Cellular Motility



<u>Course 5:</u> Collective motility in bacteria

Thomas Lecuit chaire: Dynamiques du vivant

CE

-1530-



Bacteria swim, propelled by flagella

Variations on the theme of flagella rotation in different bacteria



Howard Berg http://www.rowland.harvard.edu/labs/bacteria/movies/ecoli.php

Rhodobacter sphaeroides

- single flagellum rotates in single direction
- motor stops and flagellum relaxes and coils



E.coli

Sinorhizobium meliloti

- CW rotation of motor/flagellum in bundle
- slow rotation causes flagella to separate



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J. Armitage and R. Schmidt Microbiology. 143, 3671-3682 (1997)

The swim of the bacterium E. coli



The swim of the bacterium E. coli

Without chemoattractant

- 6 flagella bundle when they rotate counterclockwise (CCW)
- Bundles rotate and propel E. coli along runs
- Runs are followed by tumbles due to CW rotation of flagella which are no longer bundled





Howard Berg http://www.rowland.harvard.edu/labs/bacteria/movies/ecoli.php





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Howard Berg and Douglas Brown. Nature 239, 500-504 (1972)

The swim of the bacterium E. coli



- Biased random walk in a spatial gradient
- Temporal gradient sensing
- Memory



Key properties of chemotactic network

- Sensitivity Gain : output/input ratio
- Adaptation: reset after input
- High amplitude range

Bacteria are social

Bacteria colonize various surfaces (in organisms and external environments) Structural similarity of biofilms growing in hydrothermal hot springs, freshwater rivers and laboratory flow cells.

Hydrothermal hot springs (**a–c**)

Biscuit Basin thermal area, Yellowstone National Park, USA

Biofilms growing in freshwater rivers (**d**,**e**)

Gardener River, Yellowstone National Park, USA (**d**), Hyalite Creek, Bozeman, Montana, USA (**e**),



Hall-Stoodley L, Costerton JW, Stoodley P. Nature Reviews Microbiology 2:95–108 (2004)



Bacteria are social

• First sign of life on earth ~ 3.5B years ago

Stromatolites





Hall-Stoodley L, Costerton JW, Stoodley P. Nature Reviews Microbiology 2:95–108 (2004)



Swarming bacteria

• When swimming bacteria are in contact with a soft surface (eg. <1% agar), they differentiate and move at high density in 2D: they swarm



E. coli



Swarming bacteria gy

- When swimming bacteria are in contact with a soft surface (eg. <1% agar), they differentiate and move at high density in 2D: they swarm
- The bacteria Paenibacillus vortex has swarming motility

Builders grow and divide but are not motile

Explorers are highly motile colonies that spins out into new territories



Ingham, C.J. et al. PNAS 108, 19731–19736 (2011)





10 mm

Social benefits of swarming - species interactions

-Transport of antibiotic resistant bacteria by swarming antibiotic sensitive bacteria

- Paenibacillus vortex is sensitive to Ampicillin
- An *E coli* strain resistant to Amp (Amp^R) survices but cannot expand in presence of Amp
- Swarms of *P. vortex* transports Amp^R *E. coli* and both bacteria form an expanding cooperative colony where *E. coli* detoxifies the environment.
- The colony forms concentric rings with phases of expansion under low Amp, followed by phases of slower expansion when Amp is at higher concentration.
- Bet-hedging strategy: *P. vortex* sub colonies can explore new territory without *E. coli* to avoid competition of hitchhiker. Or expand more slowly with detoxyfier bacteria.
- Co-transport of *P. vortex* (red) and Ctx^R bacteria *Enterobacter aerogenes* (green) in presence of Ctx (cefotaxim).









Finkelshtein, A. et al. MBio 6, e00074-15 (1-10) (2015)



Social benefits of swarming - species interactions

—Dispersal of fungi spores by swarms of bacteria in the soil

- Heterogeneous environment of the plant root (rhizosphere) comprises bacteria and fungi
- The bacteria *Paenibacillus vortex* has swarming motility
- P. vortex transports/advects spores of the fungus Aspergillus fumigatus over >30cm at speeds of 3µm/s and can move them away from sites of adverse growth conditions (eg. anti fungal molecule)
- A. *fumigatus* spores are specifically attached/recognized by bacteria flagella.



