

SHOULD I PAUSE OR SHOULD I GO

A c-di-GMP mediated motility arrest before cell division in *Myxococcus xanthus*

Prunelle Carcassonne¹, Leon Espinosa², Tâm Mignot¹ and Dorothee Murat¹

1. Laboratoire de Chimie Bactérienne, Equipe Mignot

2. Laboratoire de Chimie Bactérienne, Equipe IAM-Instrumentation Analyse et Microscopie

Myxococcus xanthus has a complex life cycle including multicellular stages

M. xanthus is a ubiquitous soil bacterium and a model organism studied for its social behaviors and complex life cycle. Motility plays a central role in this life cycle, allowing *M. xanthus* to predate on other microorganisms. As a facultative predator, *M. xanthus* can use available nutrients in its milieu as well as kill and feed on nearby preys, thus allowing *M. xanthus* growth and division. When resources become scarce, *M. xanthus* cells converge and aggregate to form multicellular structures called fruiting bodies, which can sporulate again when conditions improve to resume vegetative growth.

Unlike other extensively studied life processes of *M. xanthus*, cell division remains poorly characterized

Before dividing, *M. xanthus* undergoes a pause (Harvey et al. 2014). Motility and its regulation are well-studied in *M. xanthus*. However, functional correlation between motility and cell division is not understood.

