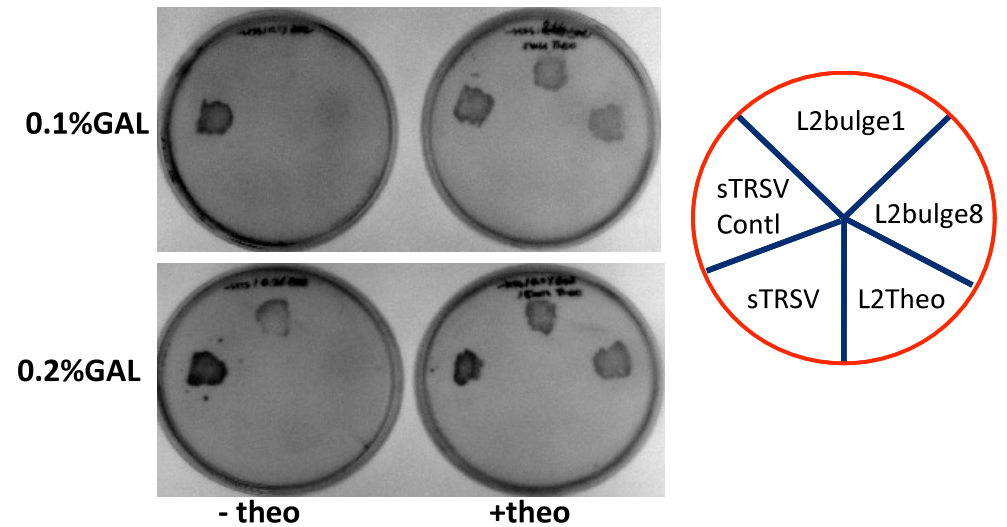
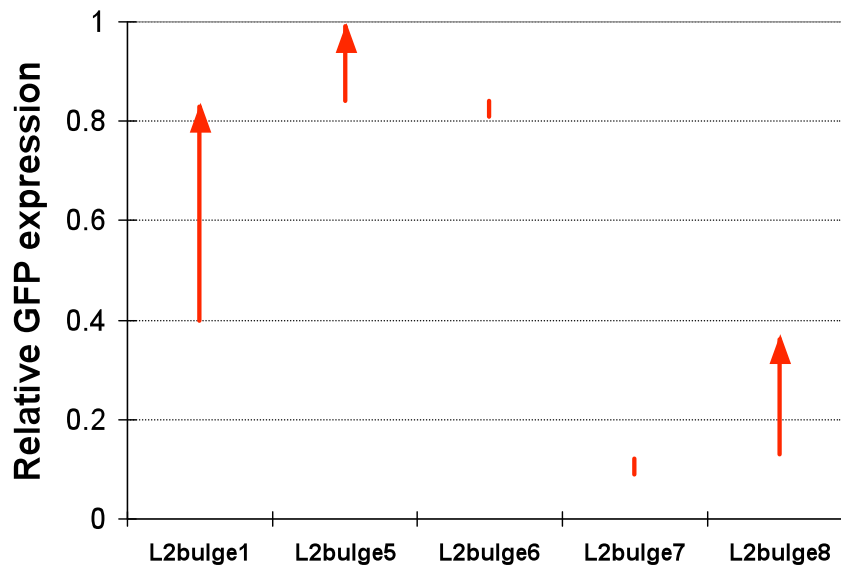
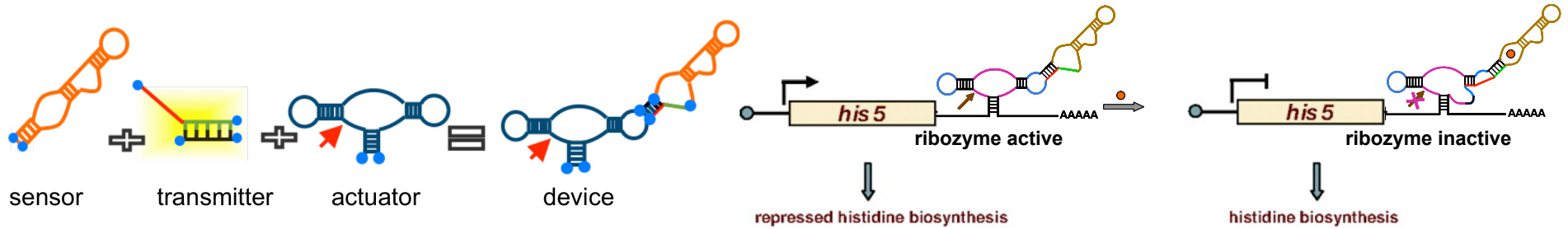
Sensor #1
Buffer gate



Sensor #2
Buffer gate



A platform for programming response properties

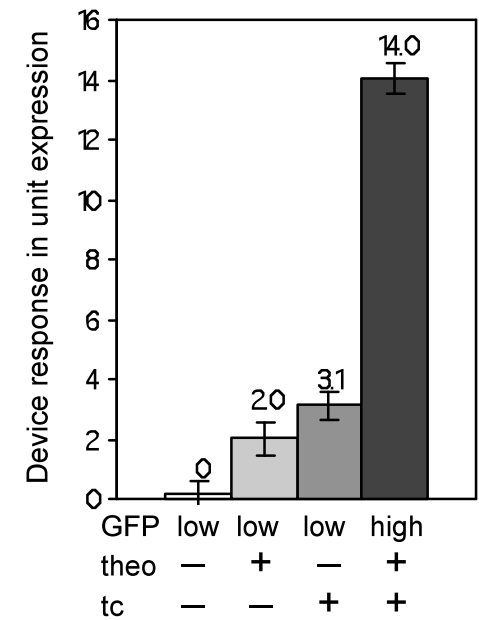
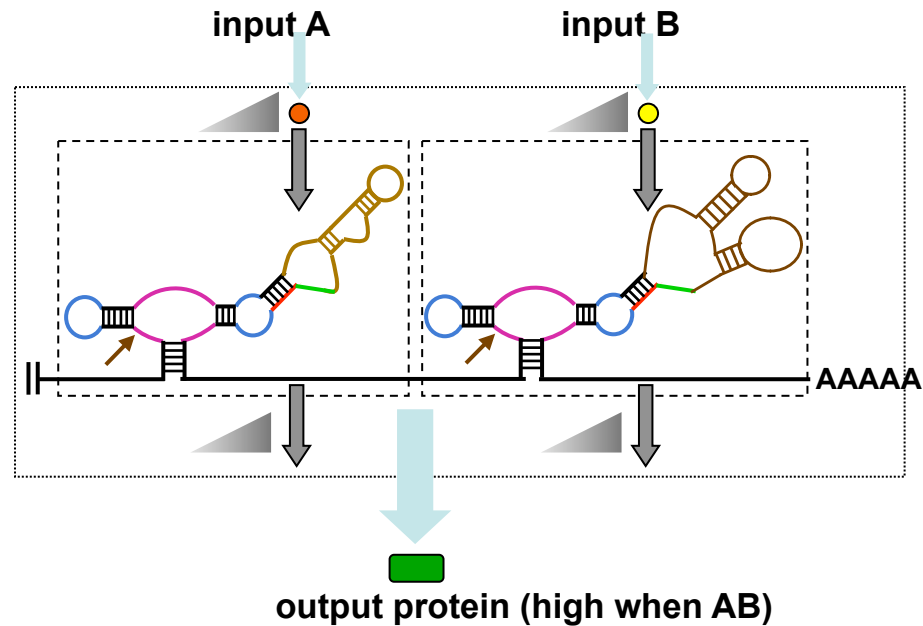


Logic gate devices

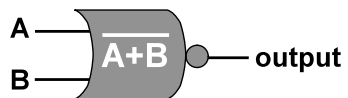
AND gate



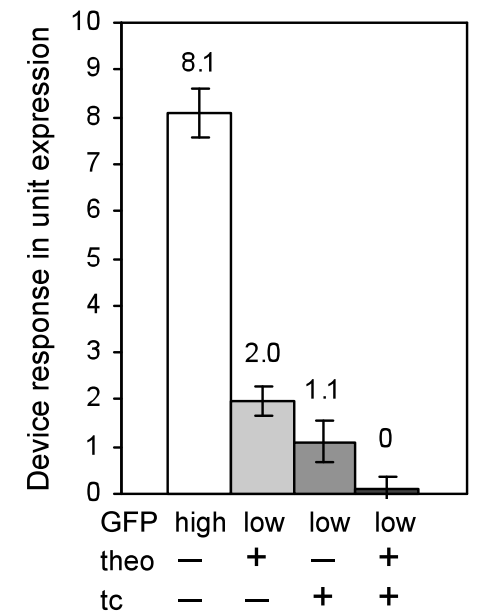
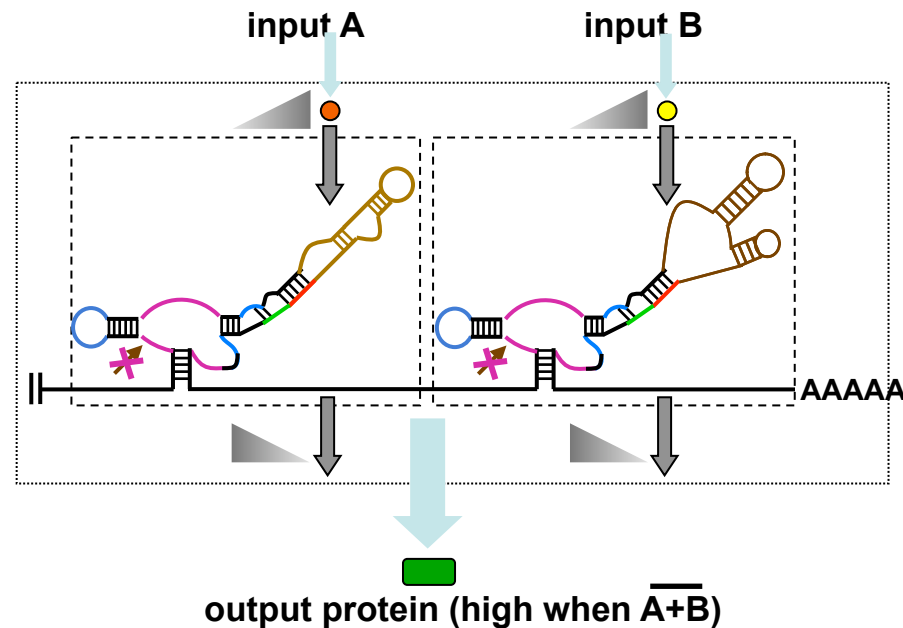
A	B	output
theo	tc	GFP
0	0	0
0	1	0
1	0	0
1	1	1



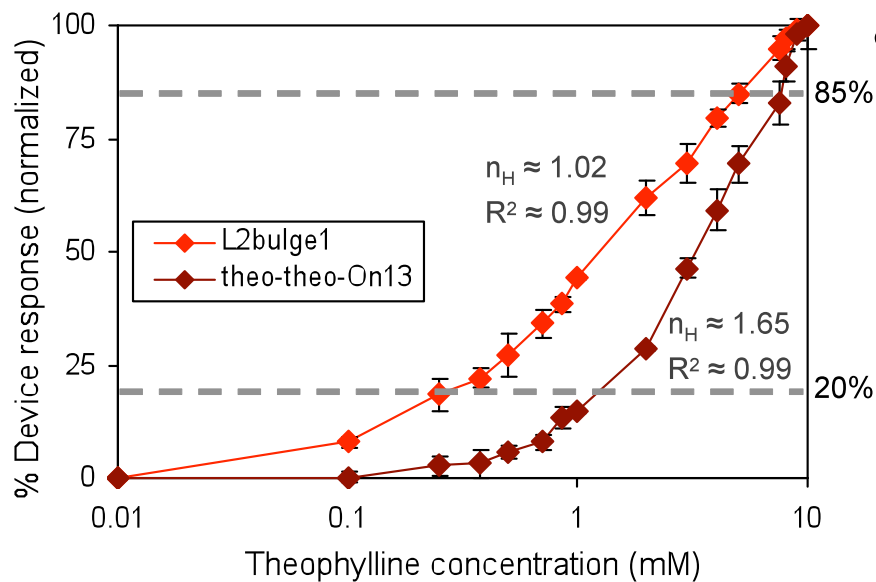
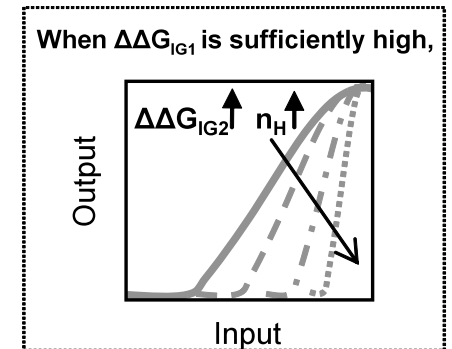
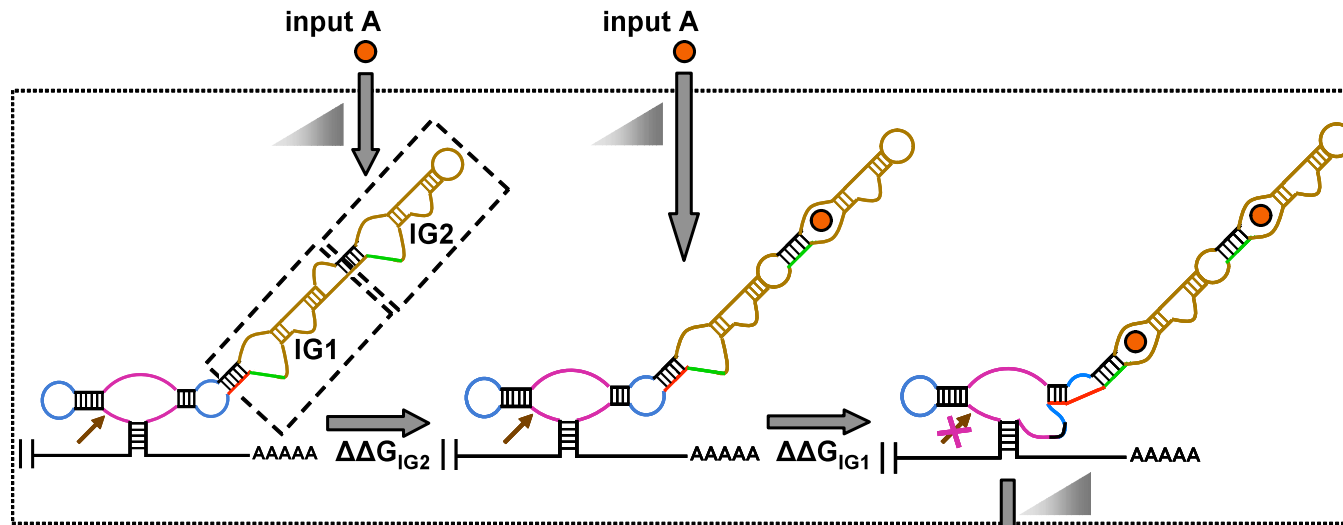
NOR gate