• In return, fungi filaments from tracts for the dispersal across gaps of bacteria swarms







Social benefits of swarming - species interactions

-Dispersal of fungi spores by swarms of bacteria in the soil





Swarming mass of *P. vortex* transporting hundreds of ungerminated conidia imaged 4 h after inoculation and 31 mm away from the coinoculation point. (Scale bar: 300 μ m.)

Swarming mass of P. vortex transporting hundreds of ungerminated conidia. Not all conidia are in motion. (Scale bar: 100 μ m.)



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Ingham, C.J. et al. PNAS 108, 19731–19736 (2011)

Social benefits of swarming - interaction with environment

-Superdiffusive transport at swarm upper layer

- The MgO particles on swarm faster than mobile particles ins of the swarm and faster than p migrated on the upper surface lower surfactant activity.
- Superdiffusive behavior of part surface in contact with surfact
- Origin of superdiffusive behavior not clear: heterogeneity in surfactant concentrations (eg. Marangoni flows)?
- Functional benefits?
 - Long range communication by transport of signaling molecules (compared to Brownian motion)
 - Flow of nutrients/oxygen from edges to inner swarms, temperature regulation









A. Be'er and R. Harshey *Biophysical Journal* 101(5) 1017–1024 (2011)

Swarming response: surface translocation of bacteria

• Swimming bacteria that contact a soft surface (eg.Agar) switch to a swarm behavior

— swarming: multicellular movement across surface powered by rotating flagella for swimming bacteria such as *E. coli* or *Salmonella*.

— This induction is reversible (if cells are placed back in solution, they de-differentiate and swim)

- It occurs within 30 min

—In some bacteria, induction of flagellar genes and **increase in number of flagella per cell** Flagella cover the entire cell (peritrichous flagella)

Peritrichous flagella bundle together when they rotate to increase the effective flagellar stiffness and make force generation more efficient in viscous liquids





Historical origins: swarming in Proteus

Differentiation of cells in contact with a soft surface (eg. Agar)

- Proteus mirabilis: Gram negative enterobacteria.
- Commensal of digestive tract in animals
- Present in soil and water
- Responsible of infections of urinary tract in humans
- Swarming behaviour of Proteus mirabilis:
 - Cycles of vegetative and swarming states gives rise to a characteristic concentric colony morphology
 - Cells become longer, filamentous due to multi nucleation (cessation of septation)
 - Development of many (~50) peritrichous flagella which allows better adhesion to substratum and motility

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COLLÈGE

DE FRANCE





John J. Farmer - CDC public health image library

Verstraeten N, et al. *Trends in Microbiology* 16:496–506 (2008) Allison, C, and Hughes, C, *Science Progress* 75: 403-422 (1991)

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Proteus mirabilis Swarm cell (50 flagella)

slime-covered agar (S). and agar without slime (A).

F.D. Williams and R.H. Schwarzhoff. Ann. Rev. Microbial. 32:101-22 (1978)



Historical origins: swarming in Proteus

- Cell differentiation associated with swarming behavior
 - Formation of multinucleate filaments (long bacteria)
 - Growth of multiple flagella (peritrichous)





Swarming behaviors in other bacteria



- Formation of multinucleate filaments (long bacteria)
- Growth of multiple (peritrichous) flagella
- Occurs on soft media (0.5-2% Agar)
- Speed of cell front expansion: 2-10µm/s and up to 30-50µm/s



Harshey RM, Matsuyama T. PNAS 91:8631-8635 (1994)



Diversity of swarming behaviors

- Cell differentiation associated with swarming behaviour
 - Formation of multinucleate filaments (long bacteria)
 - Growth of multiple flagella (peritrichous)



Fig. 3. Transmission sectron microscopy showing cell morphology of vegetative (V) and swarm cells (S) of various bacterial species. (A) Vibrio parahaemolyticus, V; (B) V. parahaemolyticus, S; (C) Serratia marcescens, V; (D) S. marcescens, S; (E) Clostridium sporogenes, V; (F) C. sporogenes, S; (G) Clostridium tetani, V; (H) C. tetani, S; (I) Bacillus cereus, V; (J) B. cereus, S; (K) Bacillus subtilis, V; (L) B. subtilis, S.