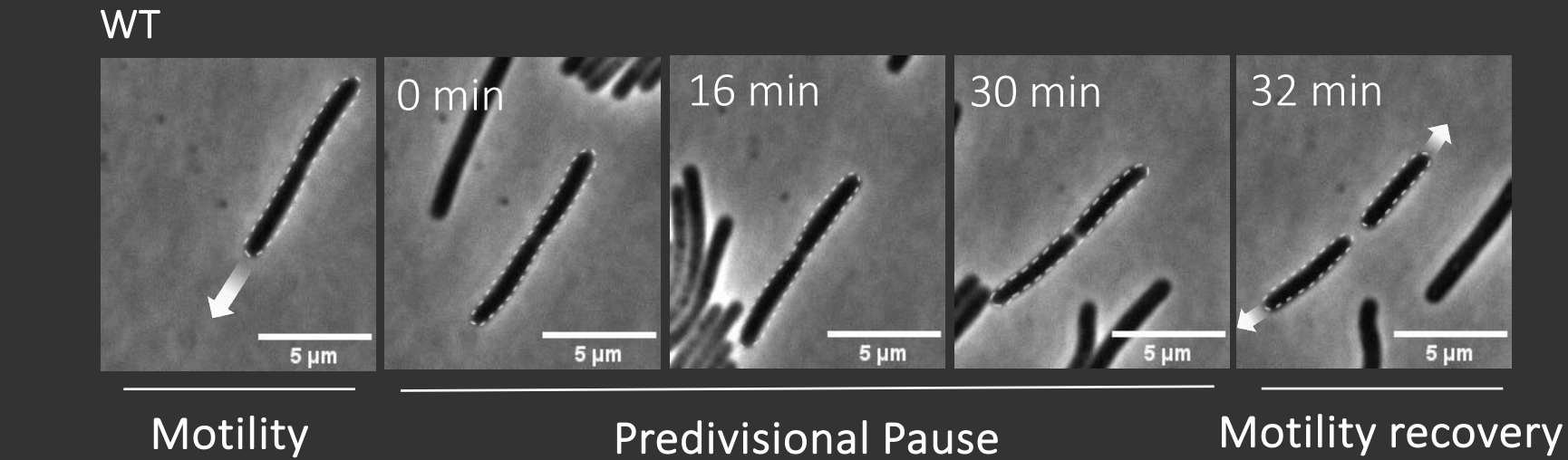
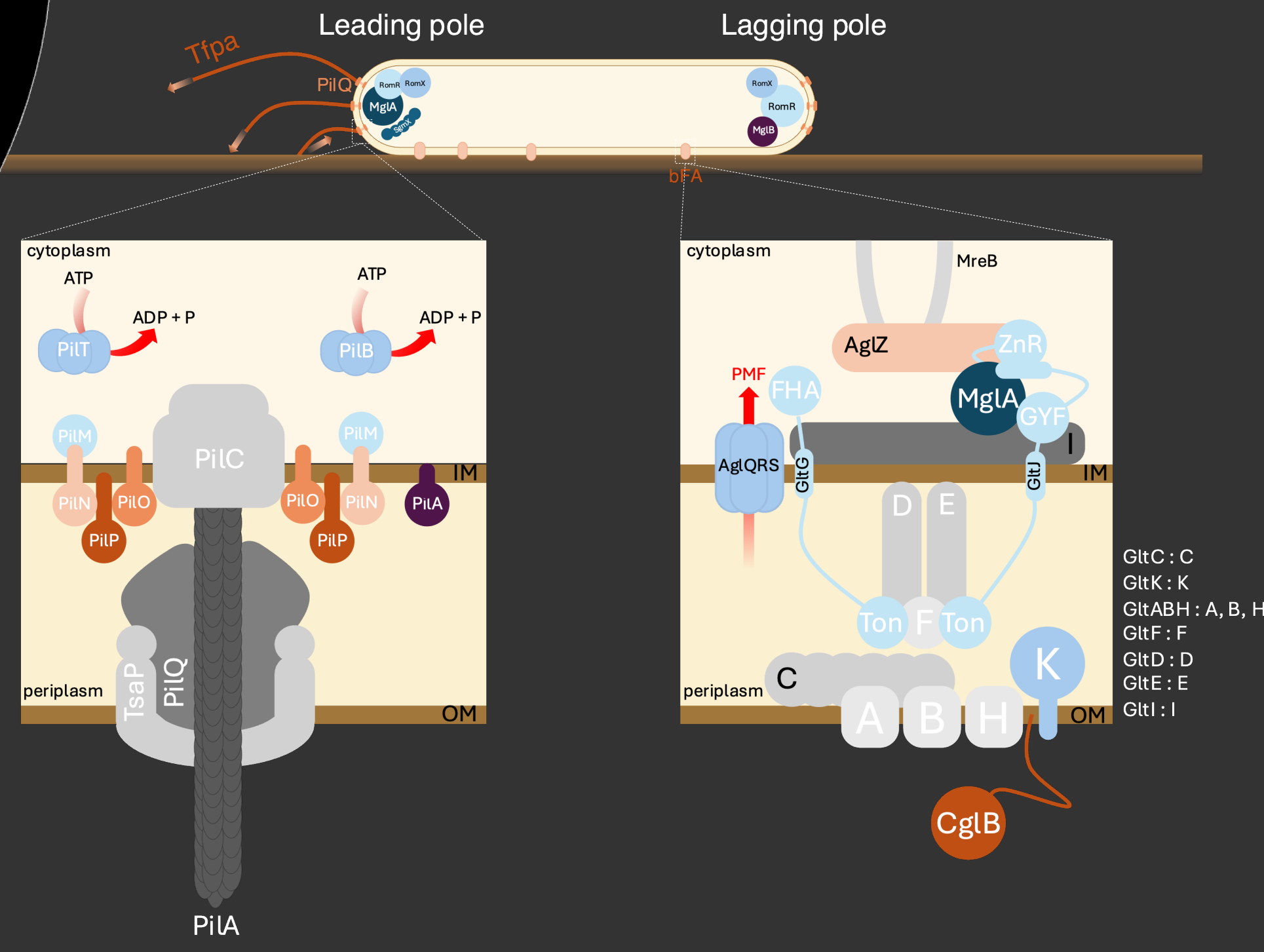
M. xanthus employs a unique PomXYZ protein complex that directly positions and stimulates cytokinetic FtsZ-ring formation.

$\Delta pomZ$: presents division defects with the formation of chromosome-free minicells and filamentous cells. Lack of PomZ also caused reduced formation of Z-rings and incorrect positioning of the few Z-rings formed. (Treuner-Lange, Aguiluz et al. 2013).

Surprisingly, It is not essential, even though it is the only known factor involved in FtsZ localization and recruitment.

M. xanthus has two motility systems

It possesses two distinct motility systems : Twitching via type IV pili (social motility) and gliding with an Agl-Glt complex linking an adhesin to the cytoskeleton (Adventurous motility).



