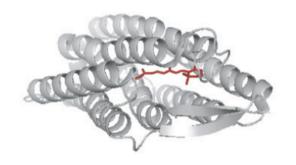
Light-sensing mecanisms in prokaryotes

An overview and possible applications

Major classes of protein domains

Protein domain

Chromophore



Bacteriorhodopsin:

- •First found in haloarchaea
- •Quite an easy mechanism to create a proton gradient through the use of light (through the isomerisation of retinal)
- Absorbing mostly green light
- •Some Bacteriorhodopsin-like genes have been found in other organisms. There are named **proteorhodopsins**. They have been expressed in *E.Coli*

Bacteriophytochrome

Bilin derivative



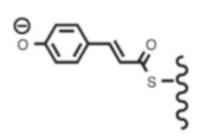
Bacteriophytochrome:

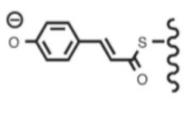
- •Found in many different types of bacteria
- •Based on the isomerisation of the chromophore
- •Most ouput domain are HK and EAL-GGDEF (involved in cyclic di-GMP hydrolysis)
- Absorbing red and far-red light

PYP

p-coumaric acid







PYP domain:

- •Based on the isomerisation of coumaric acid
- •Mostrly absorb blue light (maximum at 446 nm)
- •Usually single-domained but can also form hybrids with HisKA, GGDEF or EAL output domains

LOV-family:

- •First discovered in plant phototropin and more recently in bacteria
- •Based on the transitional formation of a covalent bond between the cysteine and the chromophore (flavin)
- •It is a subclass of the PAS domain (large classe of sensing and protein interaction based on the use of a co-factor)
- •Named after the input signal of this domain (Light, Oxygen and Voltage)
- •B. Subtilis has a LOV domain (in photoreceptor YtvA)

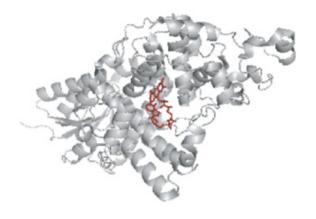
BLUF Flavin



BLUF-domain:

- •Similar struture with the PAS domain
- •Found in *Rhodobacter sphaeroides*
- •The name comes frome Blue Light sensing Using Flavin
- •Logically, this domain sense blue light
- •E. Coli has a BLUF domain (more precisely the YgcF photoreceptor protein)

Cryptochrome

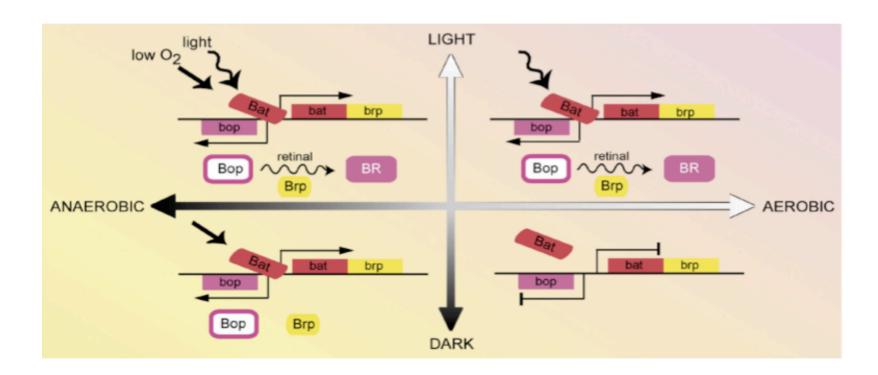


FAD

Cryptochrome:

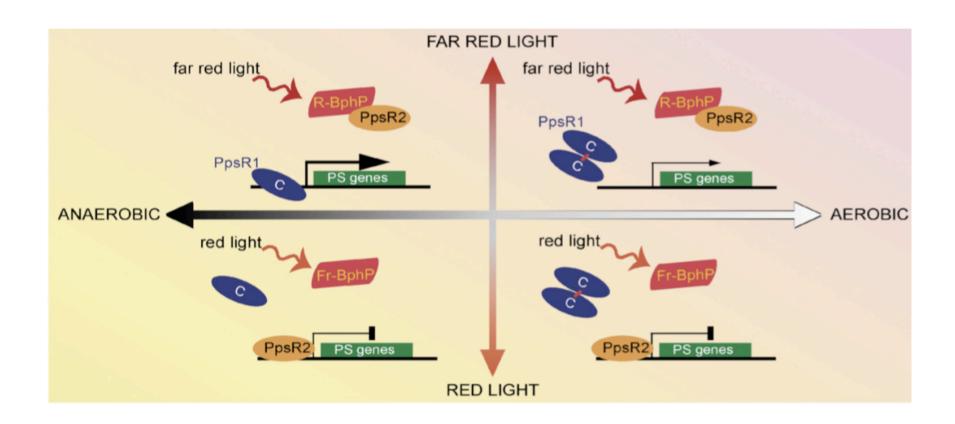
- •Found in many different organisms, like plants and animals as well as in bacteria
- •The chromophore is a flavin combined with pterin
- •Absorbs in the violet-blue domain of the spectrum
- •Light activates a phosophorylation which then might have gene regulation output

BR regulation in *H. Salinarum*



We notice that in fully aerobic and/or dark conditions, the bacteria doesn't synthetise the protein

Bacteriophytochrome and PS regulation in *C. Bradyrhizobium*



Once again we see that the expression of the photosystems genes are regulated by light and oxygen conditions

Examples of bacteria with domains

	BLUF	LOV	PYP	Phytochrome	Sensory rhodopsin		BLUF	LOV	PYP	Phytochrome	Sensory rhodopsin
Acidovorax avenae	2	1		1		Nitrosococcus oceani		1			
Acidovorax sp. JS42	1					Nitrosospira multiformis		1			
Acinetobacter baumannii	1					Novosphingobium aromaticivorans		2			
Acinetobacter sp. ADP1	4					Oceanicola batsensis				1	
Agrobacterium tumefaciens				3		Oceanicola granulosus		1		1	
Alkalilimnicola erhlichii		1		1		Oceanobacillus iheyensis		1			
Alpha proteobacterium HTCC2255	1					Oceanobacter sp. RED65	1	1			
Azoarcus sp. EbN1						Polaromonas sp. JS666	2				
Bacillus subtilis subsp. Subtilis		1				Pseudoalteromonas atlantica	2			1	
Bdellovibrio bacteriovorus	1			1		Pseudoalteromonas tunicata	2				
Blastopirellula marina DSM 3645				1		Pseudomonas aeruginosa				1	
Brucella melitensis		1				Pseudomonas entomophila				1	
Erythrobacter sp. NAP1		1		1		Rhodoferax ferrireducens	1				
Escherichia coli	1					Rhodopirellula baltica					
Exiguobacterium sibiricum					1	<i>Roseovarius</i> sp. HTCC2601				1	
Flavobacterium johnsoniae				2		Sagittula stellata	2			1	
Flavobacterium sp. MED217				1		Shewanella baltica		2			

Possible applications

Photo-sequenced protein synthesis

- •Use a pre-existing light-triggered mechanism (in *E. Coli* or *B. Subtilis*) to act on the synthesis of a protein (e.g. the YcgF mechanism).
- •A second light receptor could be added from another bacterium (if possible at the other end of the light spectrum e.g. bacteriophytochrome) that would be linked to another pathway (leading to the formation of another product/enzyme)

•

- •Industrial Use: having a feedback on the bacteria's production. A sensor in the bioreactor could calculate the concentration of a product (or toxic byproduct) and report back to the bacteria (inhibitate or activate the production) through light control
- •« Easy » and fast way to « communicate » with the cell
- •There would also be the possibility to have a sequenced production. A informatic system would switch from one light to another inducing one or expression or another

Advantages:

- Non invasive process, no risk to harm the cells (in opposite to chemical inductors)
- Faster process than chemicals (no need to wait for diffusion in the reactor for example)
- Light is easly produced