A	B	output
theo	tc	GFP
0	0	1
0	1	0
1	0	0
1	1	0



Single gain device – programmed cooperativity



output protein (high when AA)

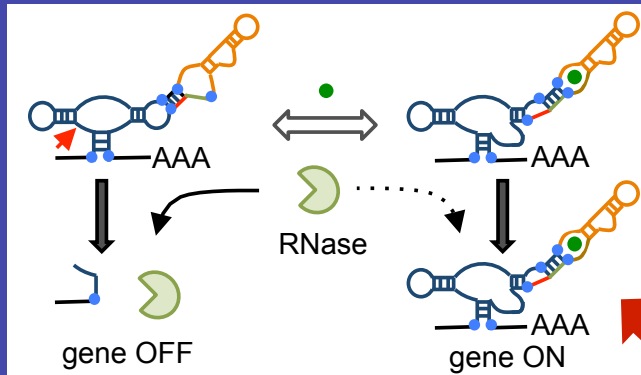
theo-theo-On13

Hill coefficient = 1.65

L2bulge1

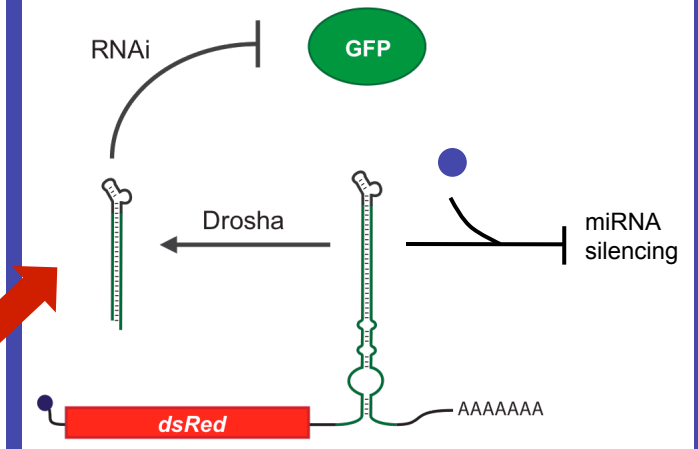
Hill coefficient = 1.02

Ribozyme Cleavage



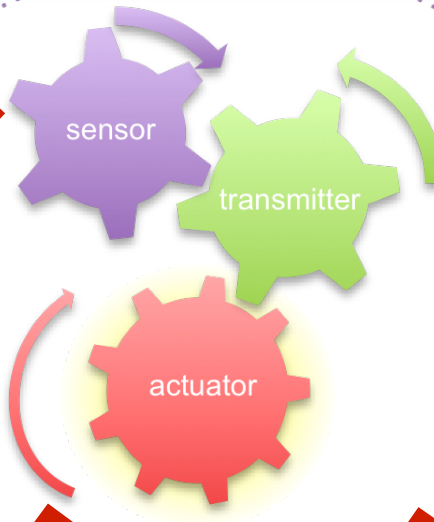
Win MN, Smolke CD. 2007. *PNAS*. 104: 14283-8
 Win MN, Smolke CD. 2008. *Science*. 322: 456-60

miRNA-Mediated Silencing

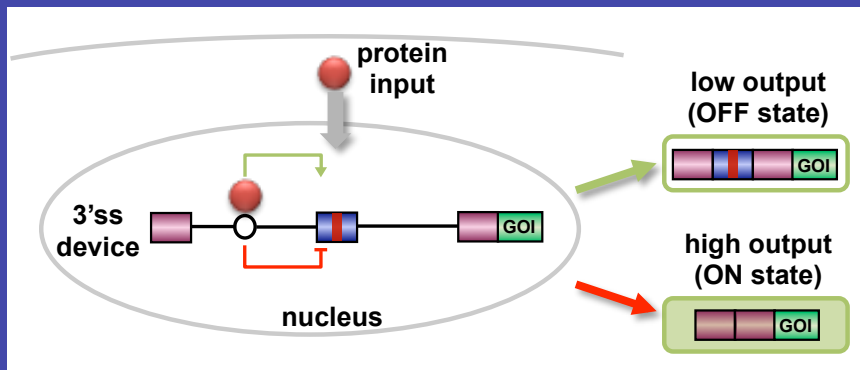


Beisel CL, et al. 2010. *In review*

RNA devices

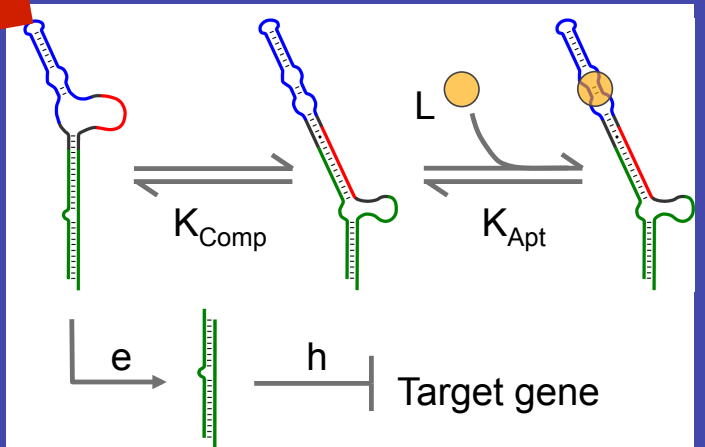


Alternative Splicing



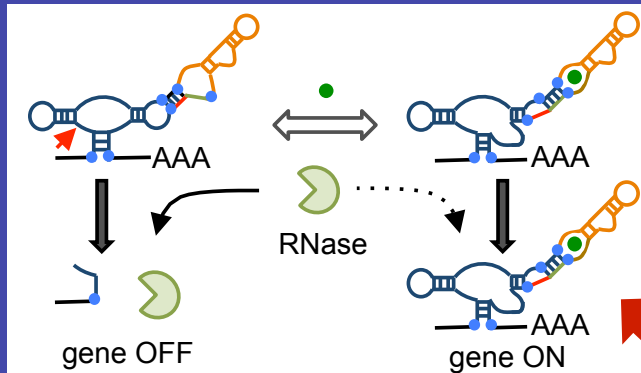
Culler SJ, Hoff KG, Smolke CD. 2010. *In review*

shRNA-Mediated Silencing

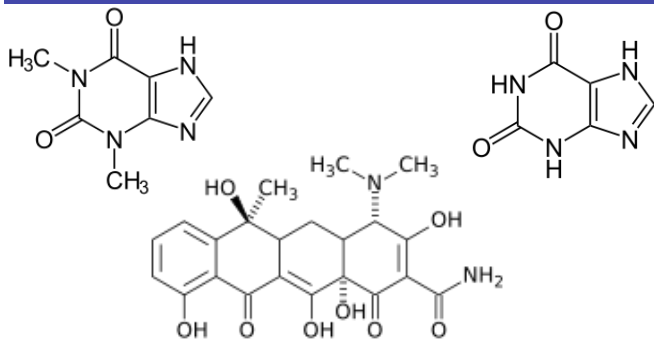


Beisel CL, et al. 2008. *Mol Sys Biol*. 4: 224

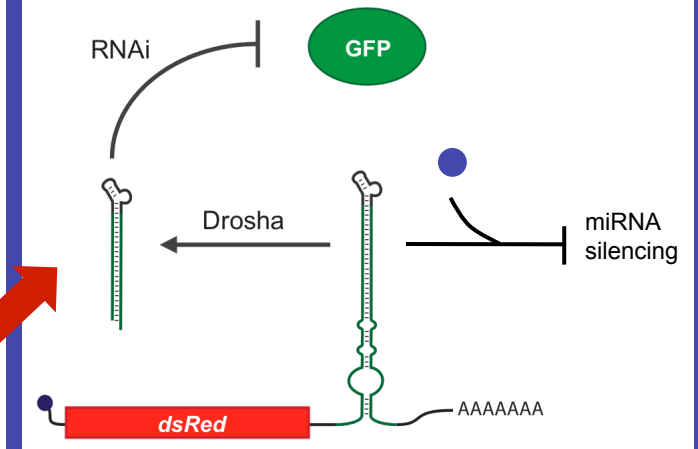
Ribozyme Cleavage



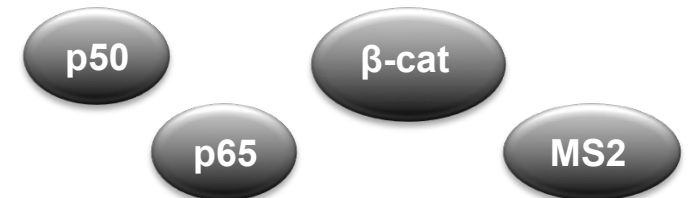
Win MN, Smolke CD. 2007. *PNAS*. 104: 14283-8
Win MN, Smolke CD. 2008. *Science*. 322: 456-60



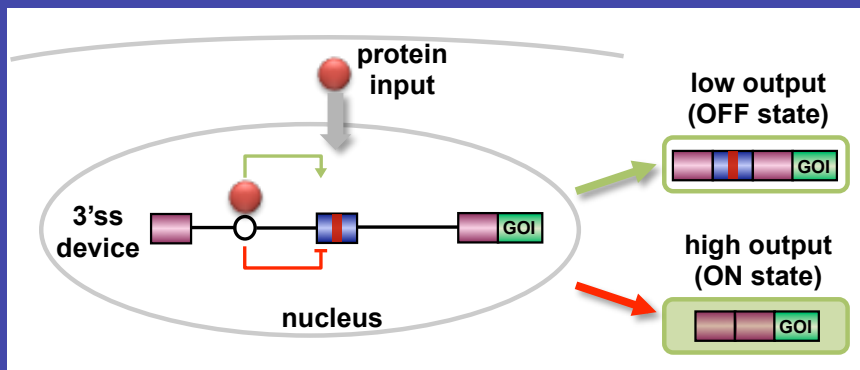
miRNA-Mediated Silencing



Beisel CL, et al. 2010. *In review*

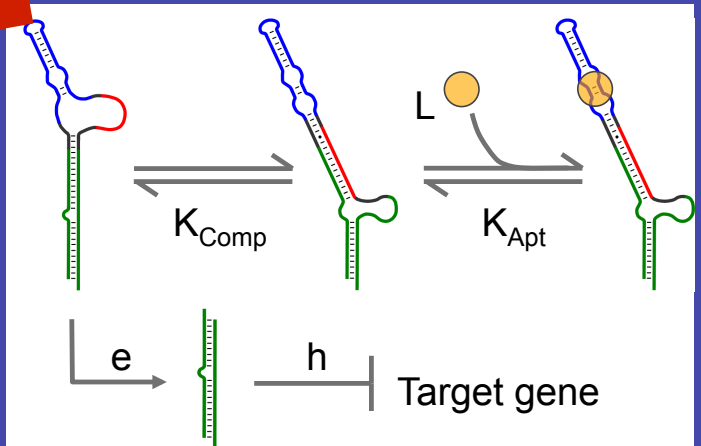


Alternative Splicing



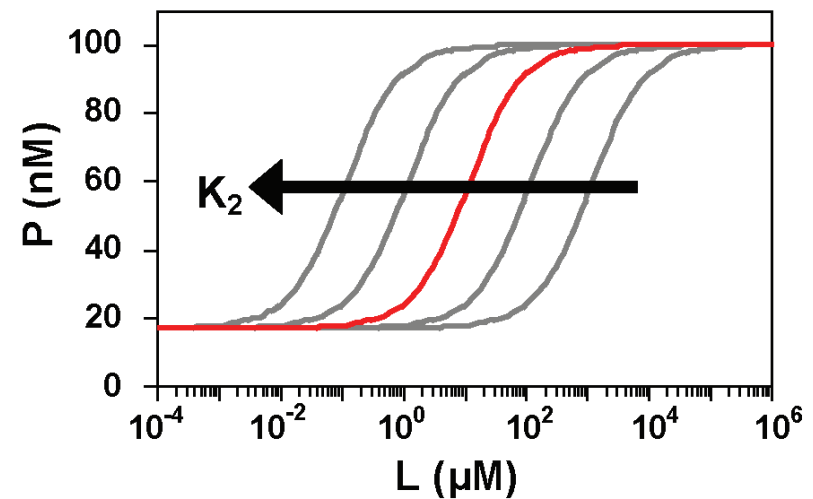
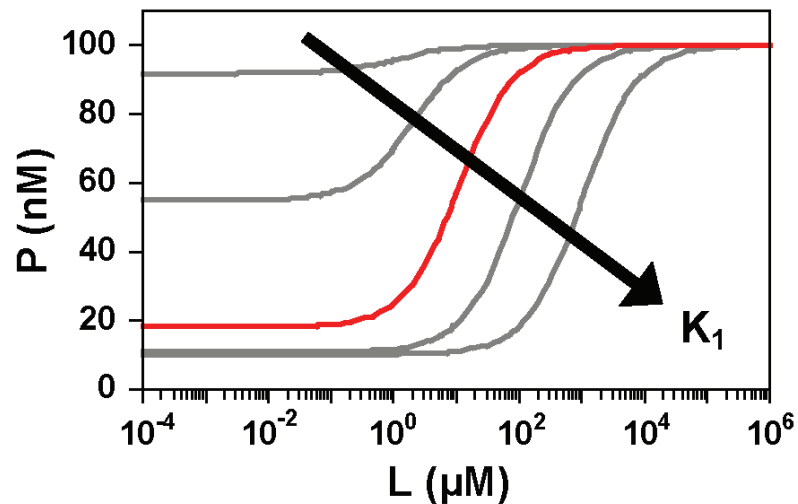
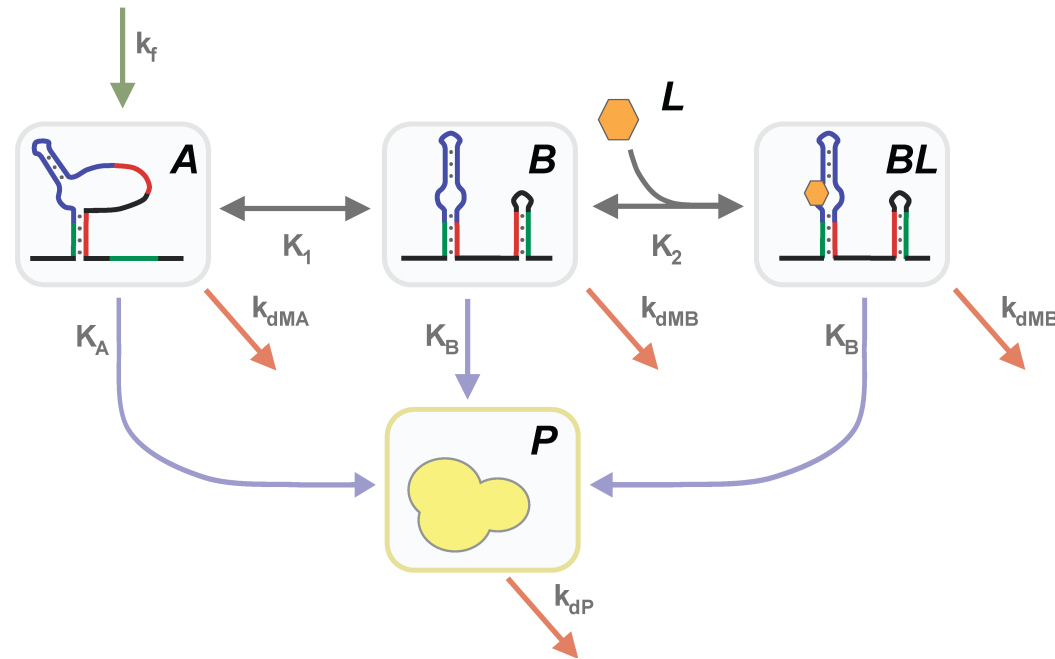
Culler SJ, Hoff KG, Smolke CD. 2010. *In review*