There is a correlation between motility systems and cell division

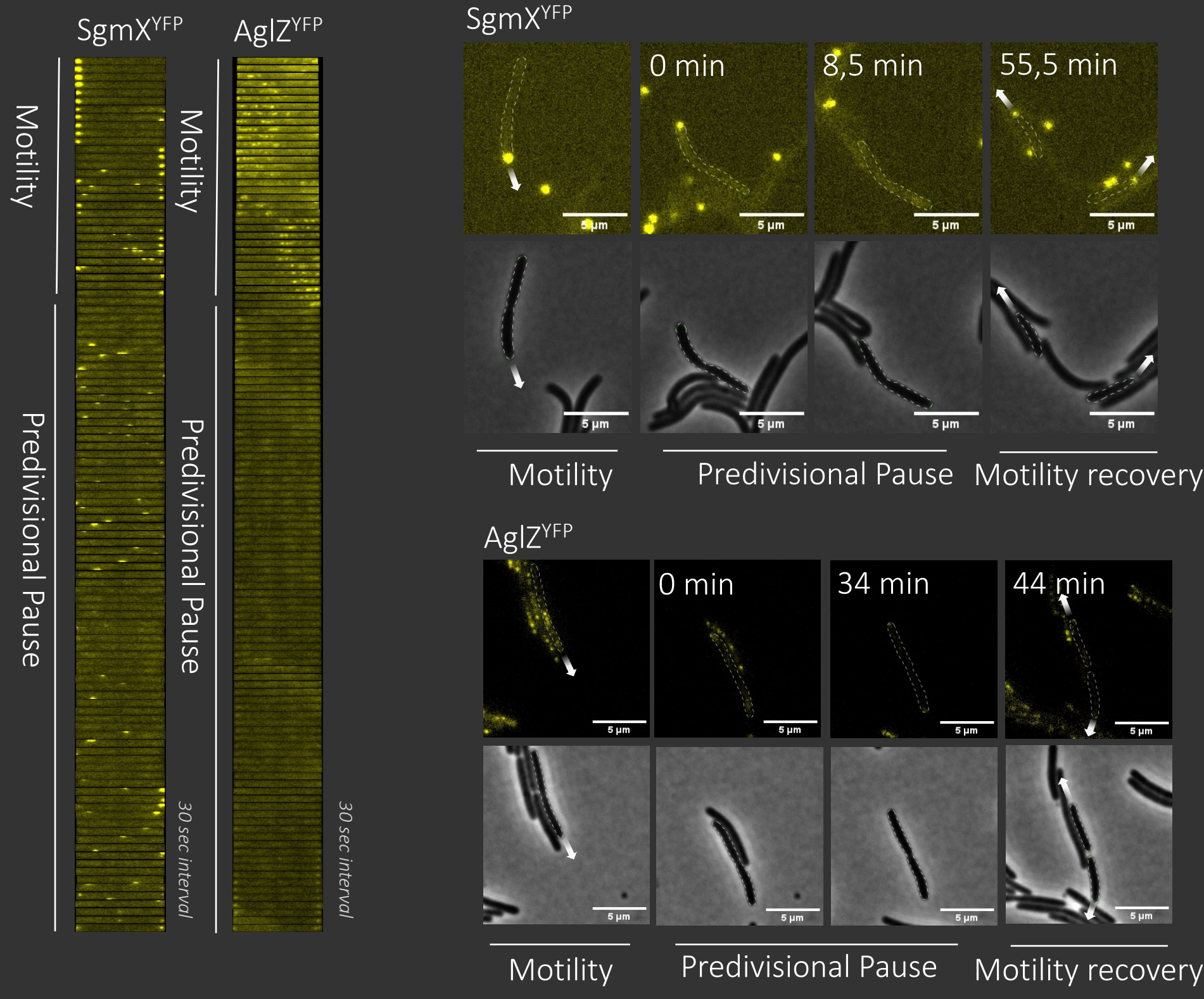
Does this suggest the existence of an additional player?

How are cell division and motility coordinated in *Myxococcus xanthus* ?