Allison, C, and Hughes, C, Science Progress 75: 403-422 (1991)

Bacterium	Flagella arrangement (vegetative/ swarm cell)	Agar concentra- tion permitting swarming (%)	Rate of surface translocation* (µm/min)	
Proteus mirabilis Proteus vulgaris	peritrichous	2.0	950	
Vibrio parahaemolyticus Vibrio alginolyticus	polar/peritrichous	2.0	125	
Serratia marcescens	peritrichous	0.7-0.8	88	
Chromobacterium spp	polar/peritrichous	0.6-1.2	92	
Clostridium tetani Clostridium novyi Clostridium bifermentans Clostridium sporogenes	peritrichous	2.0 2.0 1.0 1.0	670 nd 300 78	
Bacillus alvei Bacillus cereus Bacillus subtilis Bacillus megaterium	peritrichous	2.0 1.8 1.1	120 108 80 560	



Diversity of swarming behaviors

Macroscopic manifestations of swarming behaviors

The macroscopic appearance of a swarm colony:

- differs among bacterial species,
- depends on medium composition,
- agar concentration and water content,
- temperature.



 a Featureless mat
 b Bull's eye (Also known as zones of consolidation or terraces)
 c Dendrites (Also known as deep branches or tendrils)
 d Vortex (Also known as wandering colonies)

 Image: Consolidation or terraces)
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Fig. 4. Swarming by various bacterial species on agar-containing media, (A) Vibrio parahaemolyticus, (B) Vibrio alginolyticus, (C) Serratia marcescens,

(D) Clostridium sporogenes, (E) Clostridium tetani, (F) Clostridium bifermentans, (G) Bacillus cereus, (H) Bacillus megaterium, (I) Bacillus subtilis.

Allison, C, and Hughes, C, Science Progress 75: 403-422 (1991)

Widespread « invention » of swarming in bacteria lignages

This suggests important selective advantage of swarming behavior to colonize environments





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Harshey RM. Molecular Microbiology 13: 389-394. (1994)

Transition to swarming state: characteristics



- 1. Robust swarmers
 - Swarm on hard agar (>1.5% agar)
 - Cell elongation
 - Hyperflagellation: polar or peritriche

Azospirillum, Rhodospirillum, and Vibrio species, and Proteus





Swarmer phenotype 2

Increased cell density Cell alignment Cell elongation

- Production of osmolytes or surfactants
- Hyperflagellation (in some bacteria)

2. Temperate swarmers

- Swarm on soft agar (0.5-0.8% agar)
- No (or little) cell elongation
- No hyperflagellation

E. coli and Bacillus, Pseudomonas, Rhizobium, Salmonella, Serratia, and Yersinia species

N Wadhwa and HC. Berg. Nature Rev. Mol. Cell Biol. 20: 161-173 (2022)



Transition to swarming state: characteristics

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Slowed 2.5x

Serratia marcescens

Salmonella enterica



Swarming state: Physical constraints



- 1. Maintaining a hydrated, fluid medium
 - Osmotic regulation of fluid uptake

2. Reducing and opposing frictional forces

- Substrate interactions
- Propulsion by flagella
- Cell length and fluid drag
- 3. Reducing Surface tension: wettability
 - Surfactants and other wetting agents



WATER /



Partridge, JD. and Harshey RM. J. Bacteriology 195: 909-918. (2013)



Transition to swarming state: fluidization

- Secretion of osmolytes by bacteria give rise to water osmotic flow from agar to the bacteria media
- This decreases the viscosity of the bacteria swarm and favors motility



Gram negative bacteria secrete osmolytes, polysaccharides to draw water from agar into the swarm

- In *Proteus mirabilis, there is* secretion of extracellular matrix (ECM) composed of polysaccharides, the osmolyte glycine betaine, etc.
- Acidic polysaccharide Cmf (colony migration factor) is essential for swarming and supposed to increase colony hydration.
- Some swarmers (*P.Aeruginosa*) up regulate the osmolytes glutamate and proline.
- Evidence of altered metabolome in bacteria at the surface of agar. This induces synthesis and modification of lipopolysaccharides (LPS)

Partridge, JD. and Harshey RM. J. Bacteriology 195: 909-918. (2013)