shRNA-Mediated Silencing

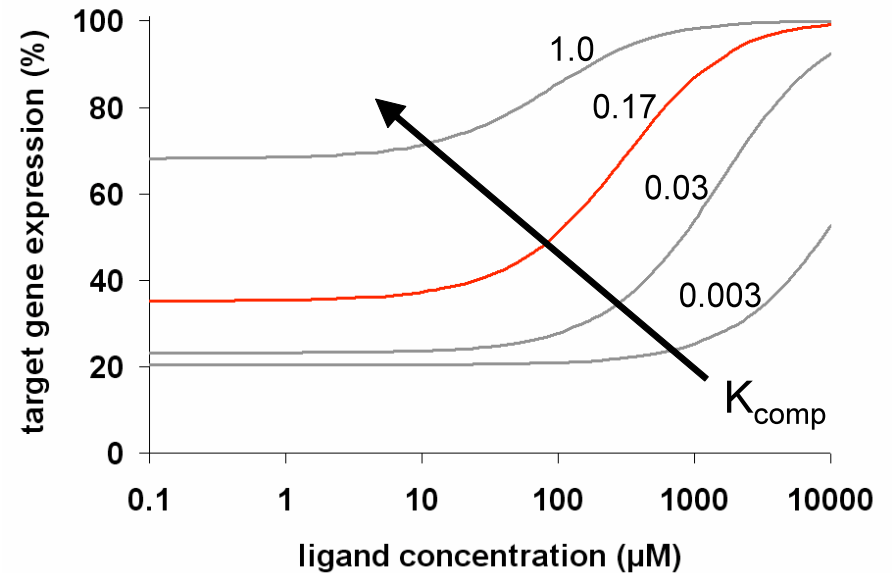
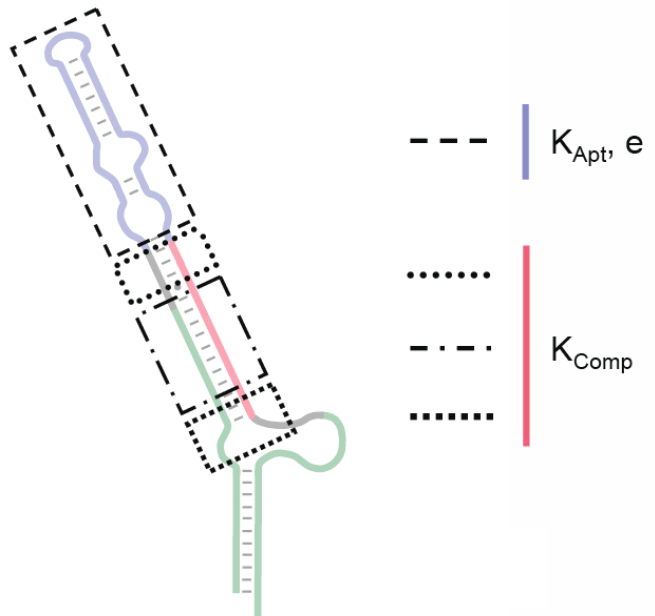


Beisel CL, et al. 2008. *Mol Sys Biol*. 4: 224

Computational models can predict parameter effects

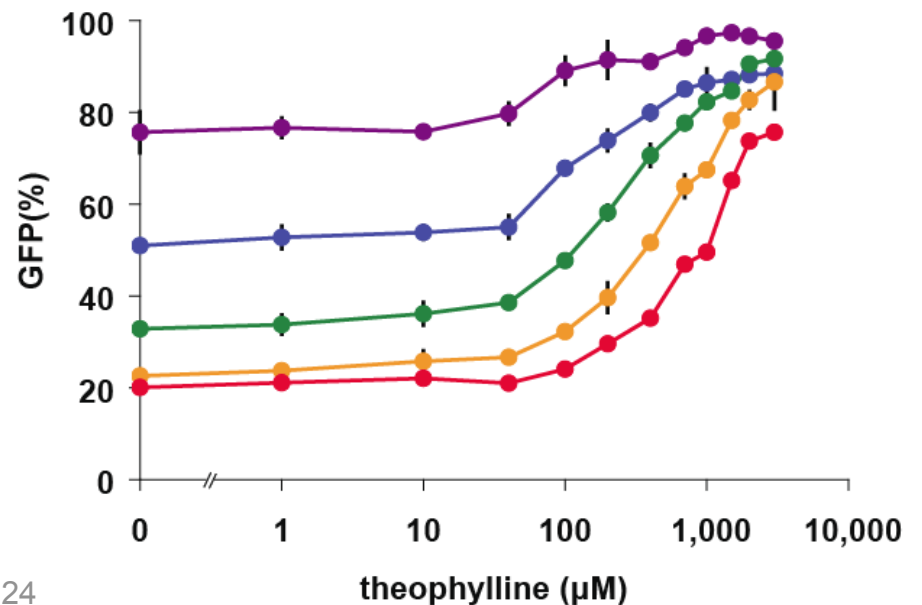
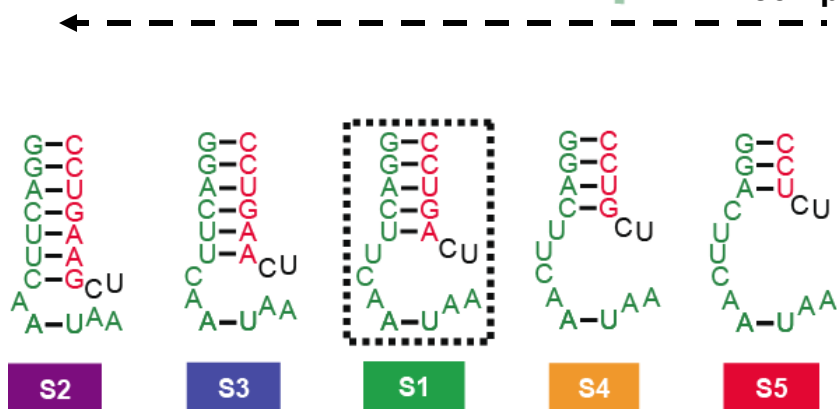


Models predict transmitter tuning strategies

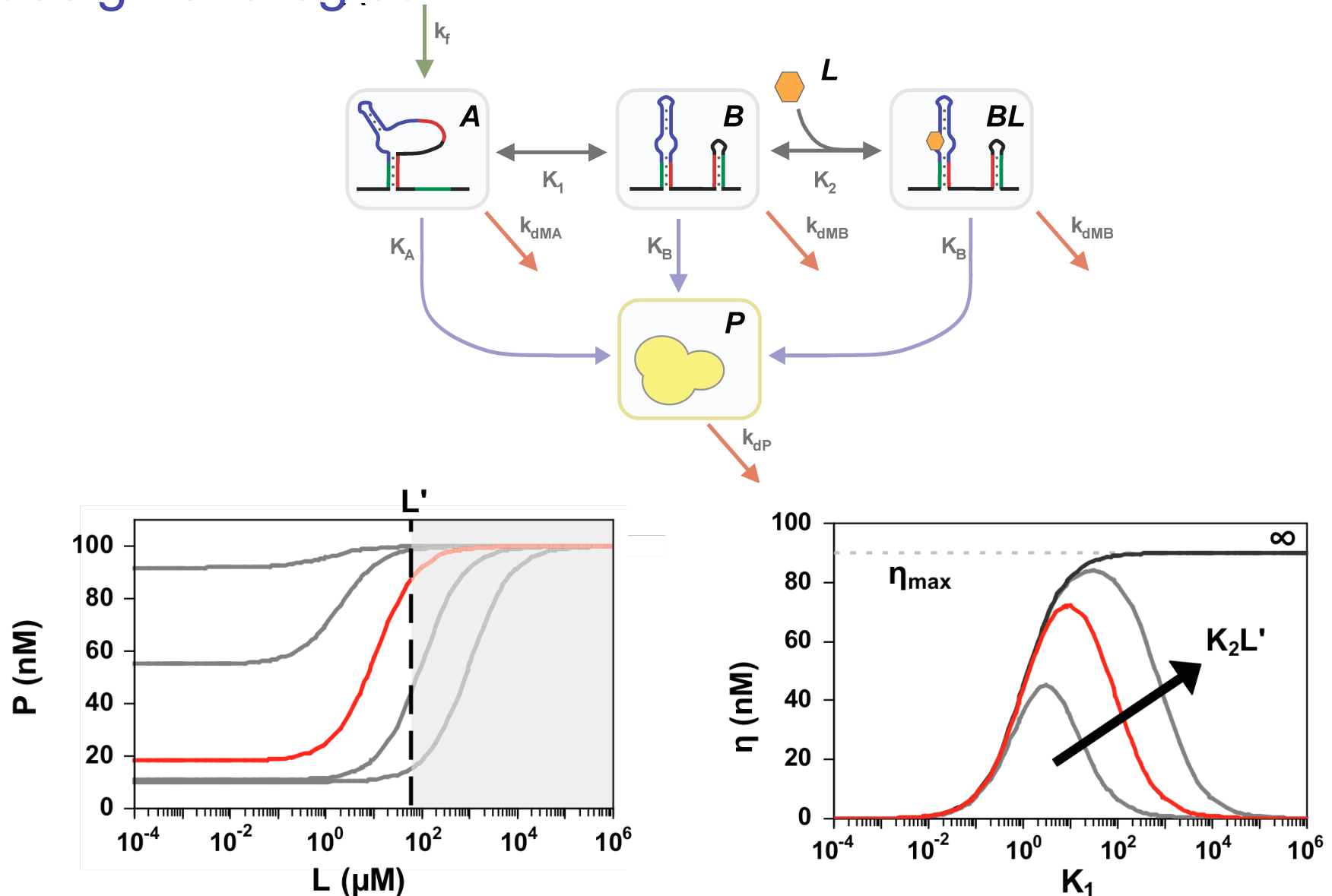


Construction and experimental validation

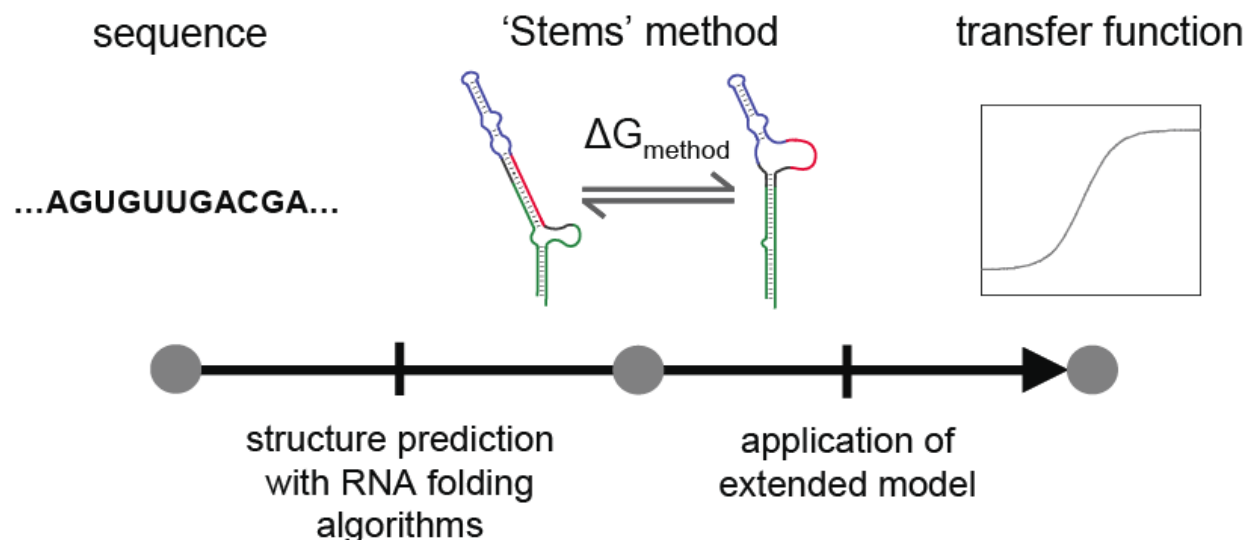
Competing strand 3' length (K_{comp})