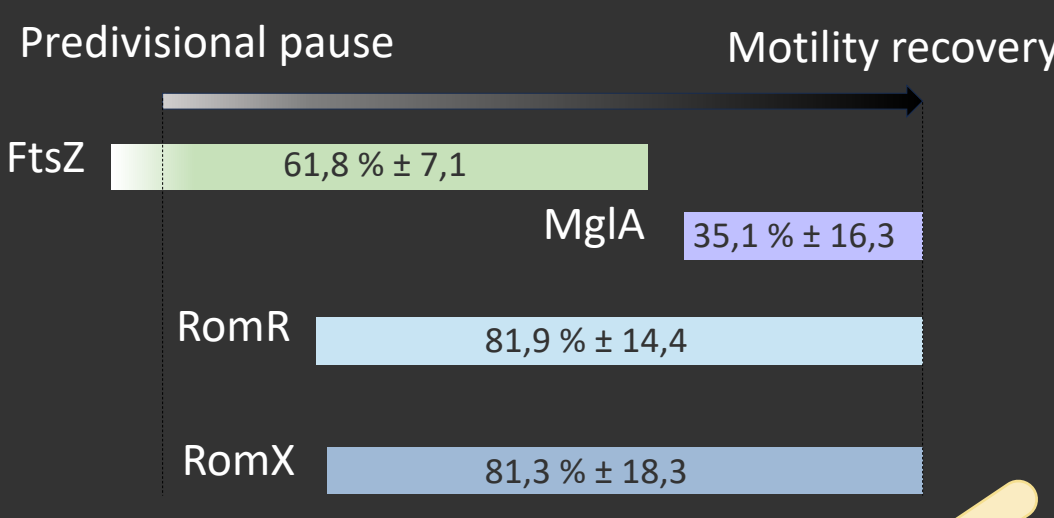
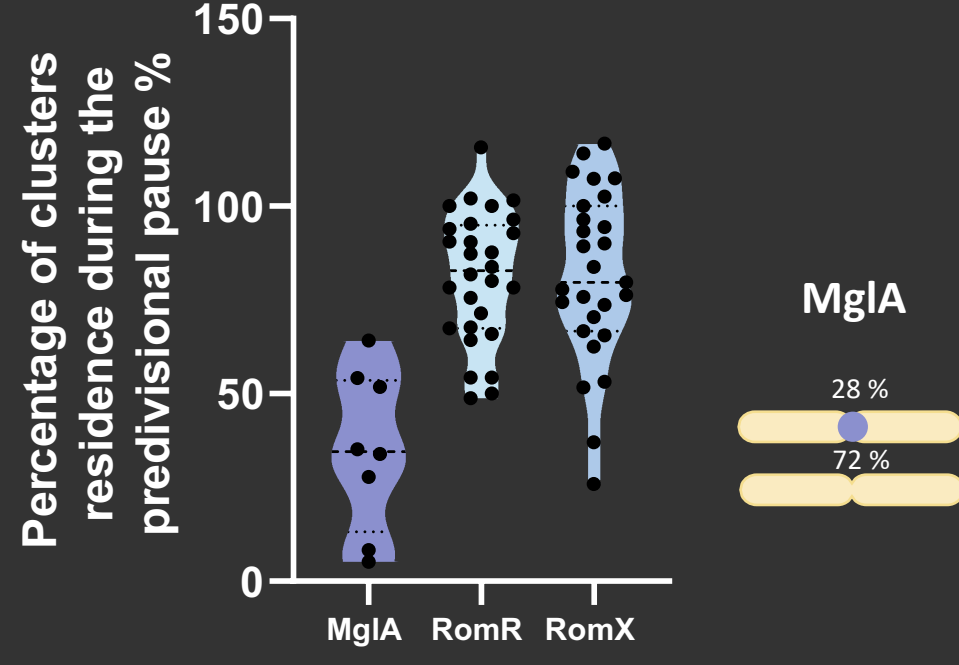