Disadvantages:

- The two spectrum domains of the different photoreceptor must be as far away as possible (to reduce the risk of interferences)
- Some mechanisms are anaerobic-dependant. The concentration of oxygen must be kept low

Combination w/ magnetotactic bacteria

- For diagnostics, or sample screening, we don't always need the whole set of receptors --> Subset of receptors expressed depending on light.
- Regulation by light could also permits a low-budget project.
 Eg: using LED or translucent paper (permit utilization of standard bulb).
- Oxygen level is very critical for vesicles' formation in magnetotactic bacteria --> We need a gene regulation system which is related to the light differences rather than the aerobic condition.

C.Bradyrhizobium

C. Bradyrhizobium sp. bacteriophytochrome regulation of PS gene expression

FAR RED LIGHT

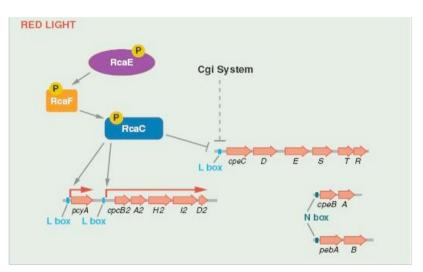
far red light

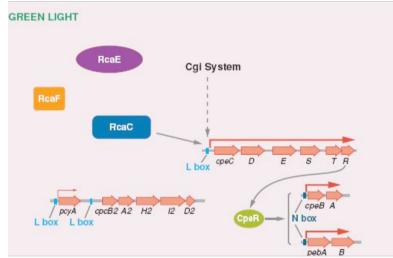
PpsR2

P

• The result is more correlated to the light differences than the oxygen level: Far-red --> Expression; Red --> Repression.

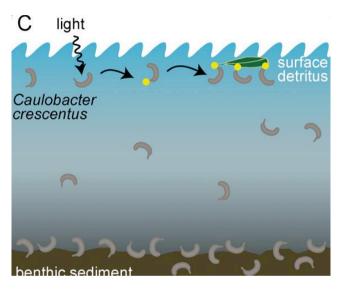
Fremyella diplosiphon

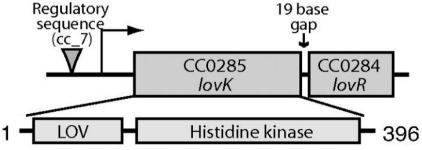




 Pigmented proteins depend on the color of ambient light --> Insert receptor genes downstream of the pigmented proteins.

Caulobacter crescentus





- Exposing cells that are coordinately overexpressing LovK and LovR to blue light enhances the attachment capacity of these cells.
- Light regulates LovK kinase activity.
- LovR regulates cell-surface attachment.

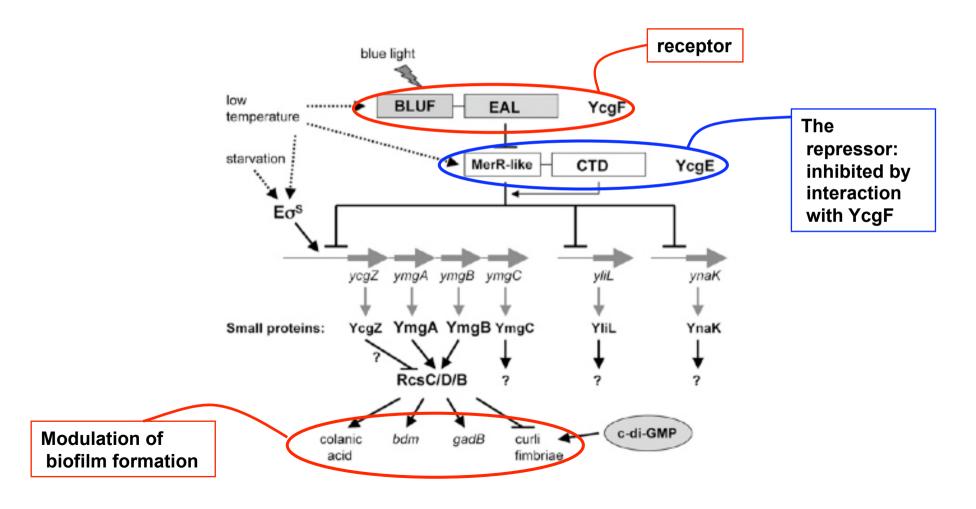
A few fast facts:

- YcgF: blue-light sensor of Escherichia coli
- BLUF: receptors that sense **B**lue **L**ight by **U**tilizing **F**AD as a chromophore
- EAL: domain involved in c-di-GMP (cyclic di-guanosine monophosphate) hydrolysis
- c-di-GMP: global second messenger used by bacteria to control multicellular behavior

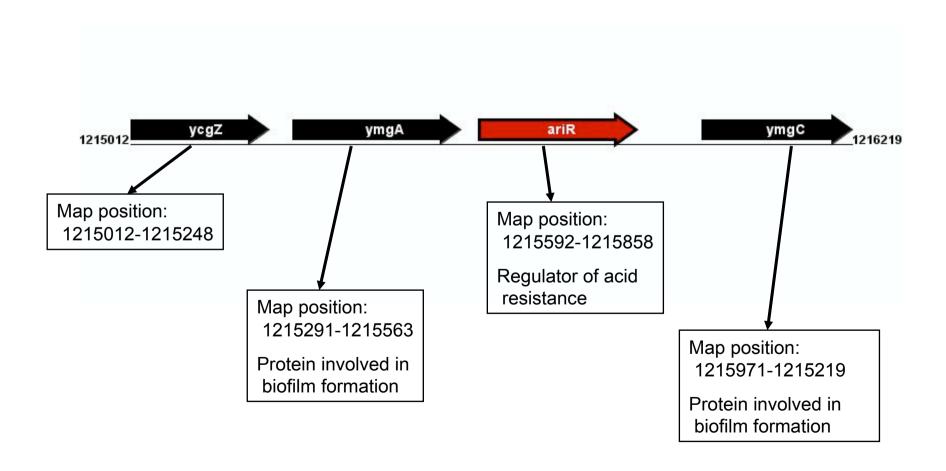


The receptor in reality doesn't interact with c-di-GMP, but instead functions as an anti-repressor in a pathway involved in biofilm formation/maturation.

The YcgF pathway:

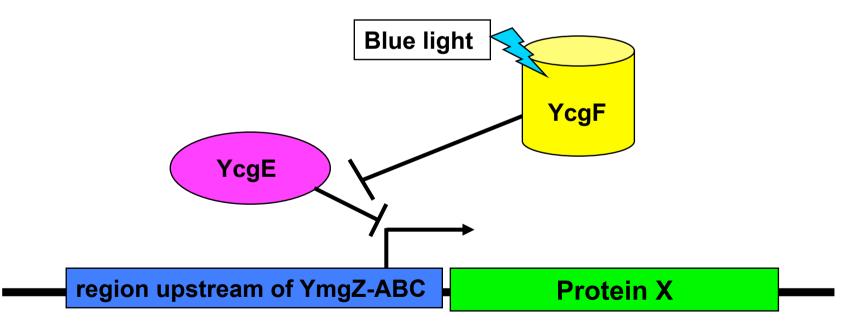


The YcgZ-ABC operon in *Escherichia coli*:



The project idea:

Link the YcgF pathway with the biofilm subject by promoting the secretion of a protein that could give interesting properties to the film to the YcgF pathway: use a promoter repressed by YcgE.