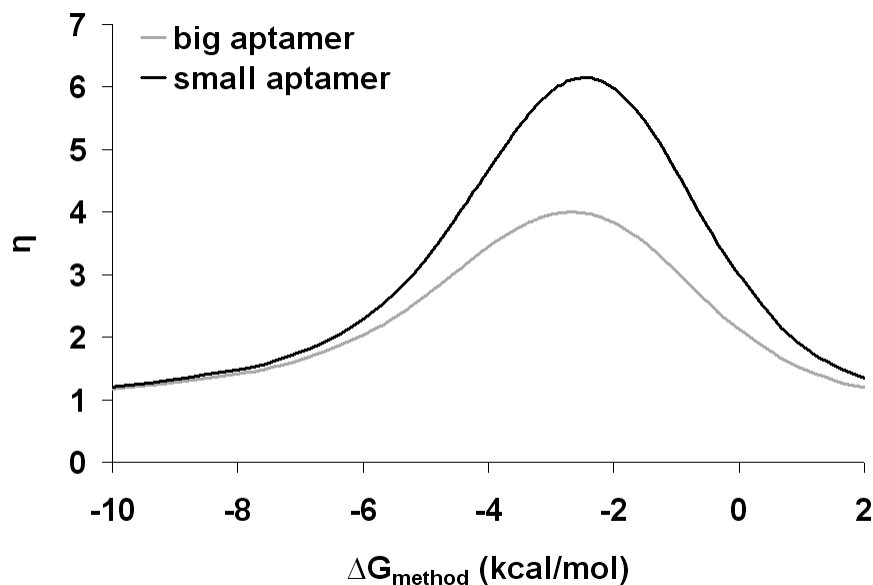
System restrictions lead to optimization in RNA device design strategies



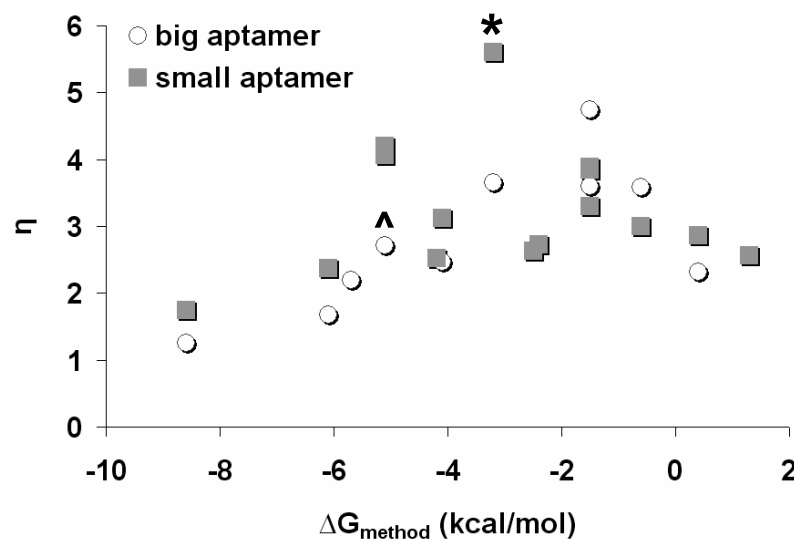
A framework for forward engineering device response



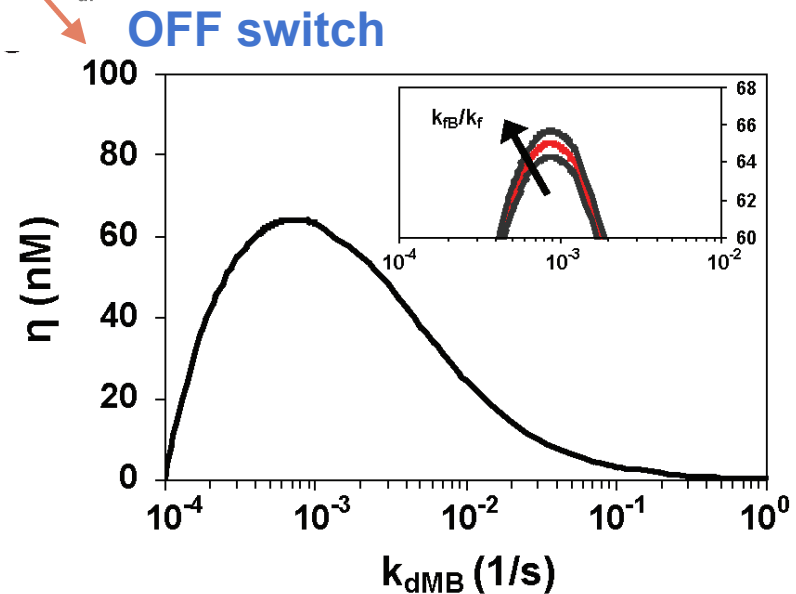
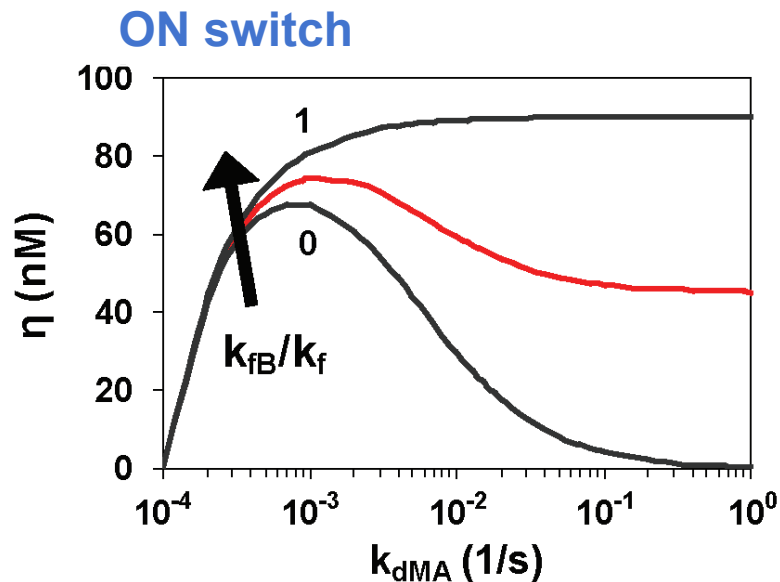
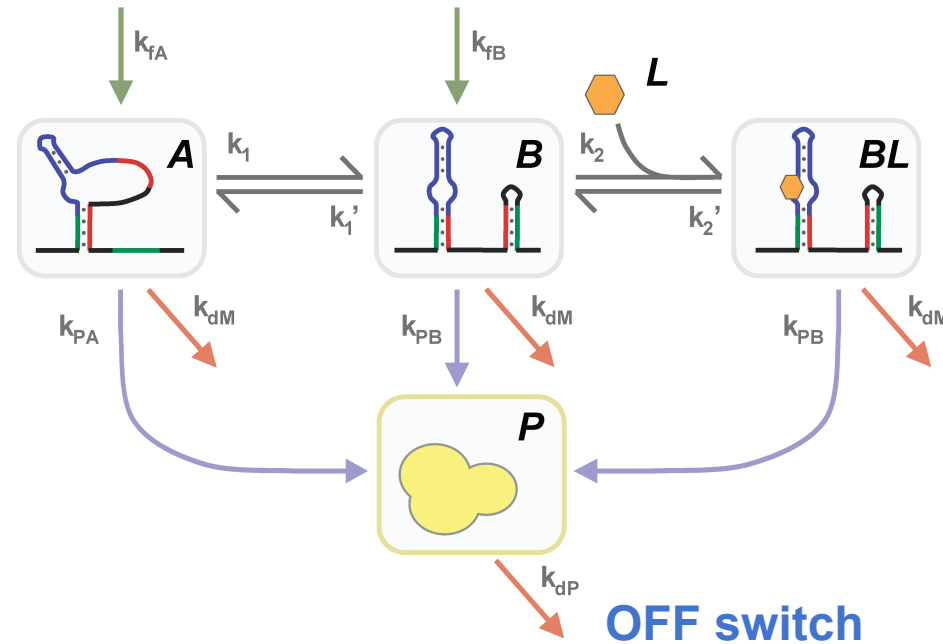
Model predictions



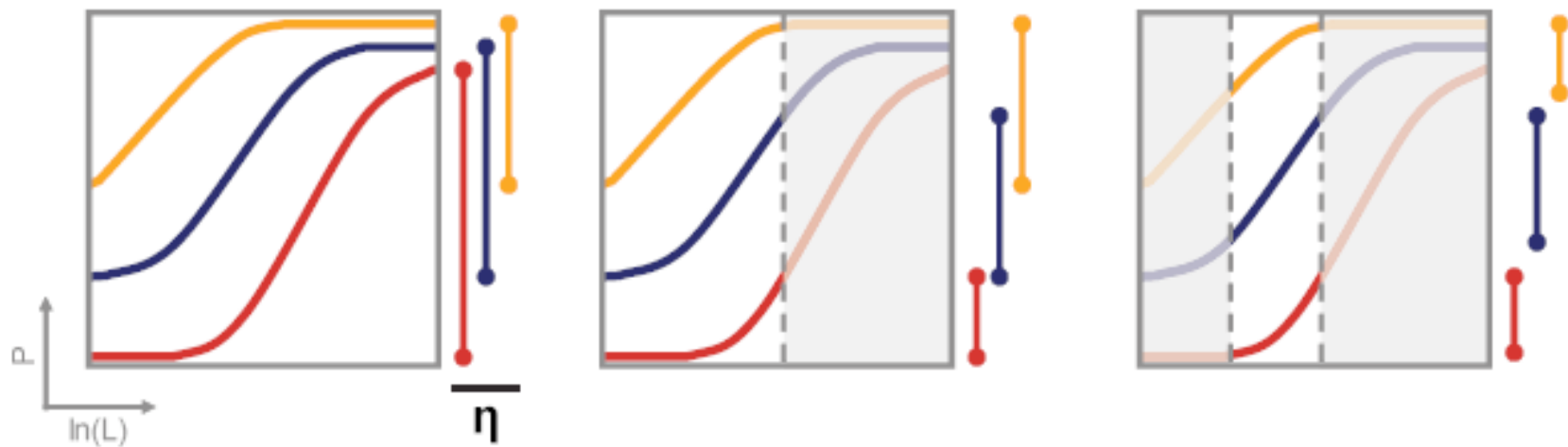
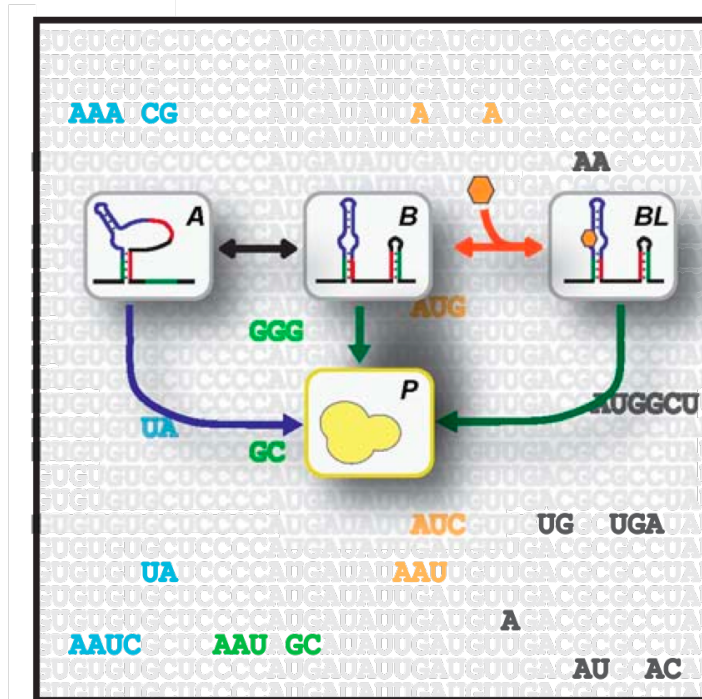
Experimental results



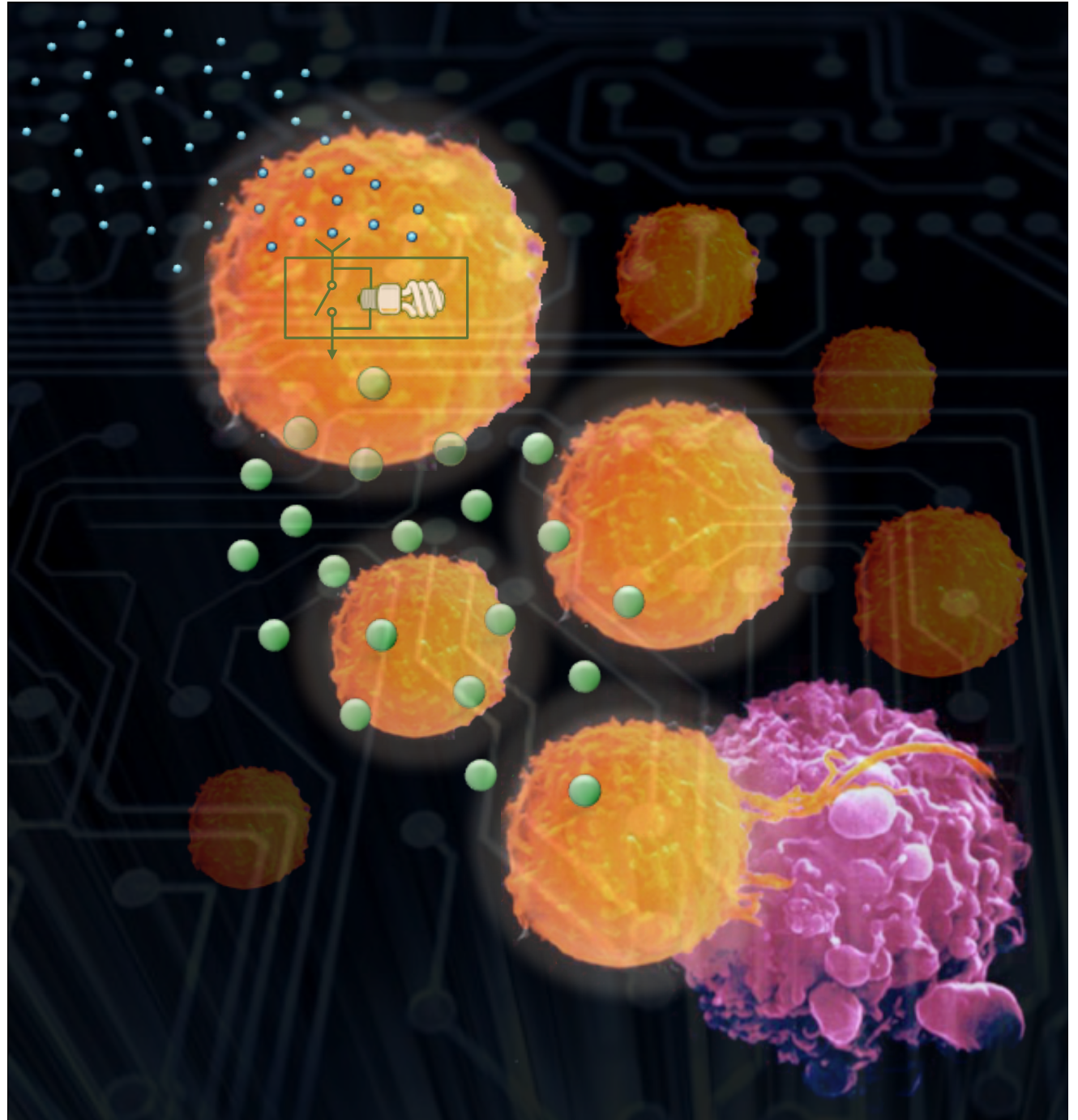
Irreversible rates can play an important role in RNA device performance