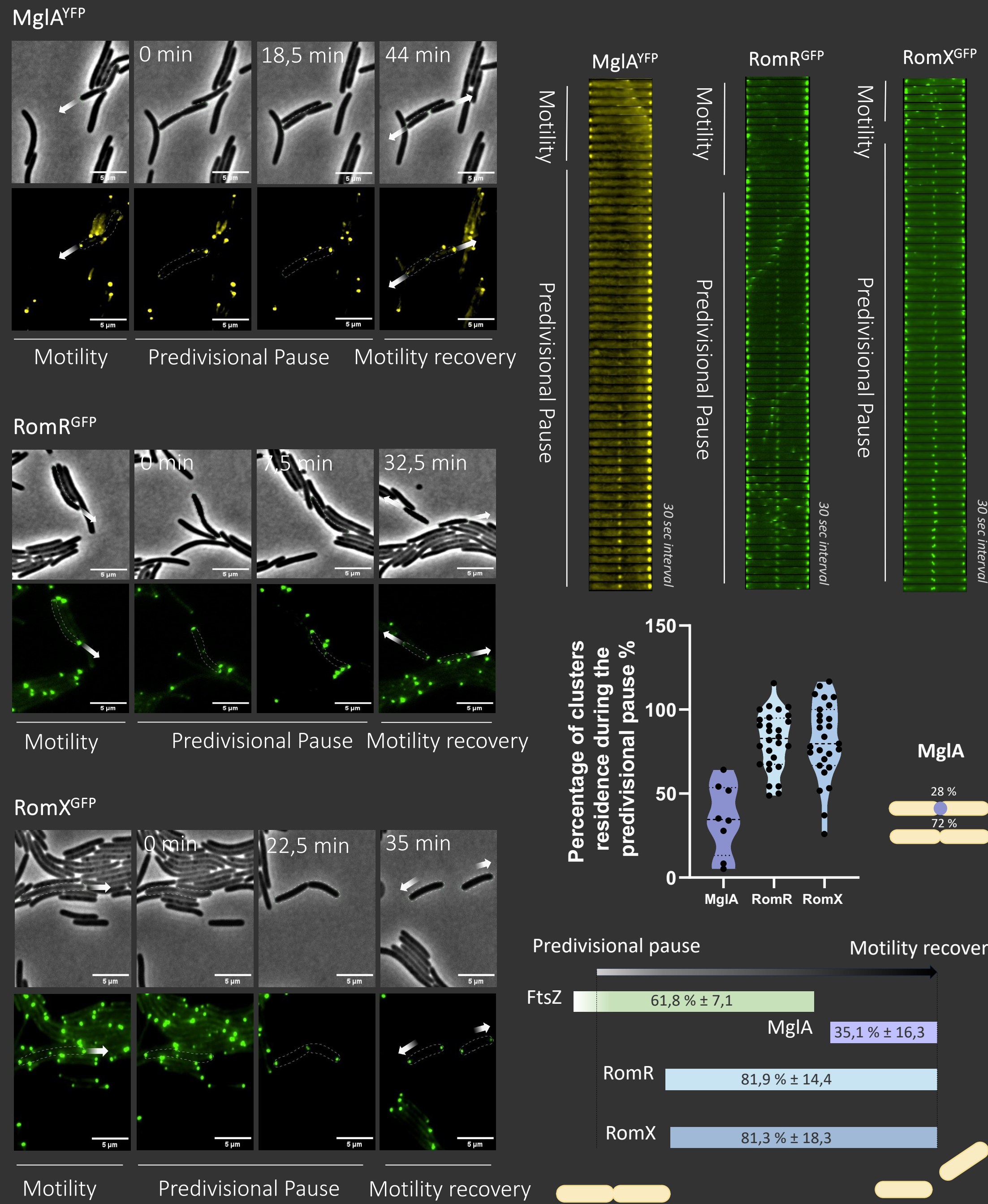
1 MOTILITY PROTEINS DURING CELL DIVISION