Advantages:

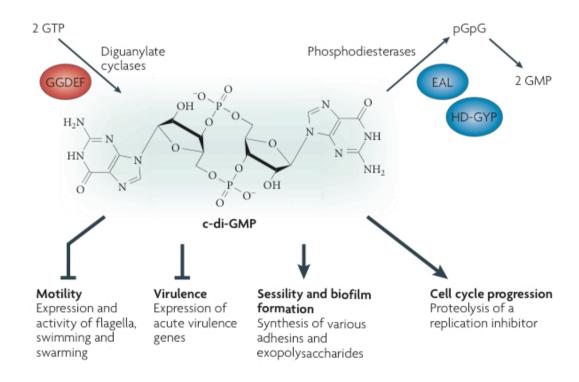
- It is a native pathway in *E. coli* (which is easy to manipulate)
- This type of mucosal biofilm is "naturally" induced by blue light: we could think of applications for an aquatic environment

Possible problems:

- Articles very recent (January & April 2009) so not much info about the sequences and proteins
- We would need to "characterize" the type of biofilm the bacteria form when the YcgF pathway is active
- The pathway seems to be part of a stress-response circuit: there could be interferences with other pathways

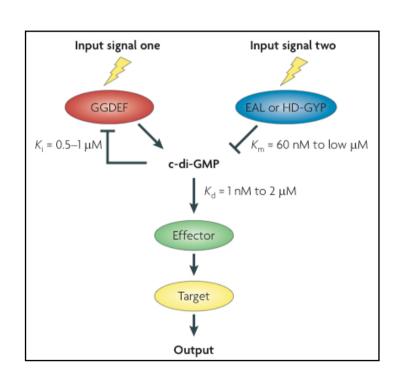
An interesting molecule for biofilms: c-di-GMP

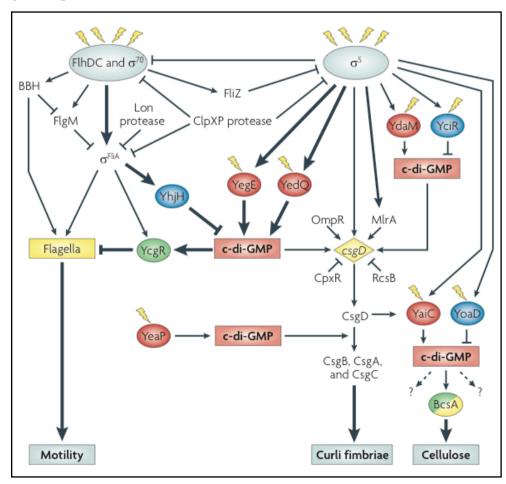
The molecule and its roles:



An interesting molecule for biofilms: c-di-GMP

Basic (left) and more complex pathways (right, in *E. coli*):





An interesting molecule for biofilms: c-di-GMP

Examples of effects on c-di-GMP in various organisms:

Table 1 c-di-GMP-binding effector components											
Effector family*	Example	Species	Functions controlled [‡]	Refs							
Protein effectors											
PilZ (+)	Alg44	Pseudomonas aeruginosa	Alginate synthesis (+)	55							
PilZ (+)	BcsA	Various Gram-negative bacteria	Cellulose synthesis (+)	54,66							
PilZ (+)	DgrA	Caulobacter crescentus	Flagellar activity (–)	67							
PilZ (+)	PilZ	P. aeruginosa	Twitching motility (–)	115							
PilZ (+)	Plz proteins	Vibrio cholerae	Virulence gene expression	68							
PilZ (+)	YcgR	Escherichia coli and Salmonella spp.	Flagella activity (–)	69							
FleQ (–)	FleQ	P. aeruginosa	Flagella expression (+) and Pel (part of the EPS) synthesis (–)	58							
PelD (+)	PelE	P. aeruginosa	Pel (part of the EPS) synthesis (+)	59							
I site effectors (+)	PopA	C. crescentus	Cell cycle progression (+)	13							
RNA effectors											
GEMM (+ and –)	Vc1 (encoded by gbpA)	V. cholerae	Intestinal adhesion	61							
GEMM (+ and –)	Vc2 (encoded by VC1722)	V. cholerae	Biofilm formation and rugosity	61							
GEMM (+ and –)	Cd1	Clostridium difficile	Flagella synthesis	61							

^{*}A + indicates that the effector is activated by c-di-GMP and a – indicates that the activity of the effector is reduced by c-di-GMP.†A + indicates that the function is positively controlled by the effector and a – indicates that the function is negatively controlled by the

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