Quantitative tuning of device response

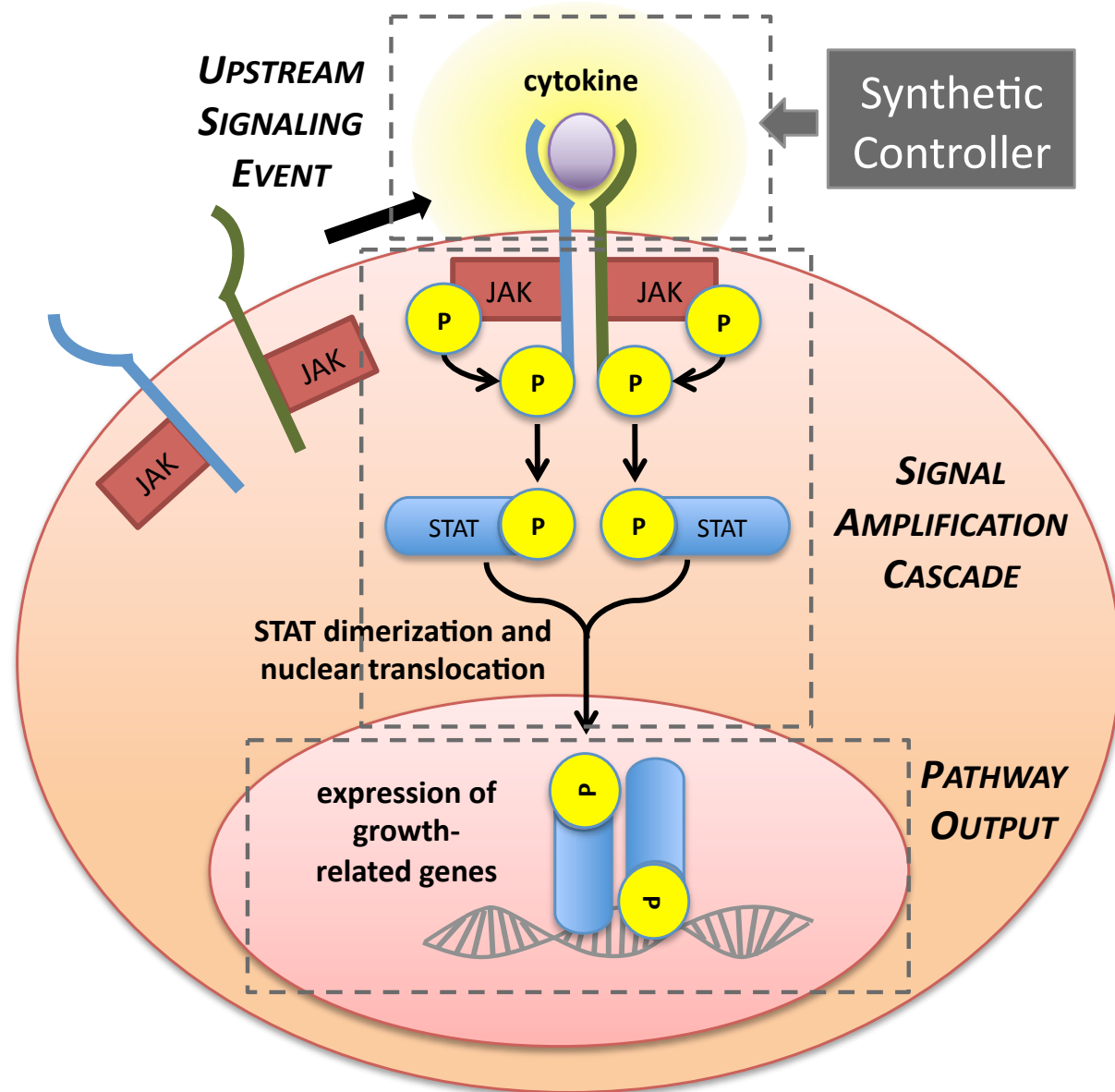


Advancing cell-based therapies

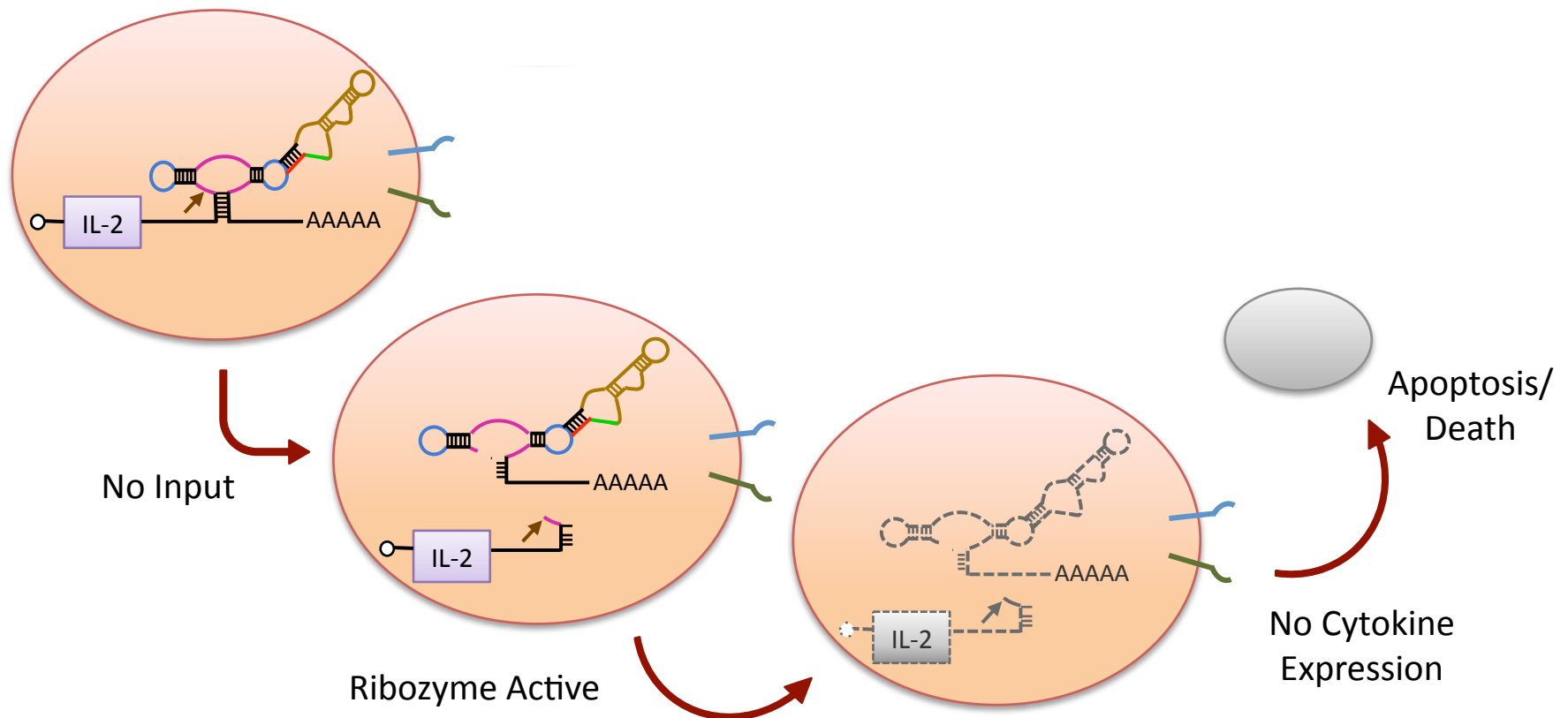


Chen YY, Jensen MC, Smolke CD. 2010. *PNAS*. 107: 8531-6

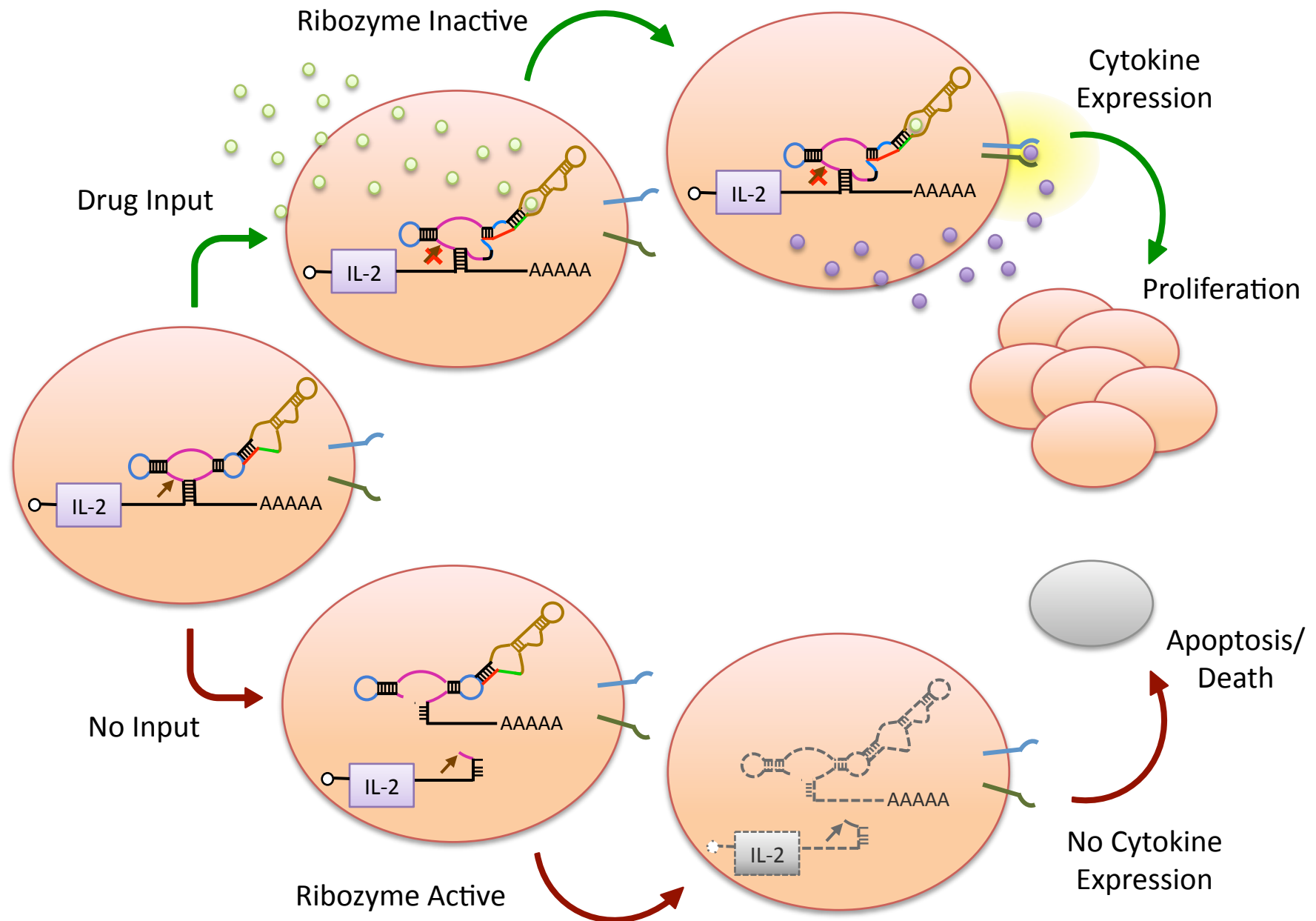
T-cell proliferation pathway



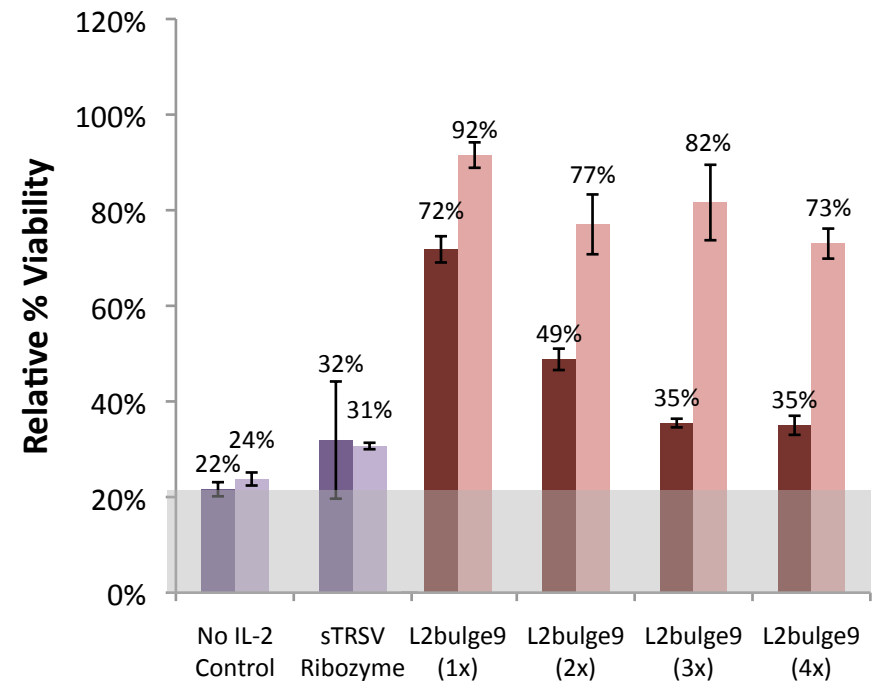
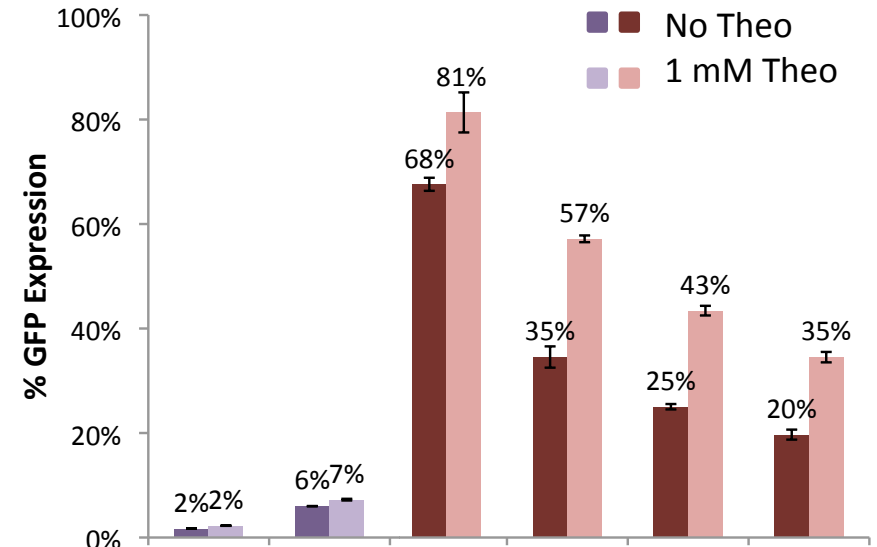
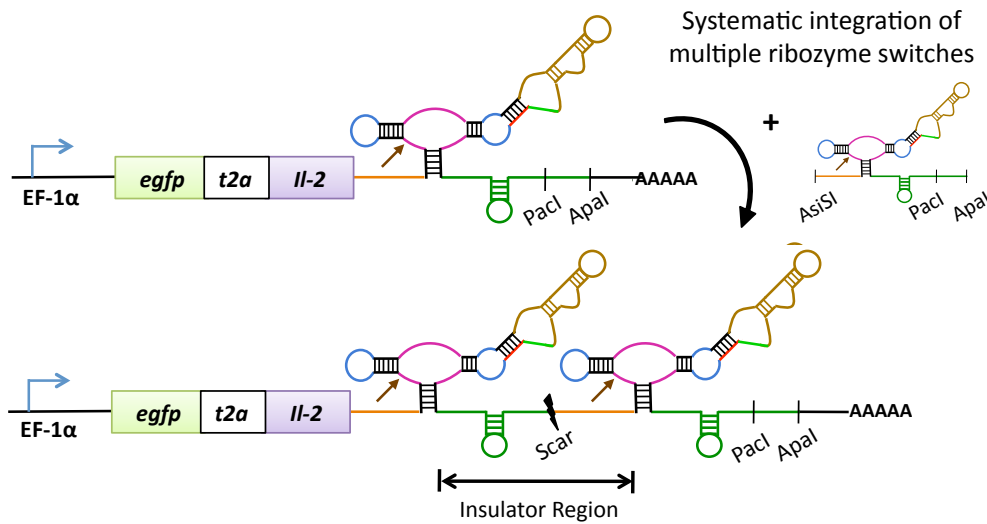
A synthetic T-cell regulatory network



