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Capacity and Delay of Bacteria-Based Communication in Nanonetworks

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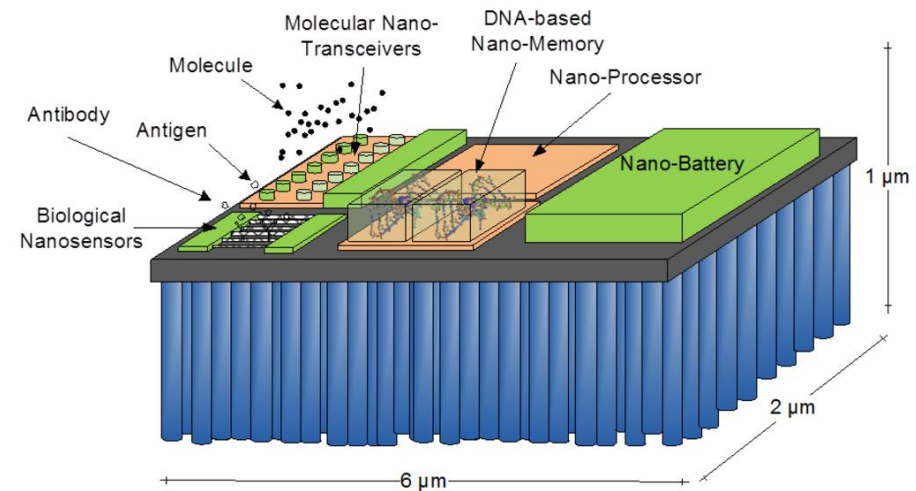
- Introduction
- Motivation and Contribution
- Effect of mutations
- Propagation model of bacteria
- Communication schemes
- Concluding Remarks

- **Introduction**
- Motivation and Contribution
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- Concluding Remarks

I. F. Akyildiz, F. Brunetti, and C. Blázquez, "*Nanonetworking: A New Communication Paradigm*," Computer Networks (Elsevier) Journal, Vol. 52, pp. 2260-2279, August, 2008.

- **Nanotechnology** is enabling the development of devices which can operate at the nanoscale.

- A **nanodevice** is the most basic functional unit, which is divided into independent nanoscale components and it is able to perform specific tasks at nanolevel, such as sensing, computing, data storing, or actuation.



○ WHY?

Due to the reduced size of the nanodevices, their limitations are clear:

- Actuation range
- Capabilities

Nanonetworks expand the possible applications of single nanodevices by a collaborative effort among them.

○ HOW?

- Bacteria-based Communication

Gregori, M. & Akyildiz, I. "A new nanonetwork architecture using flagellated bacteria and catalytic nanomotors" Selected Areas in Communications, IEEE Journal on, 2010, 28, 612 -619

● Basic concept:

- Use bacteria as carriers of encoded DNA packets

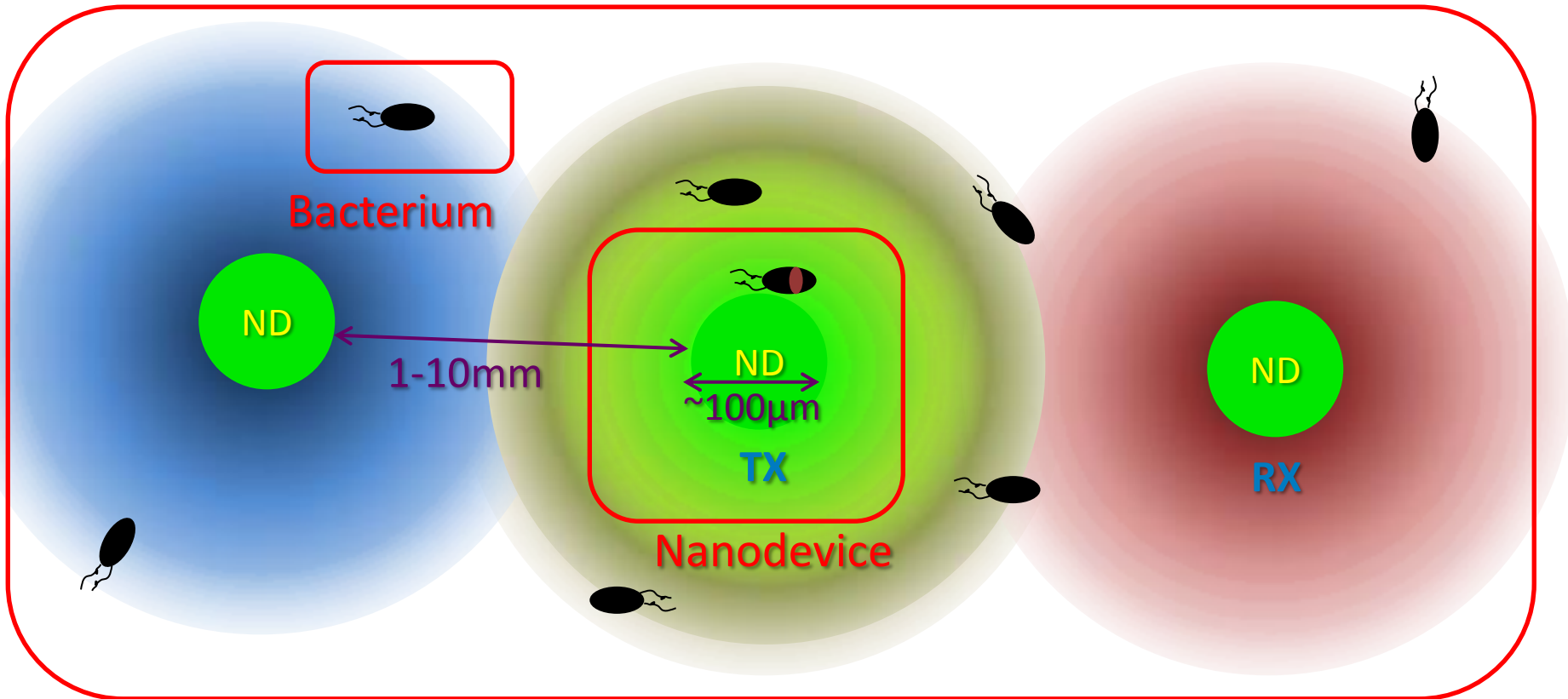
● How do bacteria move?

- Chemotaxis: bacteria random walk which follows the concentration gradient of specific substances (attractants) in the environment.