Locating and capturing division events in *M. xanthus* is difficult. Indeed, this bacterium is motile and has a doubling time of 5 hours. However, cell division is favored in a predation context. (Panigrahi et al. 2021)

Motility systems are disassembled during predivisional pause



But three motility regulators are recruited at the future division site



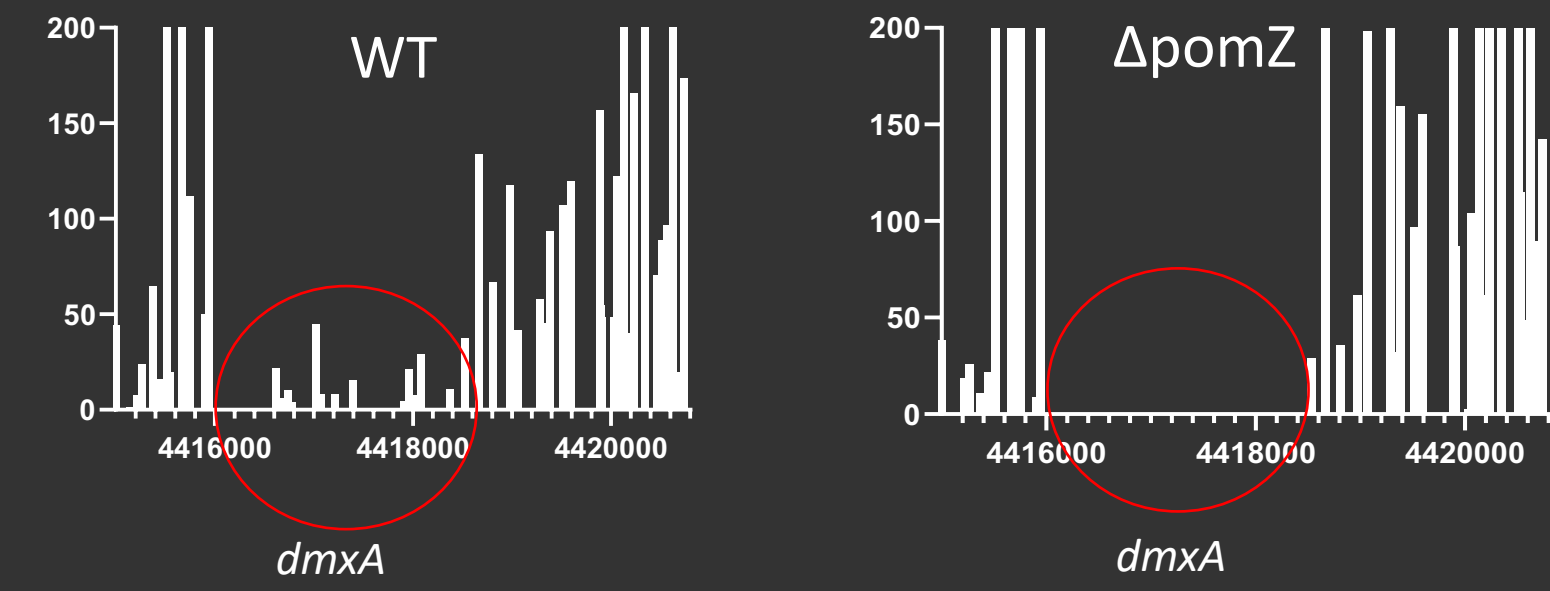
Neither the Agl-Glt complex of A-motility nor the type IV pili of S-motility are assembled in predivisional cells. However, three regulators of motility, RomX, RomR and MglA are recruited at midcell.

There is a successive recruitment of motility proteins at the division septum, suggesting a correlation between the two processes