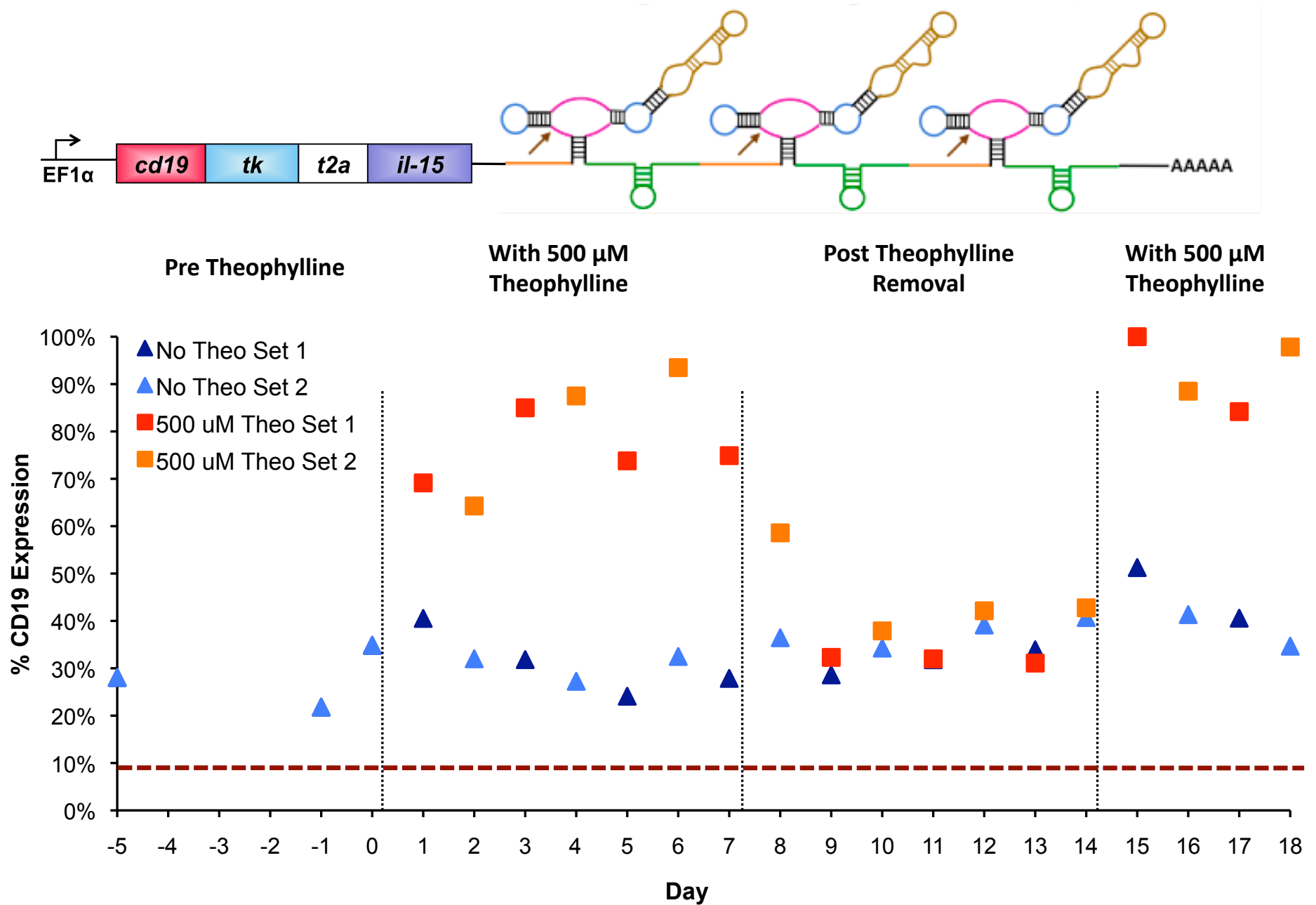
A synthetic T-cell regulatory network



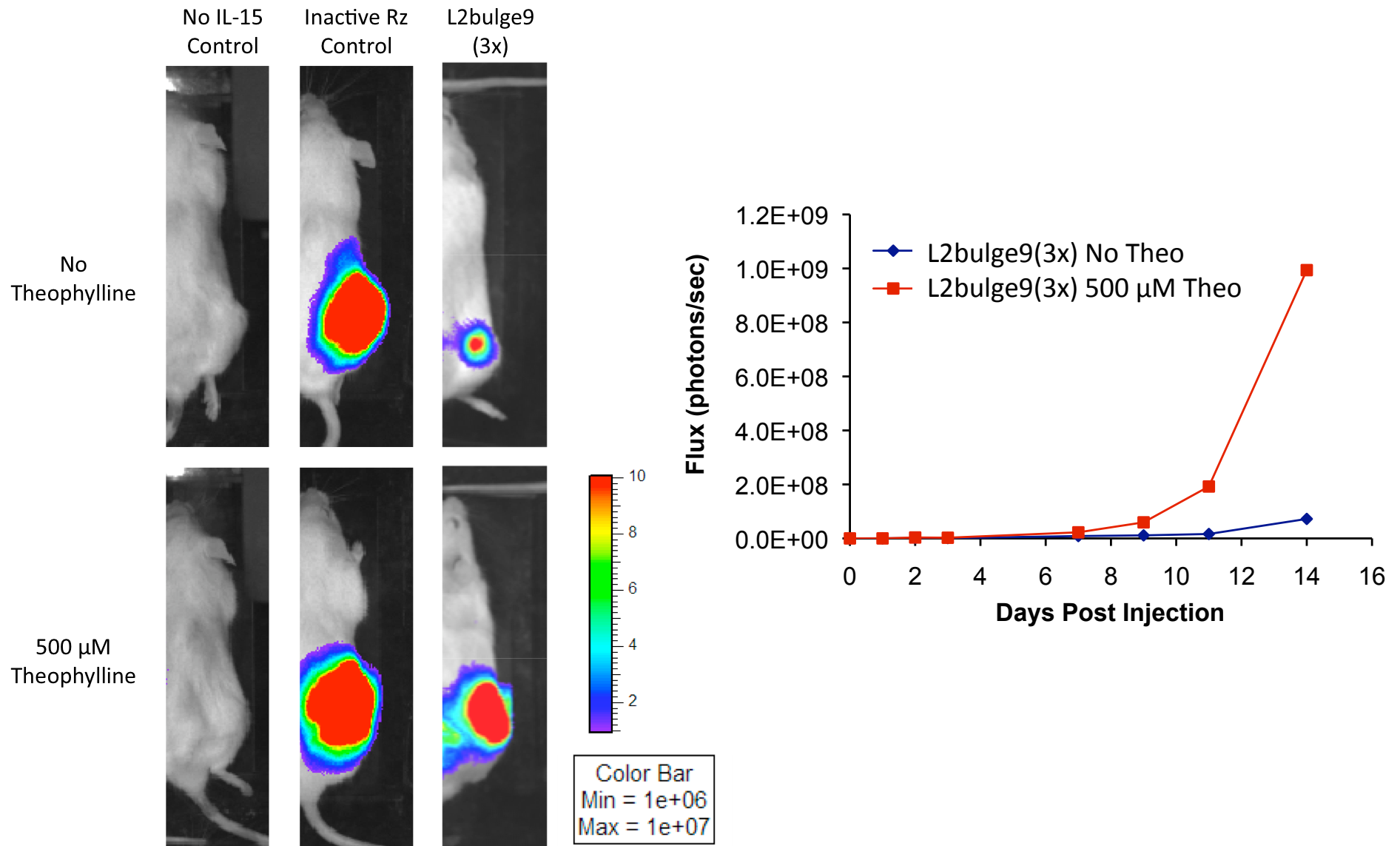
Rapid prototyping and optimization of a T-cell proliferation control system



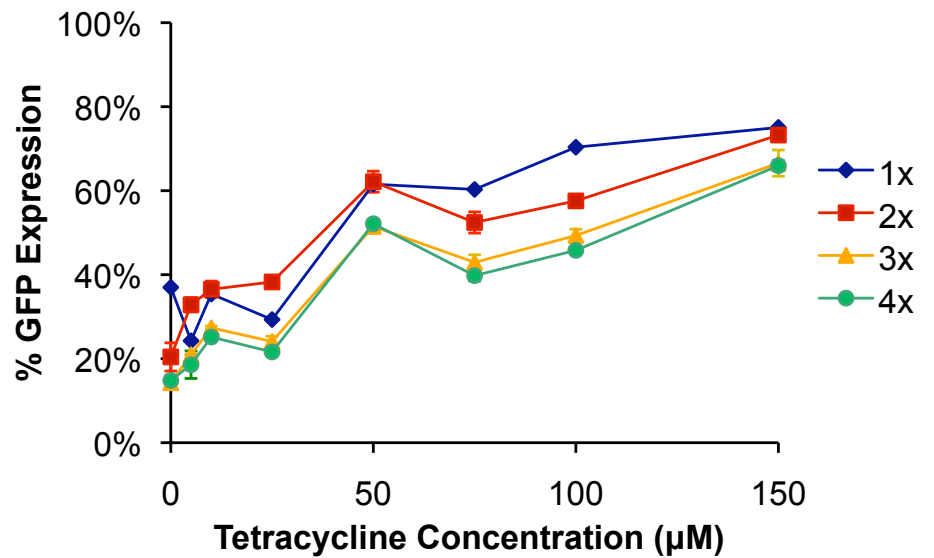
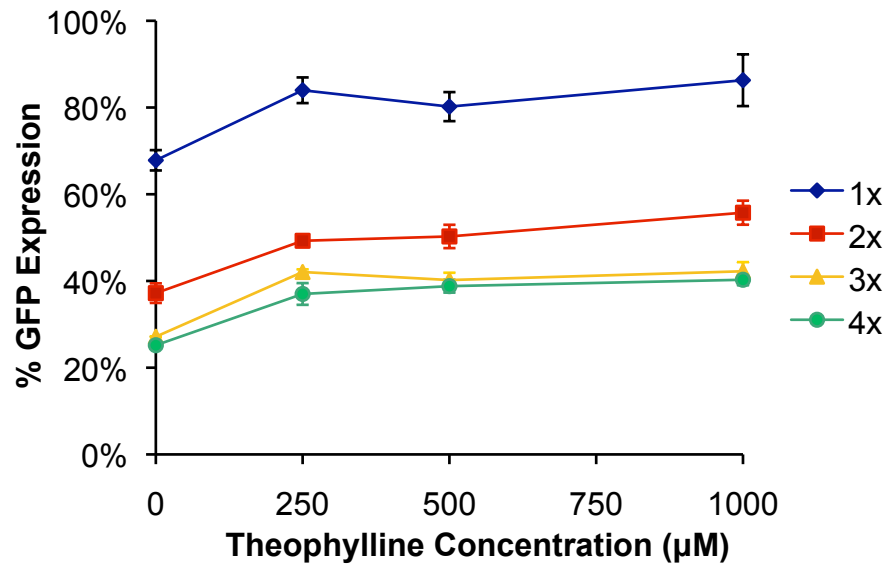
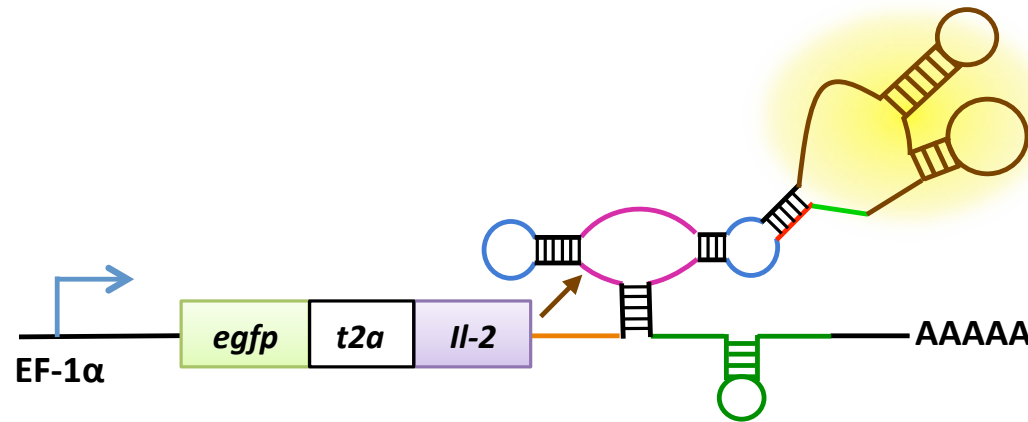
Stable cell lines show dynamic response to input