● How do bacteria reach the receiver?

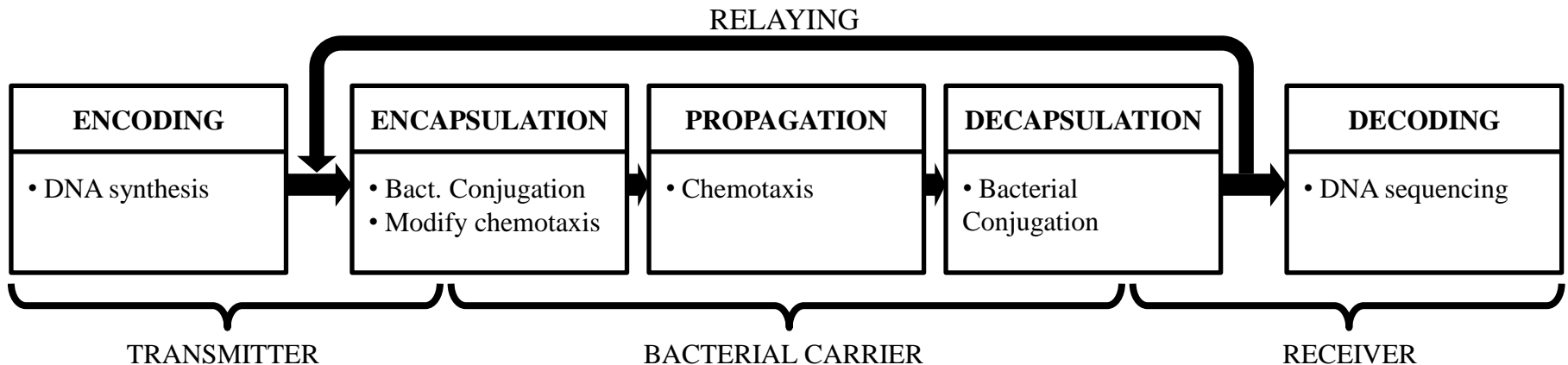
- Nanodevices stimulate the chemotaxis of bacteria towards themselves by releasing attractants.

Environment



Cobo, L. C. & Akyildiz, I. F. “Bacteria-based communication in nanonetworks”
Nano Communication Networks, 2010, 1, 244 - 256

Communication scheme



Relaying process

- The range of the attractant is limited: **Multi-hop** is needed

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- To complete an analytical model for the previous work.
 - The propagation of bacteria was only simulated.
 - The capacity model was approximated.
 - The delay was based on the simulated results.

- An analysis of the effect of the mutations during the communication process.
- A theoretical study of the propagation of bacteria.
- Definition of two different communication schemes based on bacteria.
- The study of the capacity and the delay of these systems.

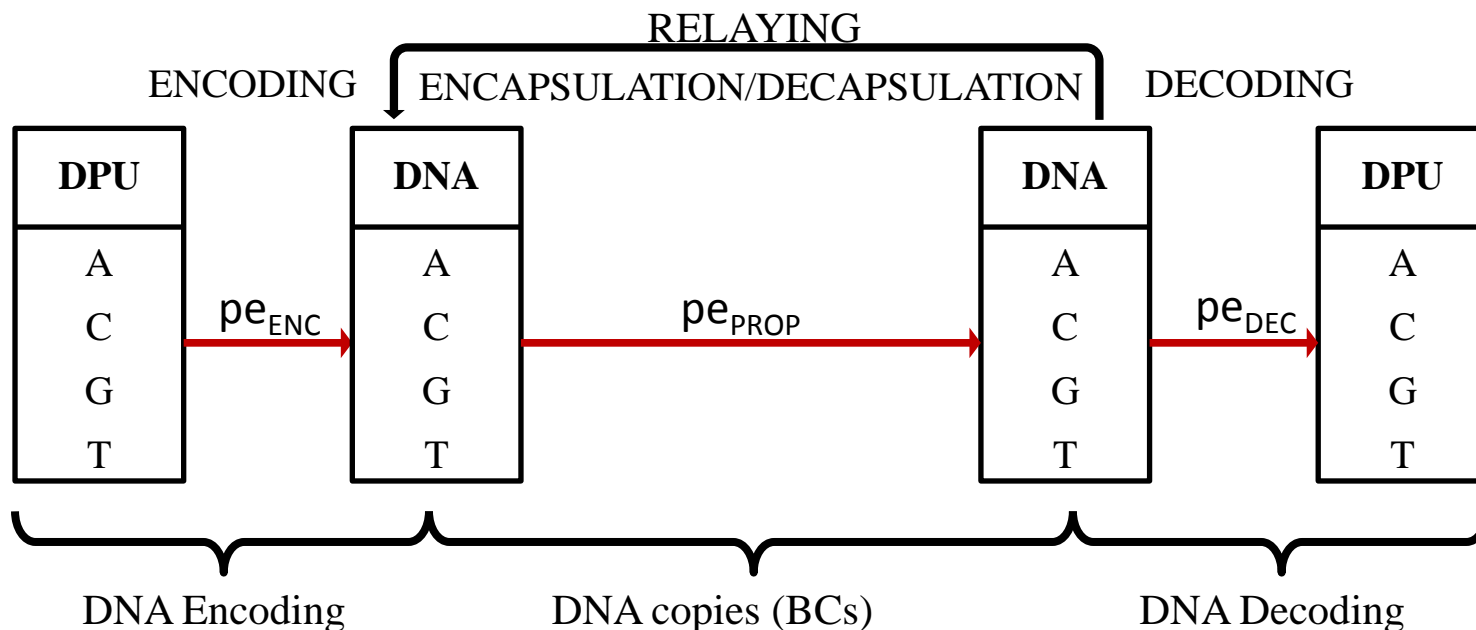
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- The information is encoded in a DNA molecule called **plasmid**.
- The basic information unit is the **base pair (bp)**.
- A plasmid can contain up to 1.6 Mbp (millions of base pairs)



Drake, J. W.; Charlesworth, B.; Charlesworth, D. & Crow, J. F. "Rates of Spontaneous Mutation"
Genetics, 1998, 148, 1667-1686