Lahaye E, Aubry T, Fleury V, Sire O. Does water activity rule P. mirabilis periodic swarming? II. Viscoelasticity and water balance during swarming. *Biomacromolecules* **8**:1228–1235. (2007)



Transition to swarming state: fluidization

Observation of fluid flows at the edge and inside a swarm of *E. coli* Prediction of osmotic fluid flows from the agar into the swarm

В

80

40 60

typical bubble tracks in the laboratory frame

-60

-70

A

15

Microbubbles fluid flow profiles as a function of distance from the edge of the swarm edge

Cell growth and metabolic activity is



 $<r_x(t) > (\mu m)$



Wu, Y., and H. C. Berg. Proc. Natl. Acad. Sci. USA. 109:4128–4133. (2012)

Transition to swarming state: fluidization

Osmolarity is patterned at the swarm edge consistent with predicted osmotic fluid flow

Wu, Y., and H. C. Berg. Proc. Natl. Acad. Sci. USA. 109:4128-4133. (2012)

Secretion of osmolytes by bacteria gives rise to water osmotic flow from agar to the bacteria media

Liposomes are prepared with 2 fluorescent dyes: one that self-quenches when liposomes shrink (G, green dye calcein) and one that does not (R, red dye sulforhodamine-101).

The G/R ratio reflects the osmolarity of the surrounding fluid.

Measure osmolarity as swarm of *E. coli* cells spread across large liposome ($450 \mu m$)

Two bands: high osmolarity at the edge (+25mOsm) and lower osmolarity behind, isotonic to agar



Ping et al. and HC. Berg. Biophysical Journal 107:871-878 (2014)



Transition to swarming state: Overcoming friction

- Reducing and opposing frictional forces
 - Substrate interactions
 - Propulsion by flagella
 - Cell length and fluid drag



- Substrate interaction:
 - role of charges (could be regulated by LPS)
 - biosurfactants are lubricants due to amphipathic (hydrophilic and hydrophobic) structure that could reduce friction. LPS (gram- bacteria), rhamnolipids (*Pseudomonas*) and lipopeptides have surfactant properties.
- Propulsion by flagella:
 - Hyperflagellation associated with increase in body length. The increased propulsion forces counteract friction
 - Specific stator units that increase active torque on rotor/flagella complex PG

P. aeruginosa has five stator proteins (MotAB, MotCD, and MotY), which likely associate into two sets of stators, whereas only two such proteins (MotAB) drive swimming motility in *E. coli* and *Salmonella*.



A. Baker and G. O'Toole. J. Bacteriol. doi:10.1128/JB.00088-17 (2017)





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N. Wadhwa, R. Phillips and HC. Berg. PNAS. 116:11764-11769 (2019)

Transition to swarming state: Overcoming friction

• Bacteria contact with a soft surface is associated with induction of hyper flagellation

—Induction of flagellar genes and increase in number of flagella per cell Flagella cover the entire cell (peritrichous flagella), eg. From 5 to 50 flagella.





Unsheathed lateral flagella (Laf genes)

L. McCarter et al. Cell 54: 345-351 (1988)



Sheathed polar flagellum (Fla genes)

Transition to swarming state: hyperflagellation

- -When bacteria sense a more viscous environment at a surface they differenciate
- Cells detect physical properties of the surface
- The induction of lateral flagella is regulated mechanically
- The polar flagella works as a dynamometer/viscometer
- Reporter of *lab* gene expression with light
- Increasing the viscosity of the medium induces Laf expression
- Blocking flagellar rotation with an antibody also induces Laf
- Laf is constitutively induced in bacteria in solution when the flagellum is defective (flagellin C or mot mutant)





L. McCarter and M. Silverman *Cell* 54: 345-351 (1988)L. McCarter and M. Silverman *Mol. Microb.* 4(7), 1057-1062 (1990)

Transition to swarming state: surfactants and wetting

Reducing Surface tension: wettability

- Surfactants and other wetting agents
- Hypothesis: Gradient in wetting agents generates Marangoni stresses and capillary flows that drive swarm expansion.