RNA devices regulate T-cell proliferation *in vivo*



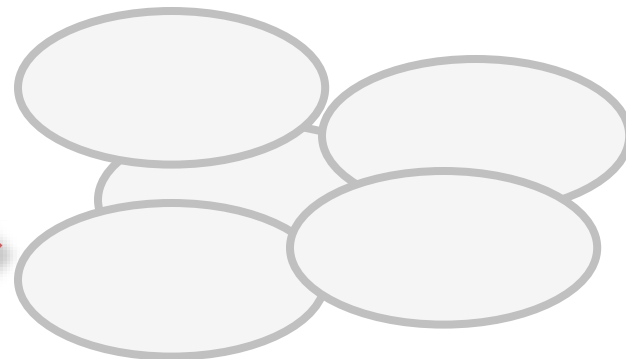
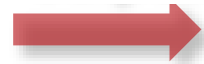
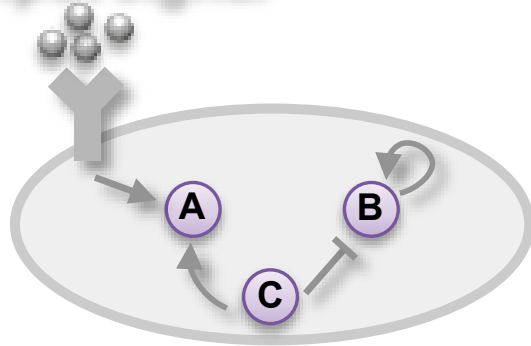
Choice of drug-sensor pair affects device performance



Moving between exogenous and embedded control

Native Network

input signal

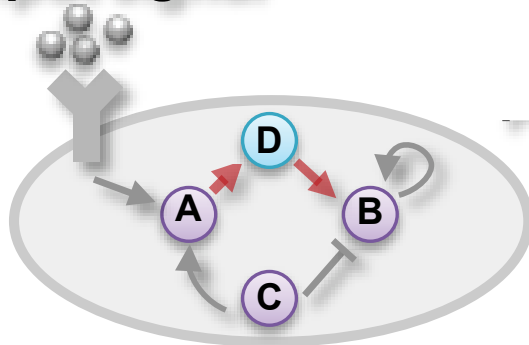


proliferation

fate 1

Altered Network

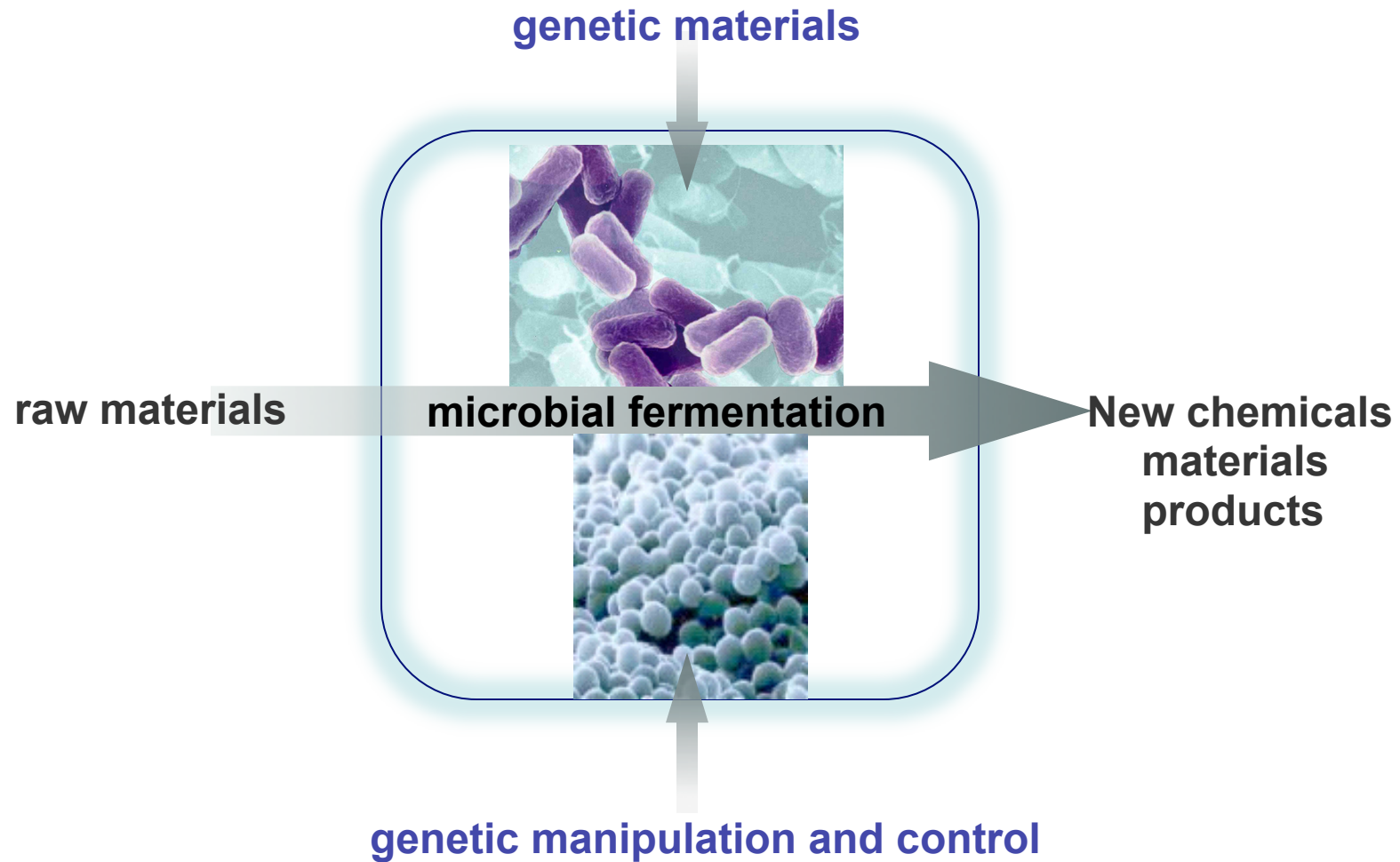
input signal



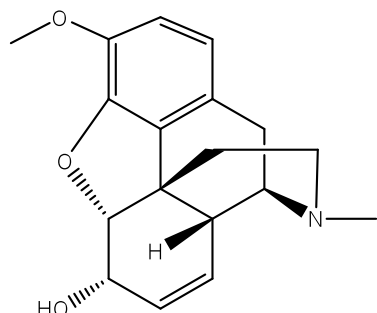
cell death

fate 2

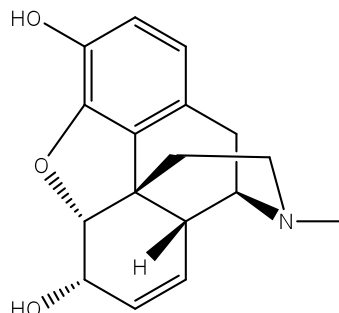
Microbial factories



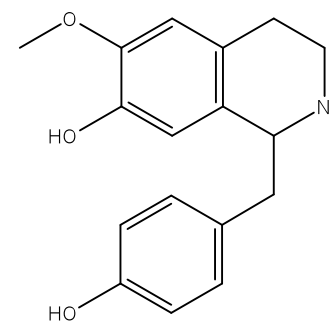
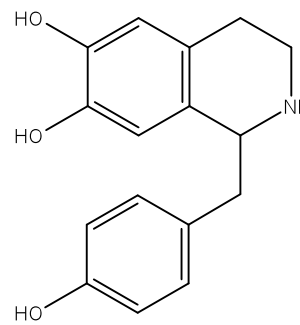
Benzylisoquinoline Alkaloids (BIAs) are an important class of natural products



Codeine

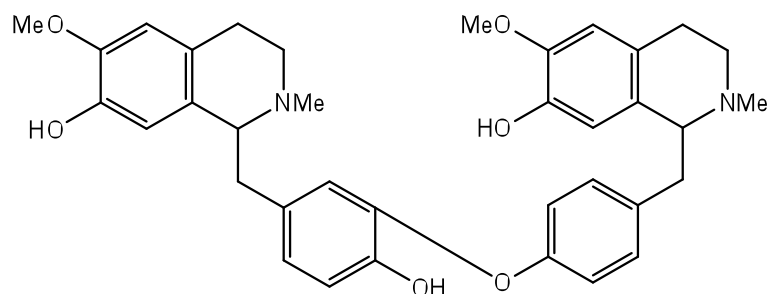
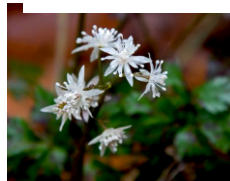