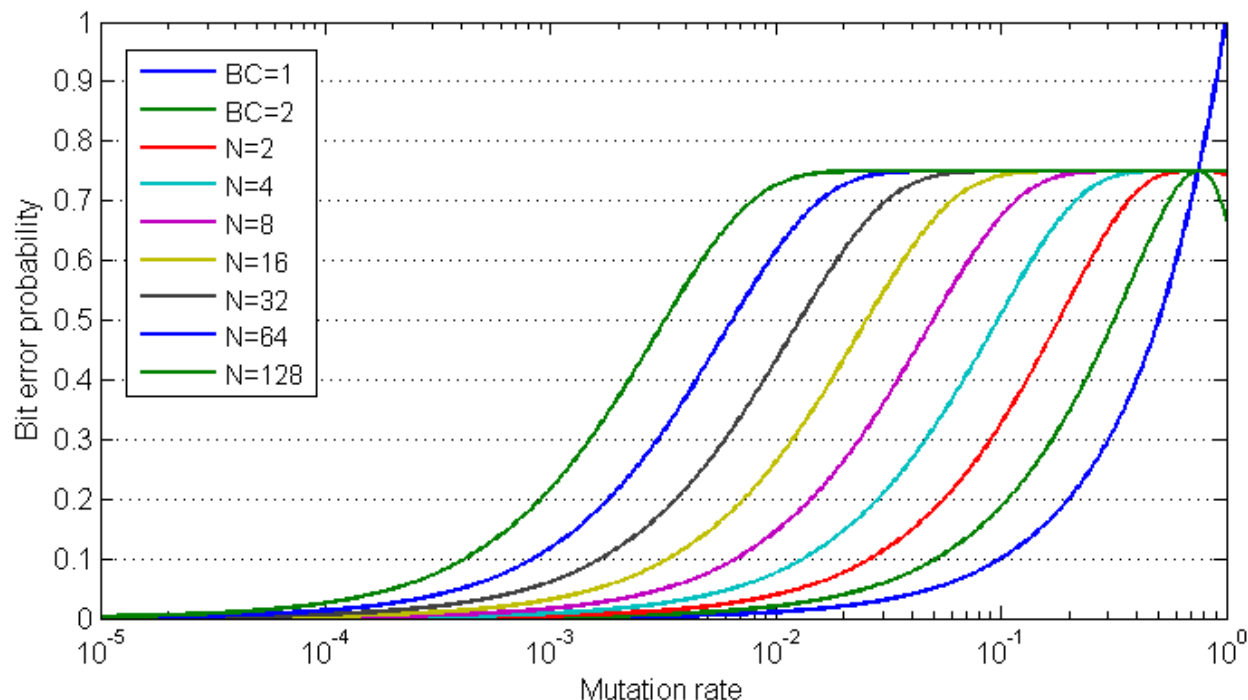
- During the communication process, **multiple copies** of the plasmid are done.
- In each copy, mutations may alter the information.



● Probability of error during the communication:

$$p_e(N) = \frac{3}{4} \left(1 - \left(1 - \frac{4}{3} p_{mut} \right)^{2N+2} \right)$$

N: Number of hops
 p_{mut} : Mutation rate

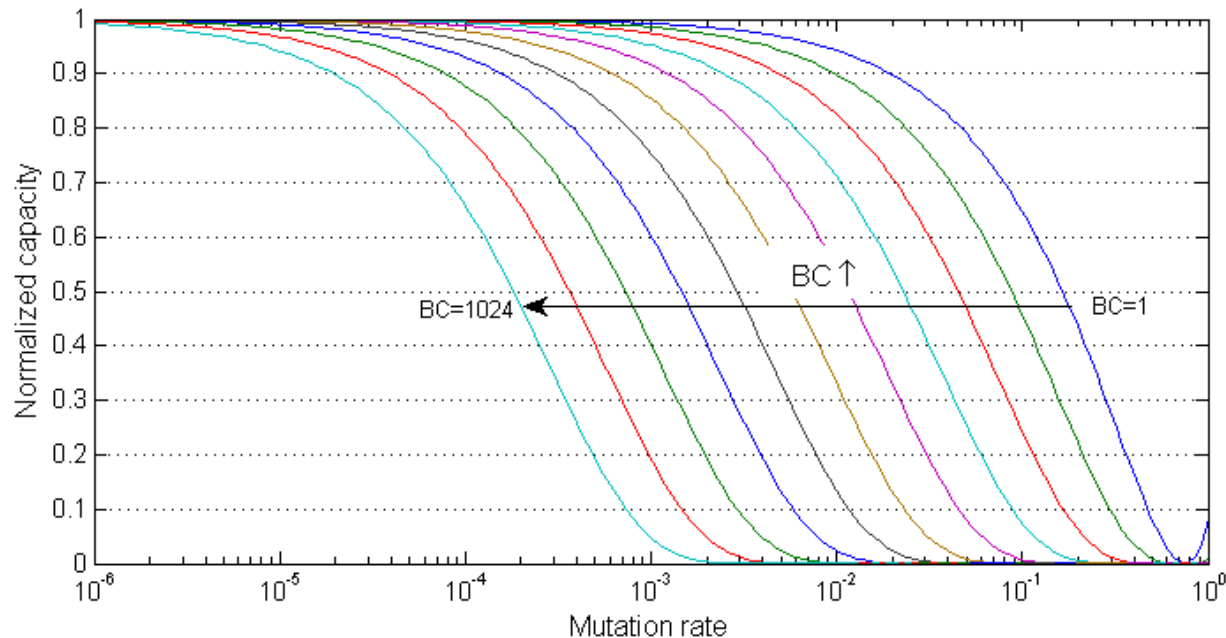


Shannon, C. E. "A Mathematical Theory of Communication" *The Bell System Technical Journal*, 1948, 27, 379-423

Capacity of a base pair (quaternary symmetric channel):

$$C = 2 - H\left(1 - p_e, \frac{p_e}{3}, \frac{p_e}{3}, \frac{p_e}{3}\right)$$

H: Quaternary entropy



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Berg, H. C. “**Random Walks in Biology**” *Princeton University Press*, 1993

● Chemotaxis

- Bacteria **random walk** which follows the concentration gradient of specific substances (attractants) in the environment.

● How does it work?

- Bacteria move in series of “runs” and “tumbles”.
- Without attractant, bacteria move **randomly** in the medium, like a particle.
- With attractant, the movement of bacteria is **biased** according to the attractant concentration.

Schnitzer, M. J. "Theory of continuum random walks and application to chemotaxis",
Phys. Rev. E, American Physical Society, 1993, 48, 2553-2568

- How to study?
 - Diffusion theory.