- The agar substrates is partially hydrophobic.
- Need of wetting agents, eg. amphipathic molecules.
- -Lipopeptides and rhamnolipids produced by Serratia, Bacillus, Pseudomonas etc
- Cyclic lipopeptides: Surfactin produced by *B. subtilis* and serrawetin produced by *S. marcescens*.
- Surfactant deficient bacteria loose swarming behavior. This can be rescued by adding exogenous surfactant.



rhamnolipid



A theoretical model of swarming



Governing equations based on mass and momentum conservation:

$$((h\phi))_t + (Q_1(x))_x = g_1(h,\phi,c),$$

$$(h(1-\phi))_t + (Q_2(x))_x = (1-\phi)V_0(x),$$

$$c_t - Dc_{xx} = g_2(h, \phi, c).$$

 $oldsymbol{\phi}_1 = oldsymbol{\phi}(x,t) \;\;$ is the active (cell) phase

 $Q_1(x)$ is the horizontal flux in the active phase, $Q_2(x)$ is the horizontal flux in the fluid phase and $V_0(x)$ is the osmotically-driven net vertical fluid influx per unit length across the permeable substrate

 $g_1(h;c;\phi)$ is the depth integrated active phase growth rate within the bacterial colony, and $g_{2(}h;c;\phi)$ is the depth integrated nutrient uptake rate



Srinivasan et al. and L. Mahadevan eLife 8:e42697. DOI: https://doi.org/10.7554/eLife.42697 (2019)



Motility properties of swarming cells





Motility properties of swarming cells

-The swarm monolayer acts like a 2D gas of self- propelled, polar particles.

• The density profile of *E. coli* cells in the swarm varies with distance from the edge.

Highest density is just behind the edge

- Cells at highest density region have higher motility and motility is broadly distributed (compared also to swimming bacteria)
- Speed is the same as in swimming bacteria
- Cell movement is mostly linear (low curvature) with a narrow propulsion angle compared to swimming bacteria that undergo tumbling and nearly random reorientation







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N. Darnton et al and H. Berg. Biophysical Journal 98(10) 2082–2090 (2010)

Motility properties of swarming cells

-The swarm monolayer acts like a 2D gas of self- propelled, polar particles.

- Microscopic properties of the « bacterial atoms of this 2D gas », such as speed distributions and correlation functions.
- Cells move in coordinated packs
- Cells tend to align laterally but less so at front/back





N. Darnton et al and H. Berg. Biophysical Journal 98(10) 2082–2090 (2010)



Motility of swarming cells

Cell tracking and time projections of cell positions Cell body: head (blue), tail (green), middle (red) Flagella (light blue)

- Forward motion
 - Bundles of flagella are aligned with long axis of the cell and propel it forward at the same speed as free swimmers in 3D (few 10 µm/s)



- Reversals
 - Cells reorganize the flagella as they stop, and rebundle them in the opposite orientation so change in direction is mostly a reversal













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L. Turner et al and H. Berg. Journal of Bacteriology, 192: 3259-3267 (2010)

Motility of swarming cells

- Lateral motion and Collisions
 - Cell collisions are responsible for lateral motion of cells and for the wide distribution of velocities



- Cell alignement
 - Cells tend to align, most likely via entrainement due to motion, interactions among flagella, and the cell elongation (long aspect ratio of cells).
 - Not due to co-bundling of flagella





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L. Turner et al and H. Berg. Journal of Bacteriology, 192: 3259-3267 (2010)

Motility of swarming cells

- Comparison of swimmers and swarmers: Similarities
 - Cells run, stall and re-run in both cases and this is due to a transition from bundled to « curly » flagella presumably due to a transition from CCW to CW rotation
 - In bot cases, run are powered by bundled flagella.

Differences

- Swimmers have a random walk in 3D
- Swarmers have a constrained movement in 2D.
- Tumbling is frequent in swimmers but rare in swarmers.
- Contrary to swimmers, swarmer cells are not free to change direction at random after they stall/stumble and undergo a new run.
- Rather cells tend to align and to reverse in the opposite direction. This is due to the high density of cells.







Other modalities of swarming...