Morphine

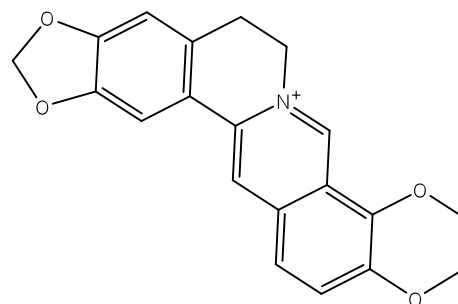


Norcoclaurine, Coclaurine

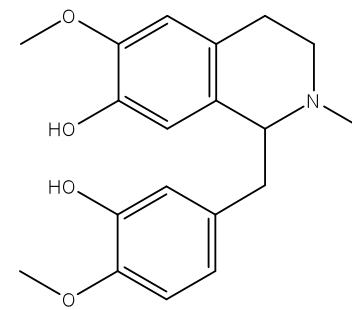
Over 2,500 known structures



bis-BIAs (Berbamunine)
Antimicrobial, anticancer

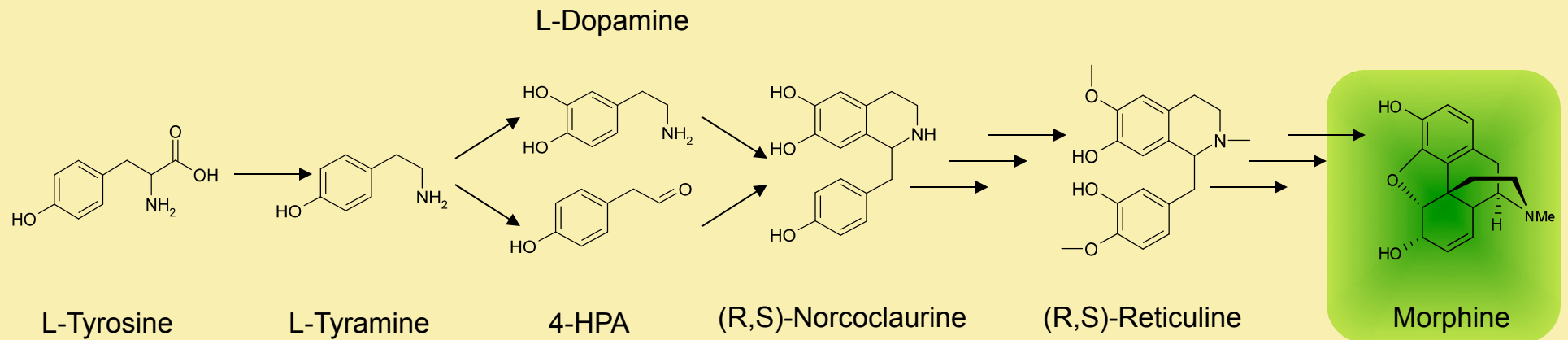


Berberine
Antimicrobial, antioxidant



Reticuline
stimulates hair growth

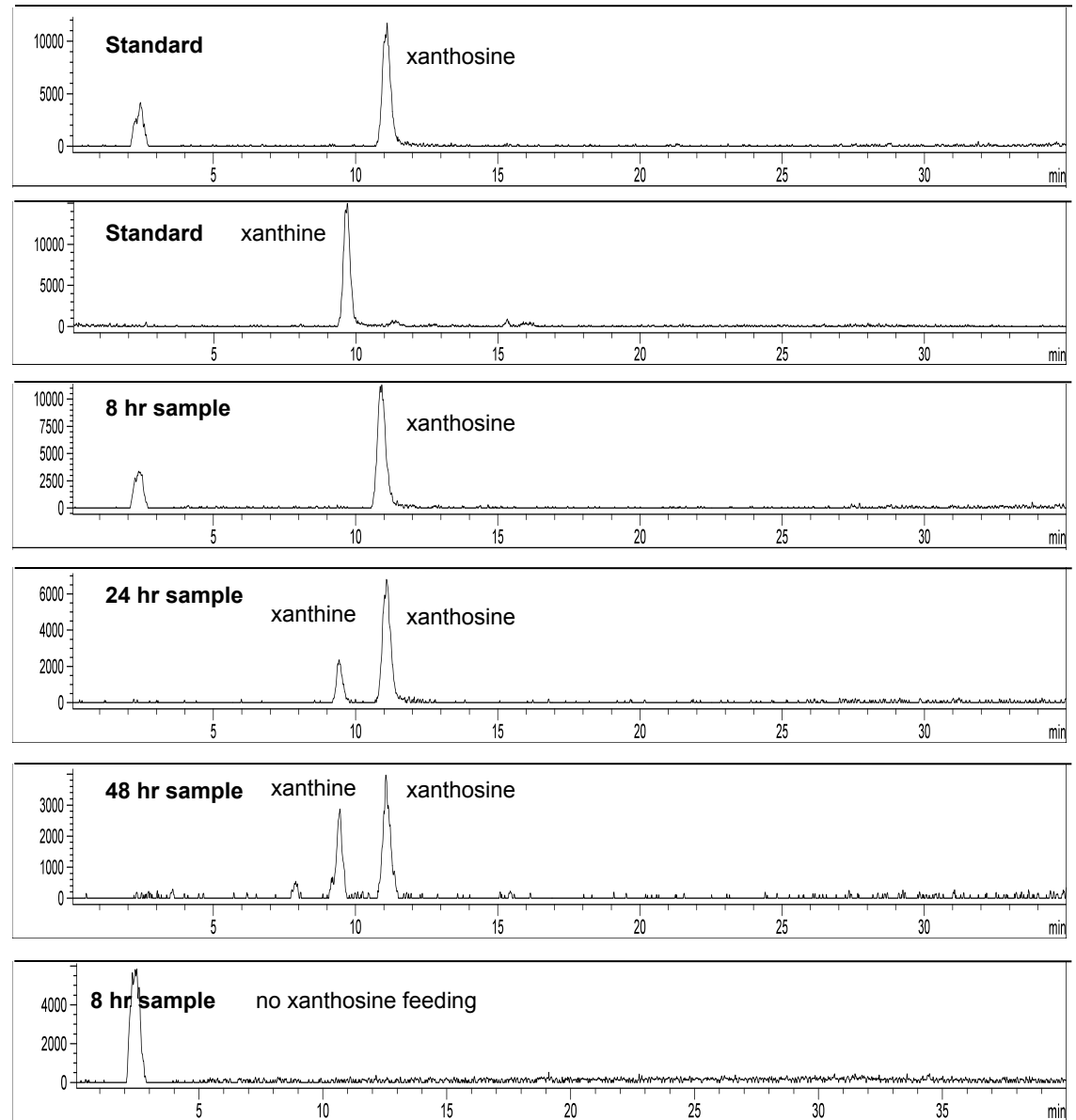
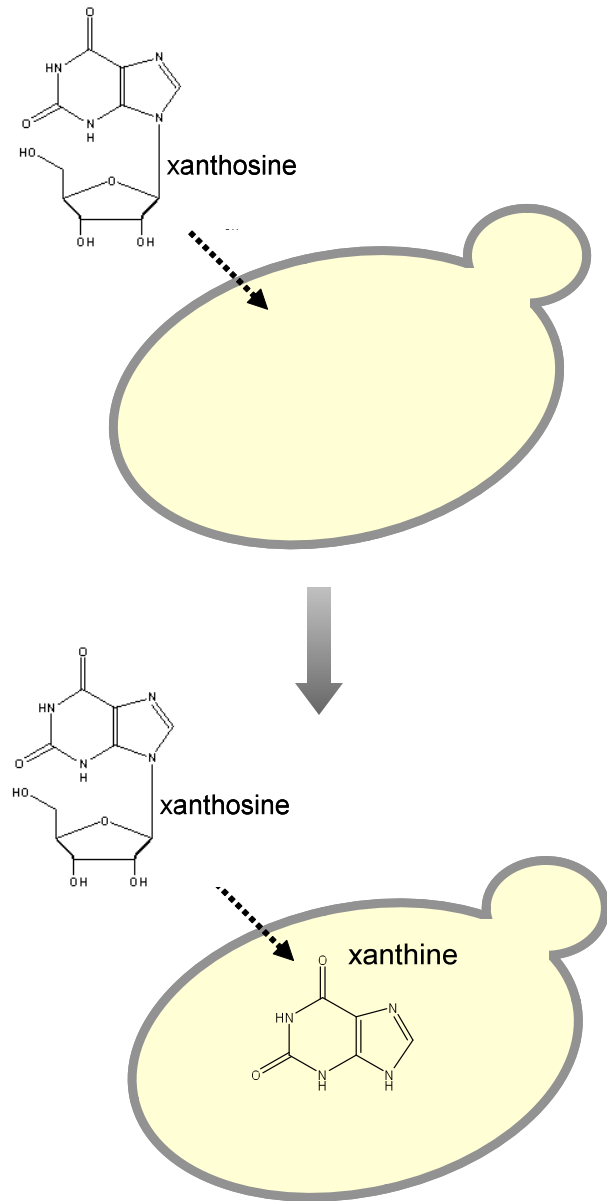
Molecular tools for optimizing metabolite production



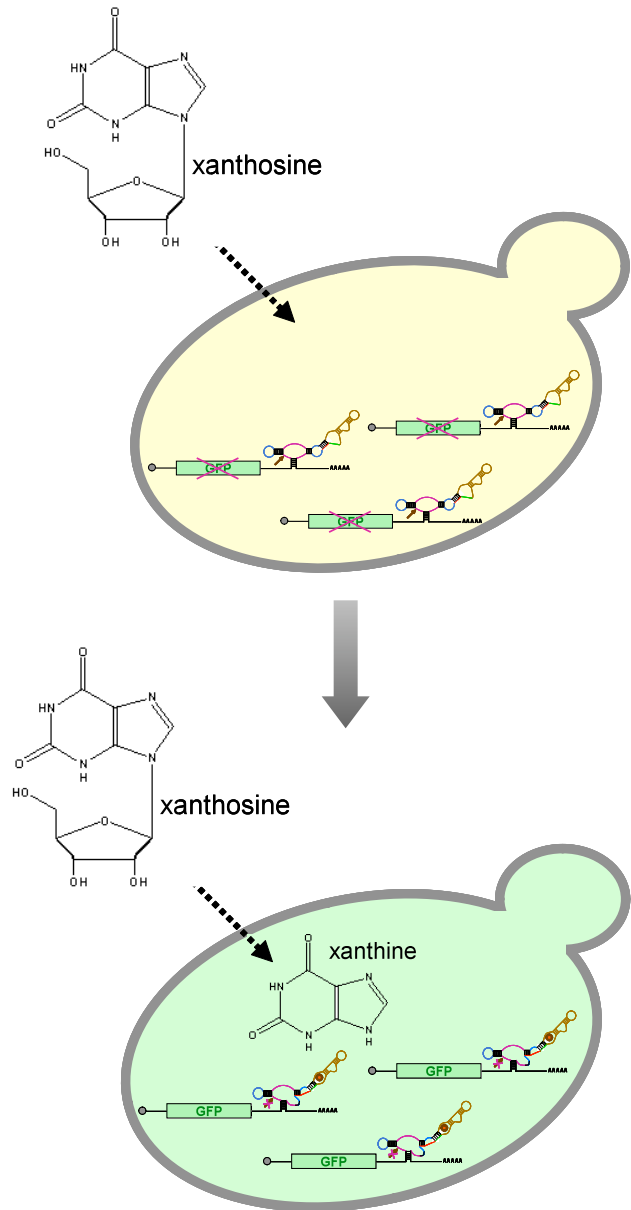
Often need to screen through large libraries of pathway/enzyme variants – invasive / analytical procedures are too time and resource intensive

Engineer noninvasive sensors of metabolite concentrations in living cells

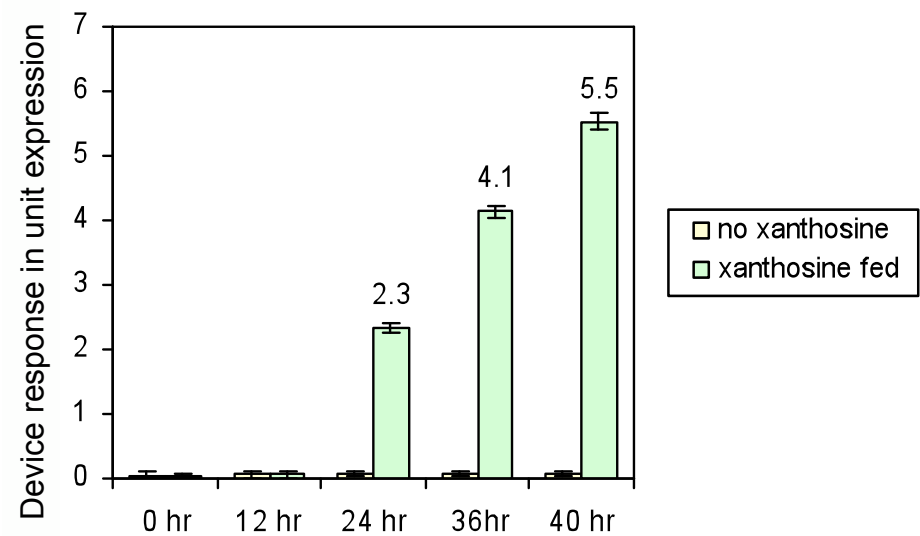
Integrating RNA devices as noninvasive sensors of metabolite concentration



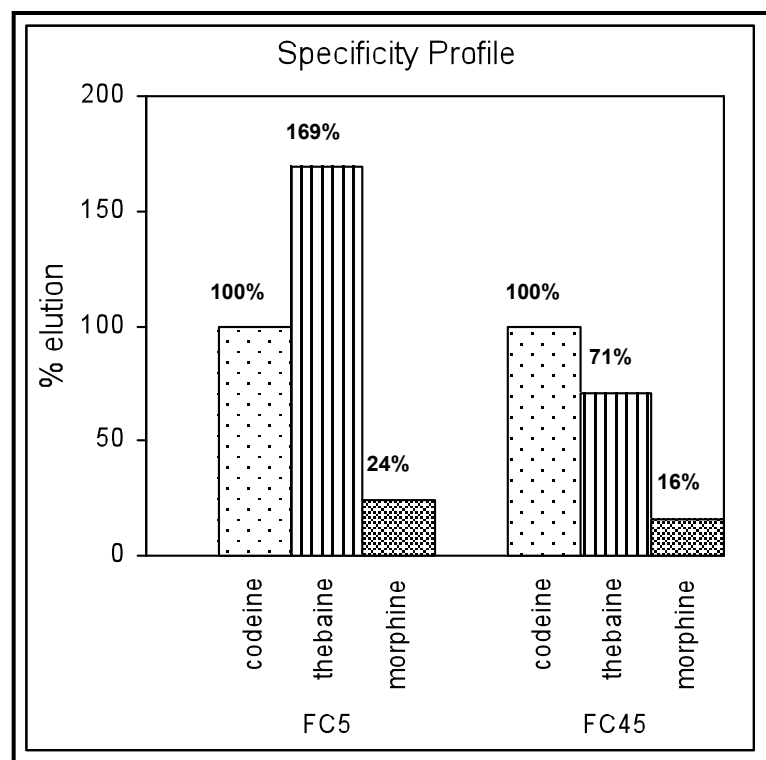
RNA devices enable noninvasive sensing of metabolite concentration



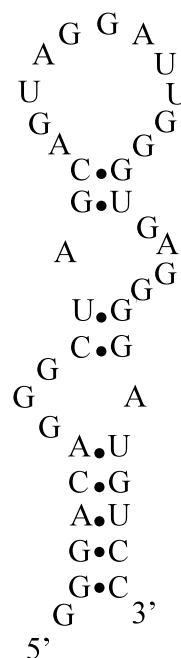
Implementation of Buffer device as a real-time noninvasive xanthine sensor



RNA sensors generated to key BIAs

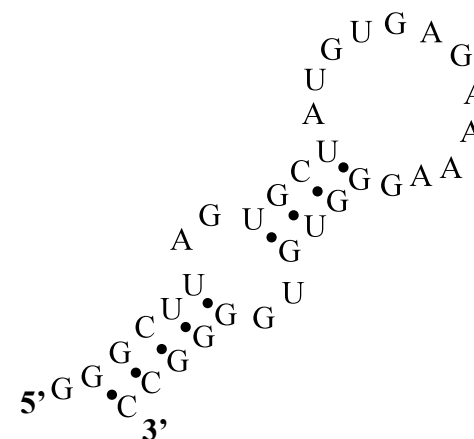


FC5

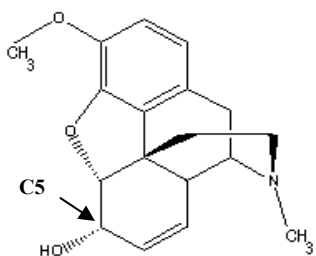


$$K_d (\text{FC5L}) = 4.55 \pm 0.14 \mu\text{M}$$

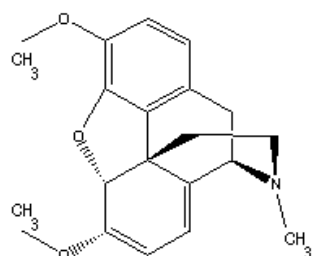
FC45



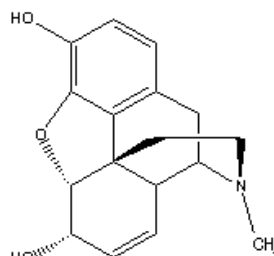
$$K_d (\text{FC45L}) = 2.59 \pm 0.09 \mu\text{M}$$



codeine



thebaine



morphine

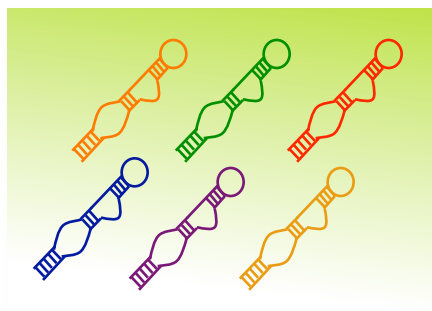
Future directions: addressing the scalability challenge

refined sensor libraries

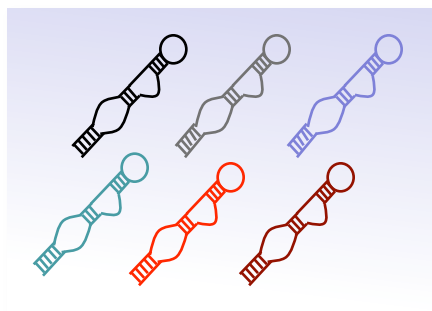
device platforms

applications

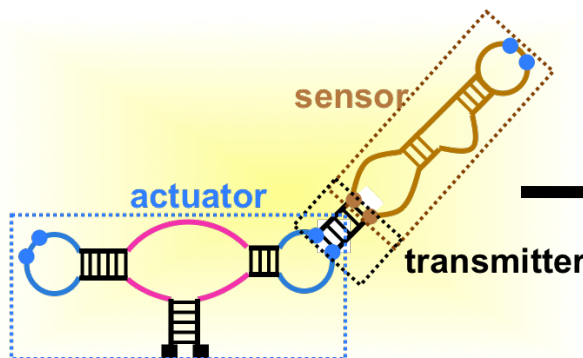
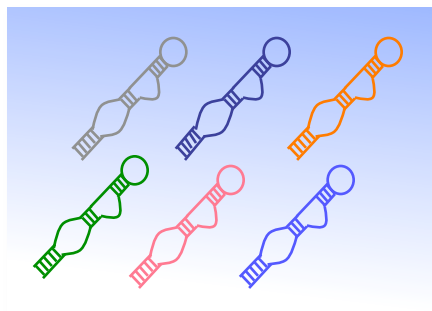
metabolites



disease biomarkers



exogenous chemicals



noninvasive
diagnostics

intelligent
therapeutics

bioprocessing /
biosynthesis

agricultural
biotechnology

*Optimization and development needed on selection
and characterization strategies*

Acknowledgements

Current Lab Members

Postdoctoral Researchers

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Graduate Researchers

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Ryan Bloom

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Andy Chang

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Leo d'Espaux

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Jay Vowles

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