- Flux of bacteria (J):

$$J = -D\nabla c(\bar{x}, t) + v_{drift}(\bar{x})c(\bar{x}, t)$$

- Where:

$$D = \frac{v^2}{n \left[\alpha_0 (1 - \Theta) + (n - 1) D_{rot} \right]}$$

$$v_{drift} = \frac{v^2 g \nabla C (1 - \Theta)}{n \left[\alpha_0 (1 - \Theta) + (n - 1) D_{rot} \right]}$$

v: velocity of bacteria
n: number of dimensions
 α_0 : tumble rate
 Θ : angular correlation
 D_{rot} : Rotational diffusion
g: Attractant sensing gain
C: Attractant gradient

Continuity Differential Equation

$$\frac{\partial c(\bar{x}, t)}{\partial t} + \nabla_{\bar{x}} \cdot (v_{drift}(\bar{x}) c(\bar{x}, t)) = -J$$

Initial condition: $c(\bar{x}, t) = 0, t > 0$

$$c(\bar{x}, 0) = \delta(\bar{x} - \bar{x}_{TX})$$

$$\frac{\partial c(\bar{x}, t)}{\partial t} = -J$$

Drift velocity: $v_{drift} \propto \nabla C$

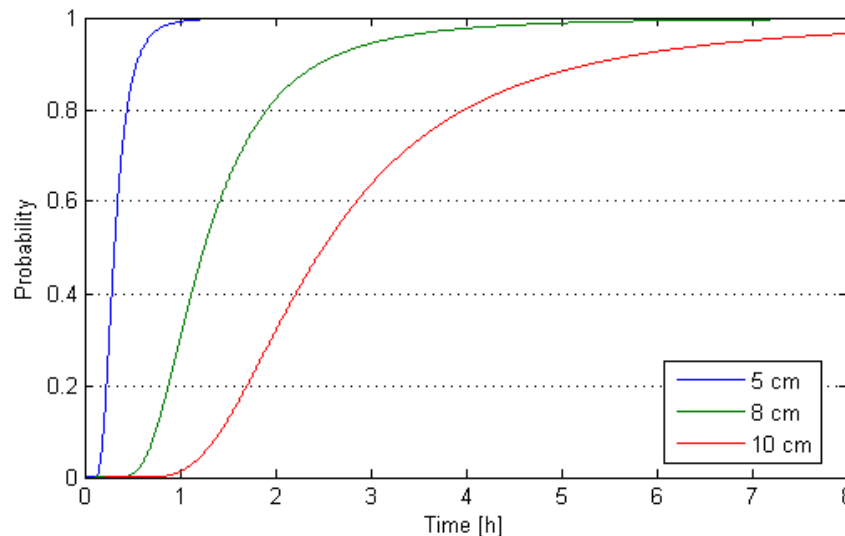
Attractant concentration: $C(r) = \frac{Q}{4D\pi r}$

Q: Releasing rate
D: Attractant diffusion coefficient

- With $c(\mathbf{x}, t)$, the probability distribution of bacteria in the medium, the distribution of the arrivals can be calculated:

$$Err_{arr}(t) = \int_{\langle Rx \rangle} c(\bar{x}, t) dVol$$

$\langle Rx \rangle$: Volume of the receiver



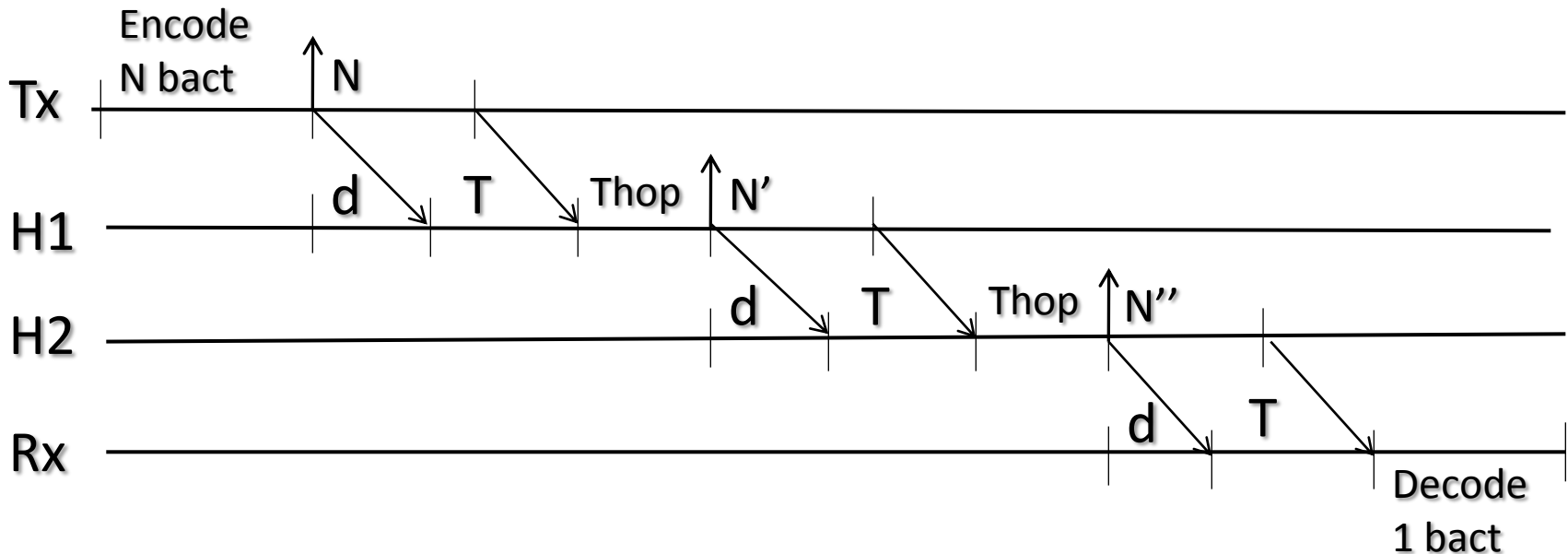
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- Communication schemes
 - ▣ Synchronous scheme
 - ▣ Asynchronous scheme
 - ▣ Comparison

● Characteristics

- The time is divided into slots (duration T).
- In each time slot we release N bacteria encoding the same information.
- Only one bacterium has to reach the receiver during the T to receive that data.
- If no bacteria reach the receptor in T , the data is lost.
- We wait a delay d , to maximize the probability of receiving one bacterium.