Ente

eriaceae



TABLE 1 Main features of various types of surface motility

Types of motility	Motive organelles	Cell differentiation	Colony expansion rates (µm/s)	Function	Bacterial genera ^a
Swarming	Flagella	Yes	2–10	Surface colonization	Aeromonas, Azospirillum, Bacillus, Clostridium, Escherichia, Proteus, Pseudomonas, Rhodospirillum Salmonella, Serratia, Vibrio, Yersinia
Twitching/ social gliding/ retractile motility	Type IV pili	bili No 0.06–0.3 Surface colonization, Ae biofilm formation, A fruiting body B development, B bphage infection, A	Aeromonas, Acinetobacter, Azoarcus, Bacteroides, Branhamella, Comomonas, Dichelobacter, Eikenella, Kingella, Legionella,		
				conjugation	Noiaxetta, Myxococcus, Neisseria, Pasteurella, Pseudomonas, Ralstonia, Shewanella, Streptococcus, Suttonella, Synechocystis, Vibrio, Wolinella

Harshey RM. Molecular Microbiology 13: 389–394. (1994)



Solitary motility of un-flagellated gliding bacteria

Bacteria: Myxococcus xanthus: $l = 5 \mu m$



A. Pelling et al. PNAS, 102: 6484–6489 (2005)



speed: 2-4 µm/min



Gliding motility: helicoidal rotation and adhesion



L. Faure et al, and T. Mignot Nature, 539:530-535 (2016)

N Wadhwa and HC. Berg. Nature Rev. Mol. Cell Biol. 20: 161-173 (2022)



Collective motility of un-flagellated bacteria

Myxococcus xanthus:

adventurous (solitary) motility: exploration

social motility (predation)



speed: 2-4 µm/min

www.youtube.com/watch?v=tstc6doiNCU

Treuner-Lange, MPI Marburg



Collective motility of un-flagellated bacteria

Myxococcus xanthus, Pseudomonas aeruginosa. And many more



Collective motility of un-flagellated bacteria



- Rippling waves are made of multilayered stacks of bacteria.
- Waves propagate, collide and reflect, in spite of their apparent persistent motion
- Individual bacteria do not follow the apparent dynamics of waves: they exhibit oscillatory motion on the substrate

Tâm Mignot lab



Oscillatory dynamics of un-flagellated swarming bacteria





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Tâm Mignot lab

Layer formation in bacteria colonies: Topological defects



Un-flagellated bacteria swarm via twitching motility

1.0 h

Pseudomonas aeruginosa move forward and reverse by twitching

- Bidirectional (oscillatory) motion in the bulk
- More directional motion at the edge



Kühn, M. J. et al. and A. Persat. PNAS 118, e2101759118 (2021).



50 µm

Cycles of pili extension and retraction, adhesion to a self-secreted matrix, drive forward and reverse movement



The pilus functions as a grappling hook

- Pseudomonas aeruginosa presents multiples thin filaments called pili at one pole of the bacterium.
- Type IV pili undergo cycles of rapid extension and retraction at rates of ~0.5-1 μm s⁻¹









JM. Skerker and HC. Berg. PNAS 98: 6901–6904 (2001)

N Wadhwa and HC. Berg. Nature Rev. Mol. Cell Biol. 20: 161-173 (2022)

Cycles of pili extension and retraction, adhesion to a self-secreted matrix, drive forward and reverse movement



D. Kaiser. Nature Rev. Microbiol. 1: 45-53 (2003)

- Cells secrete exopolysaccharides that enhance adhesion to the substrate (agar)
- Reinforcement by feedback of cell motility on tracks used by other bacteria.
- The environment thereby guides the bacteria: concept of stigmergy (PP. Grassé, *Insectes sociaux* 1959)

LA RECONSTRUCTION DU NID ET LES COORDINATIONS INTERINDIVIDUELLES CHEZ BELLICOSITERMES NATALENSIS ET CUBITERMES SP. LA THÉORIE DE LA STIGMERGIE : ESSAI D'INTERPRÉTATION DU COMPORTEMENT DES TERMITES CONSTRUCTEURS.

par Plerre-P. GRASSÉ

La coordination des tâches, la régulation des constructions ne dépendent pas directement des ouvriers, mais des constructions elles-mêmes. L'ouvrier ne dirige pas son travail, il est guidé par lui. C'est à cette stimulation d'un type particulier que nous donnons le nom de STIGMERGIE (stigma, piqûre ; ergon, travail, œuvre=œuvre stimulante).