2 DISCOVERING OF NEW PLAYERS IN CELL DIVISION USING TN-SEQ

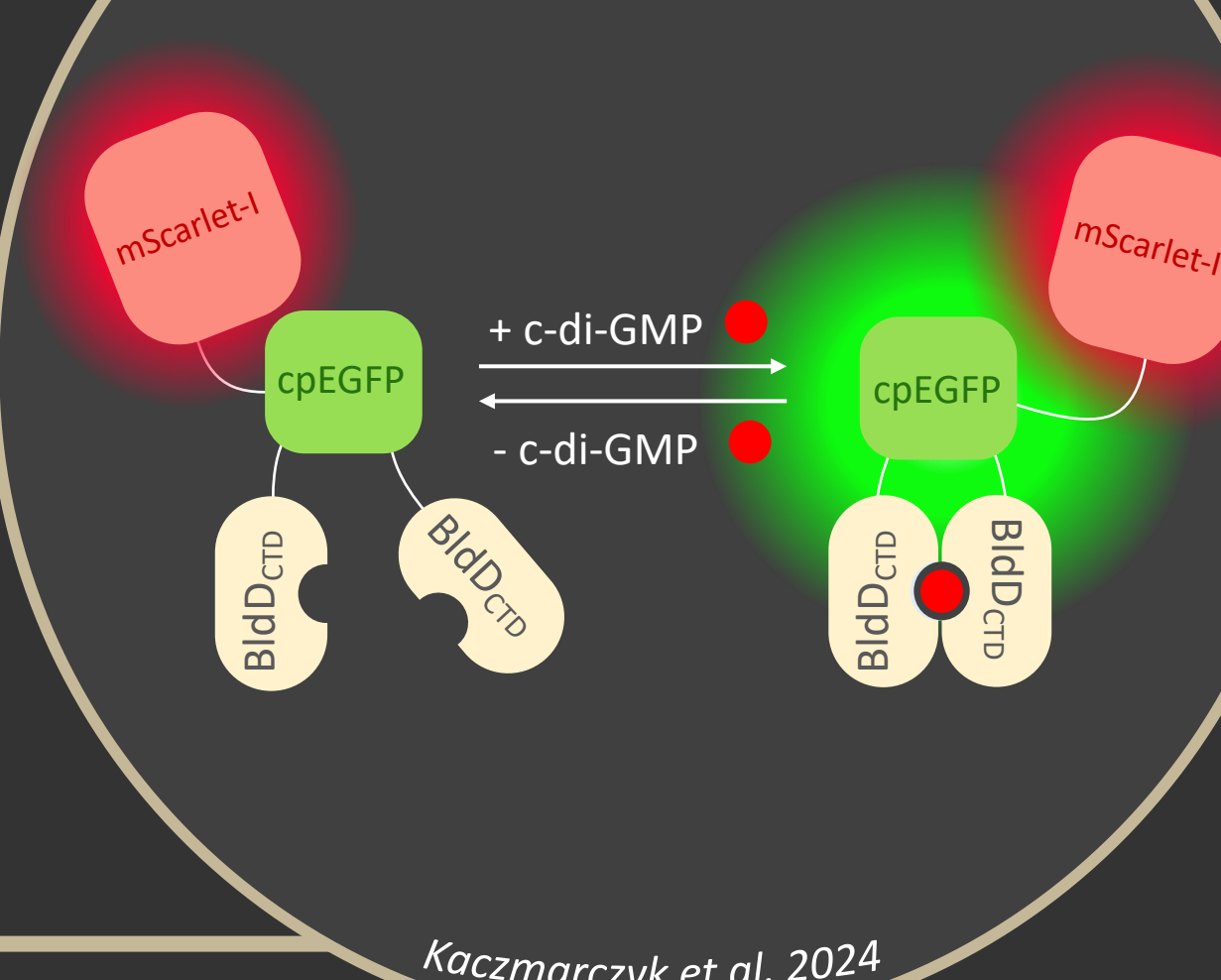
Fifty mutants are less abundant in the $\Delta pomZ$ library, suggesting synthetic lethality and potential functional redundancy.

One of these mutants is known to play a role in the distribution of motility proteins after cell division in *Myxococcus xanthus* $dmxA$. (Maria Pérez-Burgos, 2024)