System delay:



$$delay = t_{tx}(N) + S \cdot (d + T) + (S - 1)t_{hop} + t_{rx}$$

Capacity:

$$C(B, d, T, N, S) = B \frac{P_{Rx1bac}(d, T, N, S)}{T} C_{bp}$$

B: Base pairs
encoded
C_{bp}: Capacity base
pair

Probability of receiving at least one bacterium in one hop:

$$P_{Rx1bac}(d, T, N, S) = \left(1 - (1 - P_{1Rx}(d, T))^N\right)^S$$

$$P_{1Rx}(d, T) \approx arr(T + d) - arr(d)$$

Maximization of the capacity:

$$\max_{d, T, B, N} C(B, d, T, N, S)$$

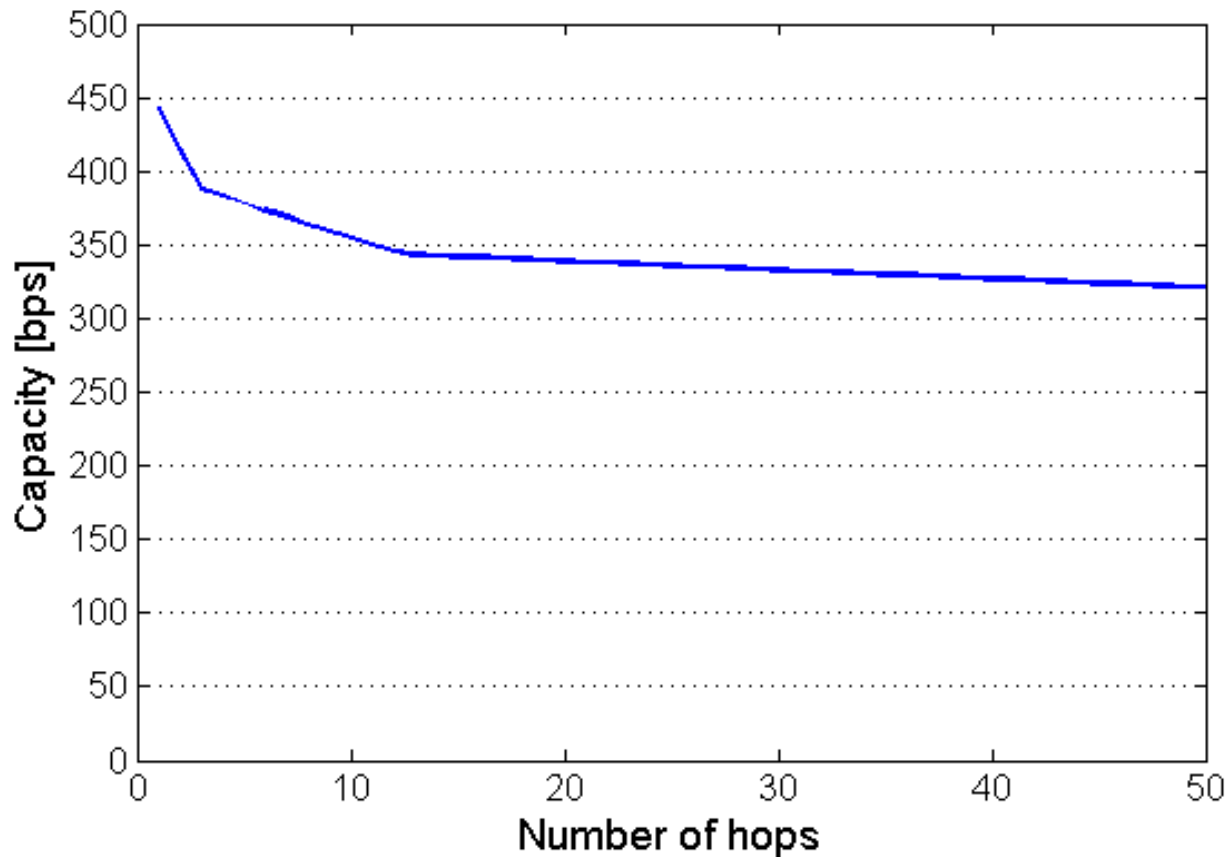
conditions:

$$T \geq T_{enc}(B, N)$$

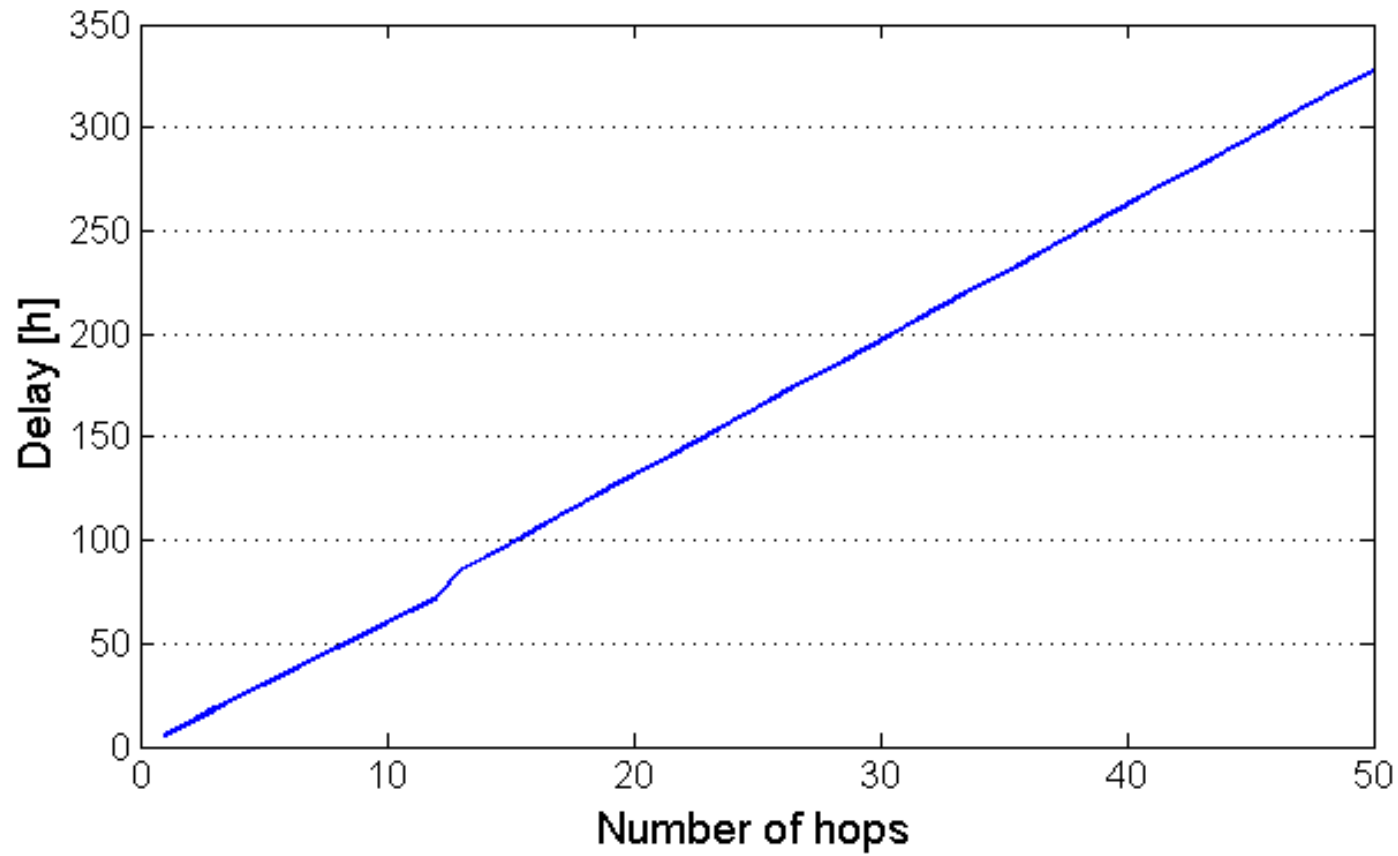
$$B \in [0, BactCap]$$

$$N \in \mathbb{N}$$

Capacity:



System delay:

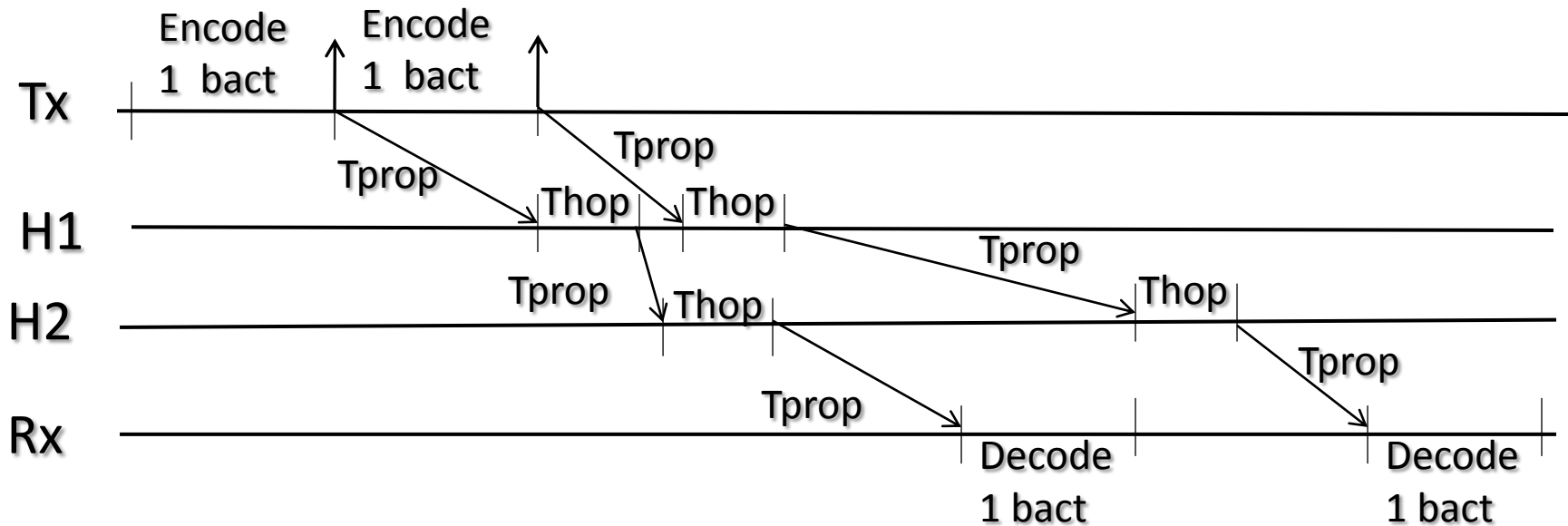


- Communication schemes
 - ▣ Synchronous scheme
 - ▣ **Asynchronous scheme**
 - ▣ Comparison

○ Characteristics

- The transmitter is continuously releasing bacteria at the encoding speed.
- The information goes from the transmitter to the receiver using other nanodevices.
- If the bacterium do not reach any of the nanodevices, the data is lost.
- The bacterium can use a Tout in each hop or not.

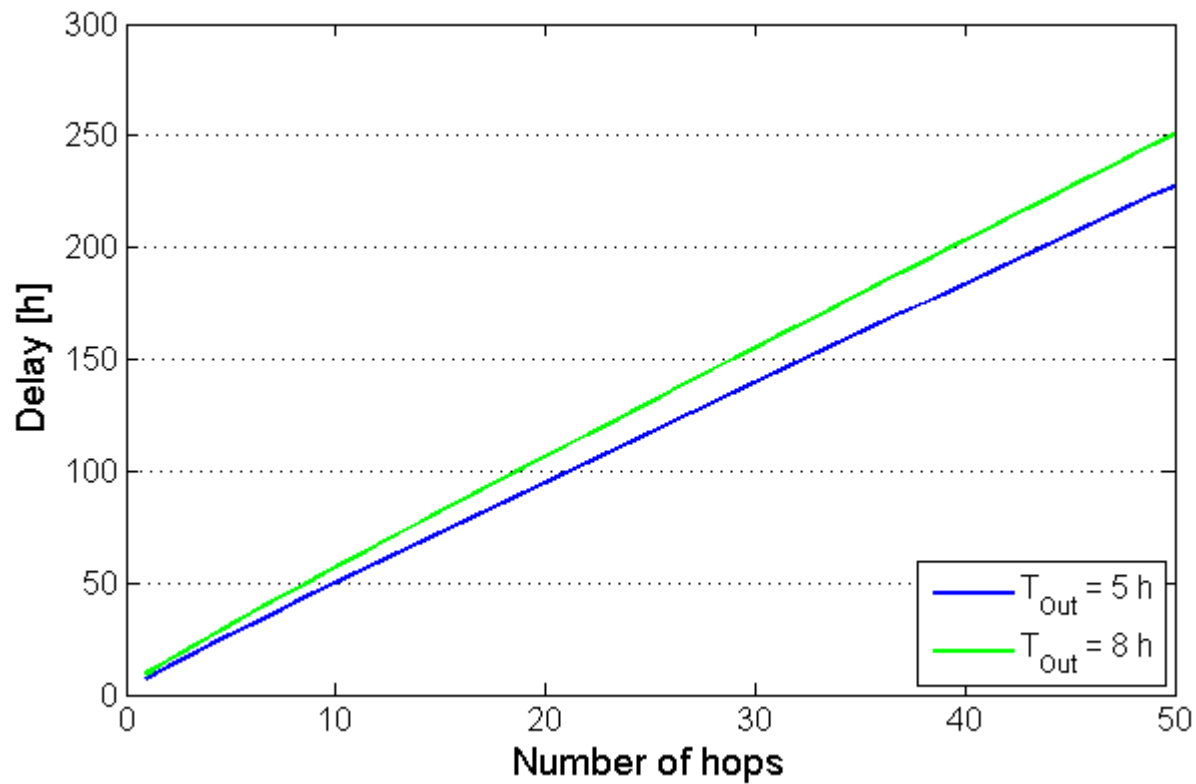
System delay:



$$Delay = t_{tx} + t_{prop}(S) + (S - 1) \cdot t_{hop} + t_{rx}$$

$$f_{t_{prop}}(S) = \overbrace{f_{arr} * \dots * f_{arr}}^S \quad [pdf]$$

○ Average system delay:



Capacity:

$$C = C_{bp} \cdot r_{encode}(B) \cdot p_{Rx}$$

r_{encode} : effective
encoding rate

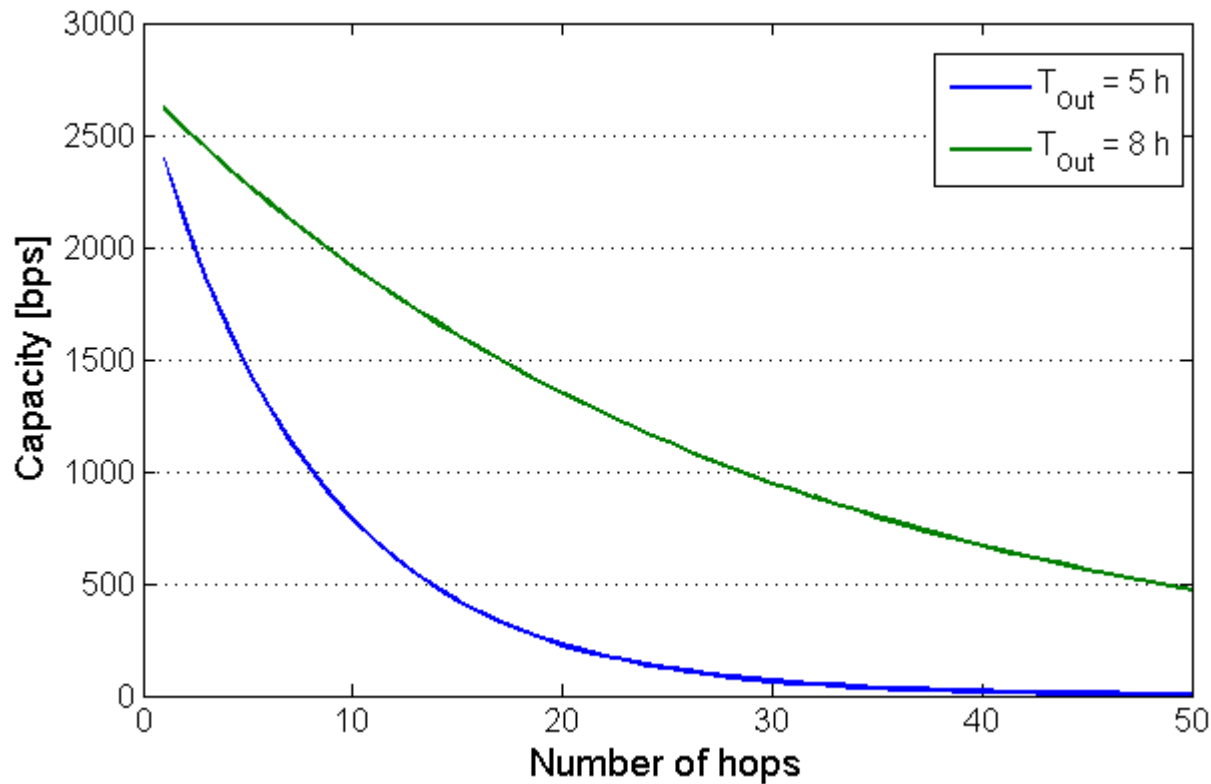
- Probability of receiving at least one bacterium in one hop:

$$p_{Rx} \approx (1 - P_{T_{out}})^S$$

$$P_{T_{out}}(T_{Out}) = 1 - arr(T_{Out})$$

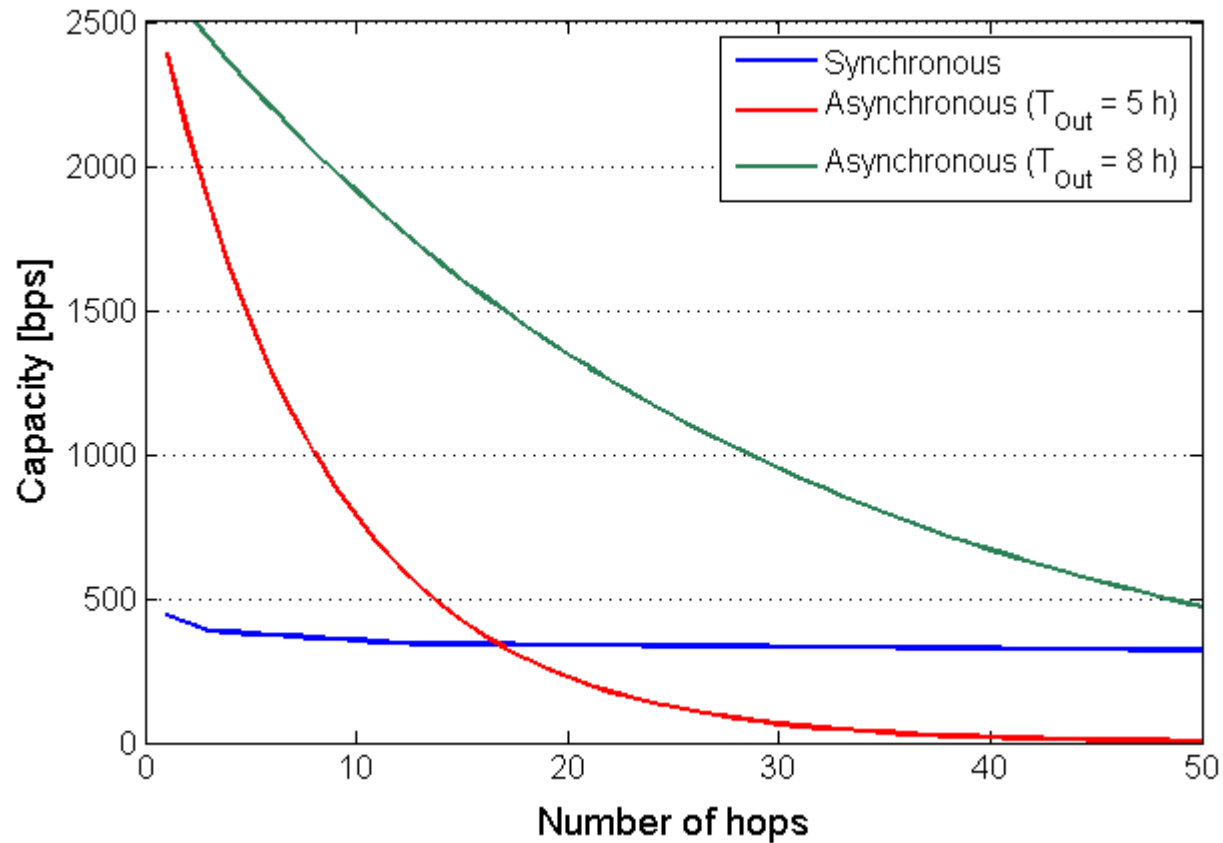
- The optimal capacity can be achieved when the plasmid encodes the maximum of information.

Capacity:

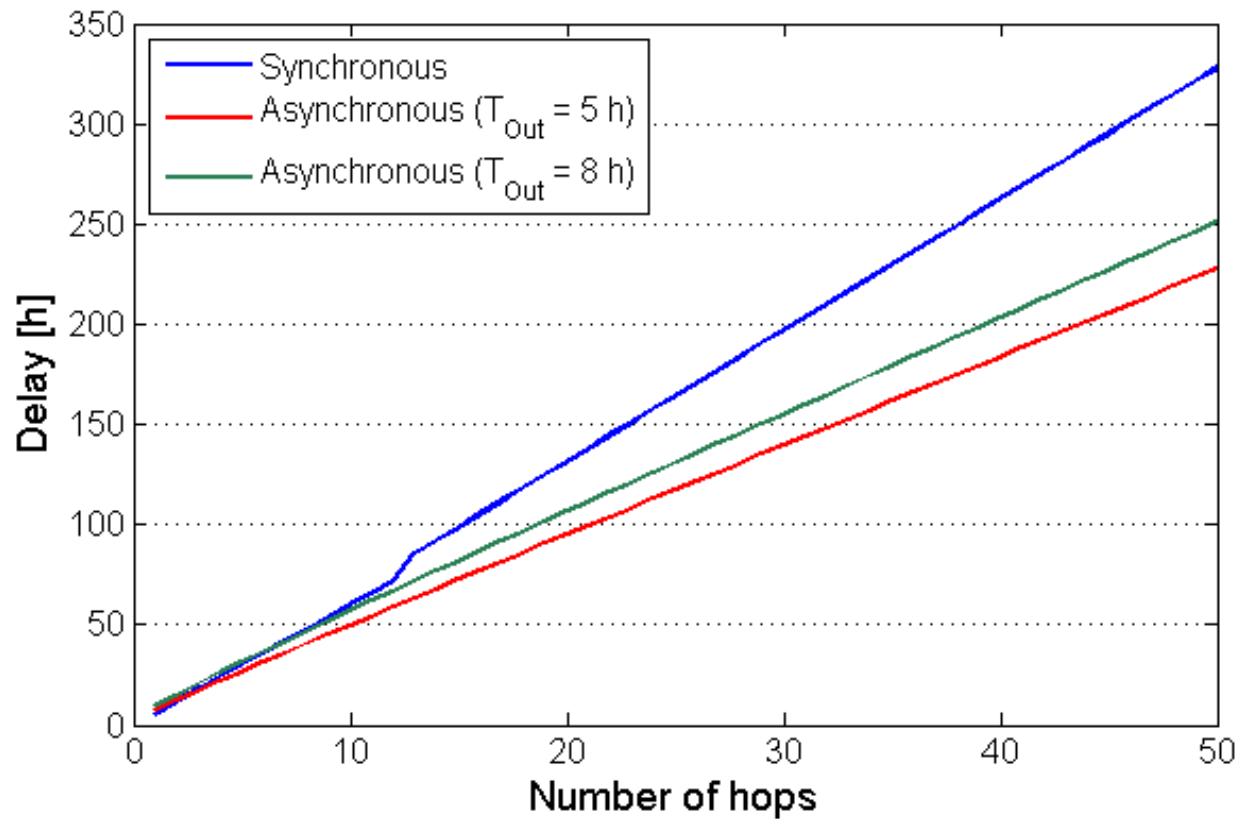


- Communication schemes
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Capacity



Delay



- The synchronous system is appropriate when:
 - *The order of the information is important*
 - *The delay has to be known and constant*
 - *The number of hops is high*

- The asynchronous system is appropriate when:
 - *Neither the order nor the exact moment when a packet is received is important*
 - *The number of hops is small*

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- A theoretical study of the effect of the mutations has been performed and validated.
- An analytical solution for the bacteria propagation has been found and validated.
- Two different communication schemes have been proposed and the capacity and the delay have been calculated.
 - Synchronous scheme
 - Constant delay and high performance in long range transmissions
 - Asynchronous scheme
 - High performance in short range transmissions

Thank you!



Thank you very much for your attention.
Any question?