S.T. Islam, T. Mignot / Seminars in Cell & Developmental Biology 46 (2015) 143-154

Pili extension and retraction

\rightarrow	Type IV pilus
	Retracting pilus
Solid surface	



- Type IV pilus machinery is closely related to the type II secretion system of Gram-negative bacteria and the archaeal flagellar machinery
- Type IV pili are ubiquitous in bacteria and archaea and mediate adhesion, DNA uptake, predation, virulence, phototaxis, chemotaxis and surface sensing.
- Single pili are polymers of PilA, or Pilin monomers assembled from the cytoplasmic side.
- PiIT and PiIB at hexameric motors that induce assembly and disassembly of PiIA, and thereby elongation and retraction of the pili.
- The is an energy consuming (ATP dependent) active process.



N Wadhwa and HC. Berg. Nature Rev. Mol. Cell Biol. 20: 161-173 (2022)



Pili retraction powers bacterial twitching motility: evidence

- Neisseria gonorrhoeae are spherical ~lµm long bacteria that undergo twitching motility via pili.
- Cells form small swarms
- Cells move at a speed of ~ I $\mu m/s.$





Pili retraction powers bacterial twitching motility: evidence

- Use optical tweezer to hold a cell at the edge of a small colony
- The cell moves towards the colony and escapes the trap revealing retractile forces between cells.

Cells usually move back towards the trap revealing the transient nature of retractile force

b

- Cells immobilized on a bead anchored to the substrate:
- An optical tweezer was used to hold a cell a few microns away and measured the retractile forces
- Retractile forces are in the range of I00pN







A. Merz, M. So and M. Sheetz. Nature 407:98-102 (2000)

• Oscillatory dynamics of twitching cells

- *P. aeruginosa* shows oscillatory dynamics but exhibit biased direction.
- This involves in part chemotaxis.
- The ChpA (Chemosensory protein A) sensory system is similar to the CheA/Y system (in *E. coli*) and is likewise involved in chemotaxis









Kühn, M. J. et al. and A. Persat. PNAS 118, e2101759118 (2021).



- Twitching cells detect collisions and reverse motility
- The ChpA/PilG/PilH sensory system is required for the detection of collision
- Detection of collision allows uniform spreading and density of bacteria in swarms.
- consistent with Cells detect a syr • mechanotaxis instead of chemotaxis













initial leading pole



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Kühn, M. J. et al. and A. Persat. PNAS 118, e2101759118 (2021).

• PilB and Pili tend to concentrate at the same cell pole

PiliB presents a polarized distribution at the poles of *P. aeruginosa*. PilB is concentrated at the pole where Type 4 Pili are also concentrated



Pili:T4P

Kühn, M. J. et al. and A. Persat. PNAS 118, e2101759118 (2021).



• PilB and Pili concentrate at the leading, forward moving cellular pole

The localization of the extension motor PilB sets the direction of twitching and the polarization of Pili activity The retraction motor PilT has no biased distribution





90% of cells move towards PilB pole





Kühn, M. J. et al. and A. Persat. PNAS 118, e2101759118 (2021).

Model of mechanotaxis in *P. aeruginosa*

0

0 Normalized distance from midcell



Kühn, M. J. et al. and A. Persat. PNAS 118, e2101759118 (2021).

 Chp mechanosensing induces a positive feedback on T4P motors to favor extension at the

same pole leading to polarization of the cell and persistent forward motion

After collision (or loss of T4P attachment/retraction) at the leading pole, a negative feedback down regulates T4P activity. T4P at the opposite pole can then attach and generate a positive feedback that reverses cell polarization and lead to persistent reverse twitching.

COLLÈGE DE FRANCE

Implications for swarming dynamics in P. aeruginosa



- Hypothesis:
- Chemotaxis:
 - The swarm expands towards a chemoattractant that lowers the frequency of reversal. Persistant random walk in ID/2D.
- Mechanotaxis:
 - The density of collisions is expected to be lower at the edge of the swarm and favors persistent twitching down the density gradient
 - In the bulk of the swarm, reversals are more frequent and collision detection causes a more uniform density distribution.



Oscillatory dynamics of *Myxococcus*





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Oscillatory dynamics of twitching Myxococcus

• A module of cell polarization driving motility

• Cell polarity:

MgIA, a Ras family GTPase, is localized in the leading pole and MgIB in the lagging pole.

• Control:

Accumulation of MgIA at the leading pole is controlled by:

- The RomRX complex which works as a GEF, that produces MgIA-GTP and allows its polar recruitment
- and by MglB which works as a GAP and inhibits MglA
- MgIA, RomRX and MgIB define a polarity module that controls the direction of movement.
- During reversal, MgIA relocalizes to the opposite pole, switching the polarity axis
- This allows the cells to move in the opposite direction.





Oscinatory dynamics of twitching Myxococcus

- An oscillatory module underlies reversai[™]
- MgIA, and MgIB are relocalized at opposite poles during reversal.
- This depends on RomRX oscillations:
 - RomRX recruits MgIA to leading pole and subsequently dissociates
 - RomRX relocalises to and primes the lagging pole
- Oscillatory dynamics based on delayed negative feedback.
 The refractory period is set by the time scale of RomRX relocalization
- Yet, RomR:GFP relocalization does not coincide strictly with reversal, suggesting another signal to trigger reversal.





P

MgIA

MglB

Guzzo M, et al. and M. Howard, T. Mignot Nat Microbiol 3:948-959 (2018)

Oscillatory dynamics of twitching Myxococcus

• A gated relaxation oscillator: a switch and an oscillator

- FrzX inhibits MglB and works as a SWITCH
- Once RomRX is in lagging pole, the lagging pole is primed for activation by FrzX, which inhibits MglB and allows MglA to be recruited, thereby causing reversal
- FrzX reduces the RomR threshold and the effective refractory period for reversal





Current Opinion in Cell Biology



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Guzzo M, et al. and M. Howard, T. Mignot Nat Microbiol 3:948-959 (2018)

Potential implications for swarming *Myxococcus*

Hypothesis:

Potential implications for swarming dynamics:

- Each cell has an intrinsic oscillator that induces reversals
- This oscillator is gated by a signal (FrzX)
- Presumably collisions induce/activate FrzX and thereby collisions could bias the oscillatory dynamics of twitching cells
- The frequency of collisions would bias the direction of movement.
- Rippling waves are associated with reflection of cells via this mechanism
- On the edge of a swarm where cell density is lower, cells could be biased down the density gradient towards regions with fewer cells, allowing expansion of the swarm (similarities with *Pseudomonas*)



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Similarities between Pseudomonas and Myxococcus



- T4Pili dynamics underlies twitching
- PilB and PilT ATP dependent motors
- Cell reversal induced by collisions.

• Similar regulation

- ChpA is a homologue of FrzE
- They regulate PilG/PilH, response regulators similar to FrzZ/FrzE (CheY type)
- PilG controls the motors (PilB) in Pseudomonas, while FrzZ/X control MgIA/B in Myxococcus
- MgIA controls Pili via other proteins

Mercier R, et al. & Mignot T. PNAS 117(45):28366-28373 (2020) Bautista S, et al, Mignot T, Mercier R. *EMBO J*. 8:e111661 (2022)



Regulation — Mechanics



Emergence of collective dynamics in swarms of confluent cells

Jamming





www.youtube.com/watch?v=IZupwFOhjl4

Hi Chi Minh city - Vietnam

• Collective dynamics



Slowed 2.5x

Bacillus subtilis swarm Partridge, JD. and Harshey RM. *J. Bacteriology* 195: 909-918. (2013)



Confluent MDCK cell monolayer

D. Cohen et al PNAS 113: 14698–14703 (2016)



Comparisons between eucaryotes & procaryotes

- <u>Eukaryotes:</u>
- Following collisions, cells reorganize their polarities to induce local ordering
- high density induces mechano-chemical coupling that allows global symmetry breaking (leaders) or local symmetry breaking (boundary and leader free).
- Positive density-dependent regulation of motility: coordination of velocity (see Toner Tu model course #1)
- <u>Prokaryotes</u>:
- Following collisions, cells move in opposite directions
- Density dependent biased random (oscillatory) ID walk
- Negative density-dependent regulation of motility?
- But collisions and possible lateral adhesion to promote local ordering?
- Lateral alignement based on morphology of rigid body
- Mechanotaxis: stiffness
- Chemotaxis