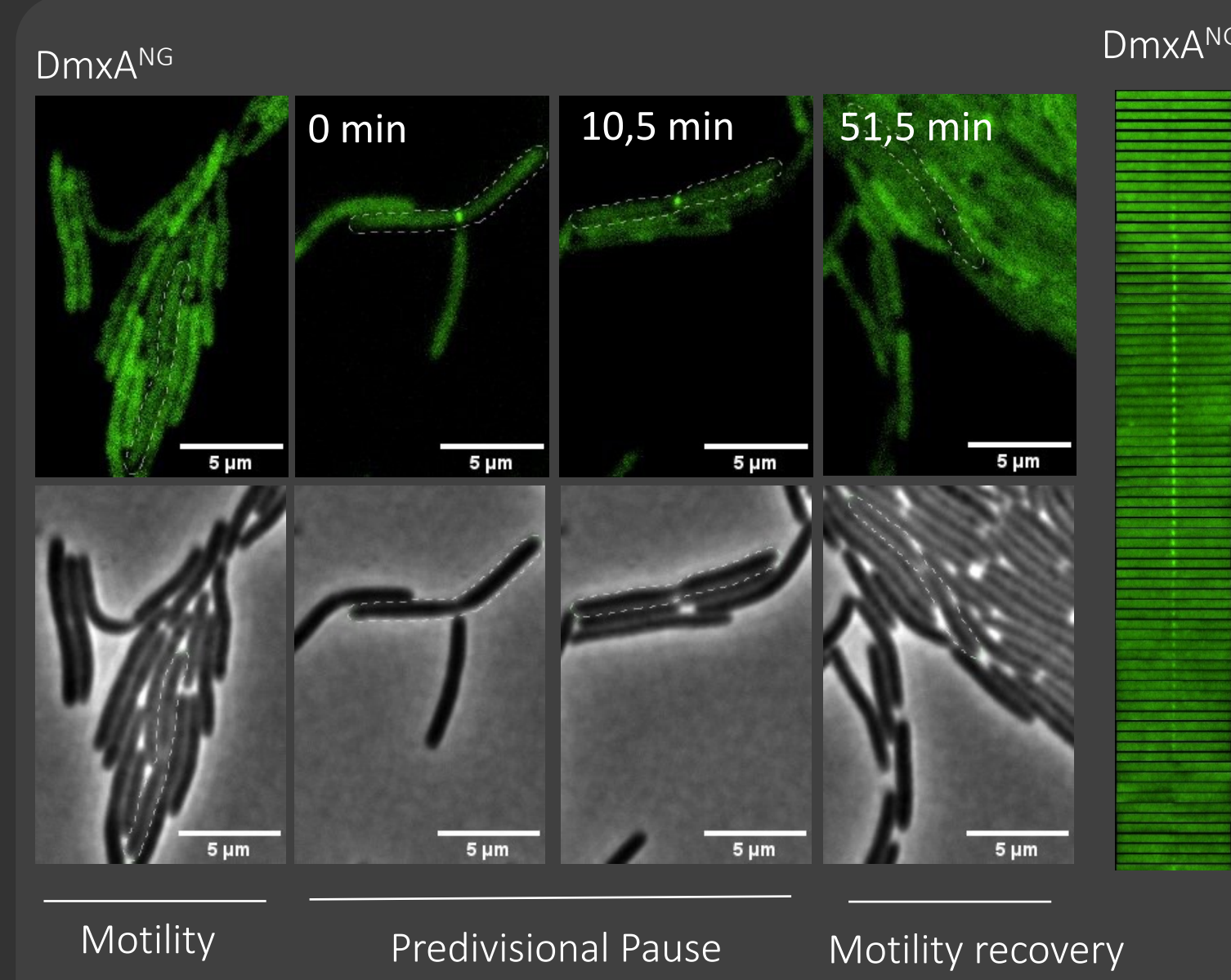
The $dmxA$ mutant is synthetically lethal in a $\Delta pomZ$ background

c-di-GMP biosensor



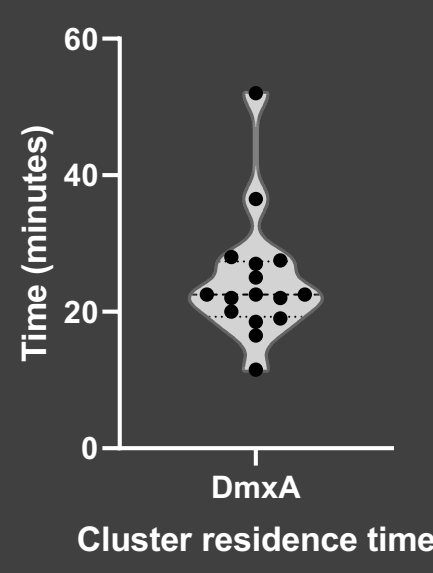
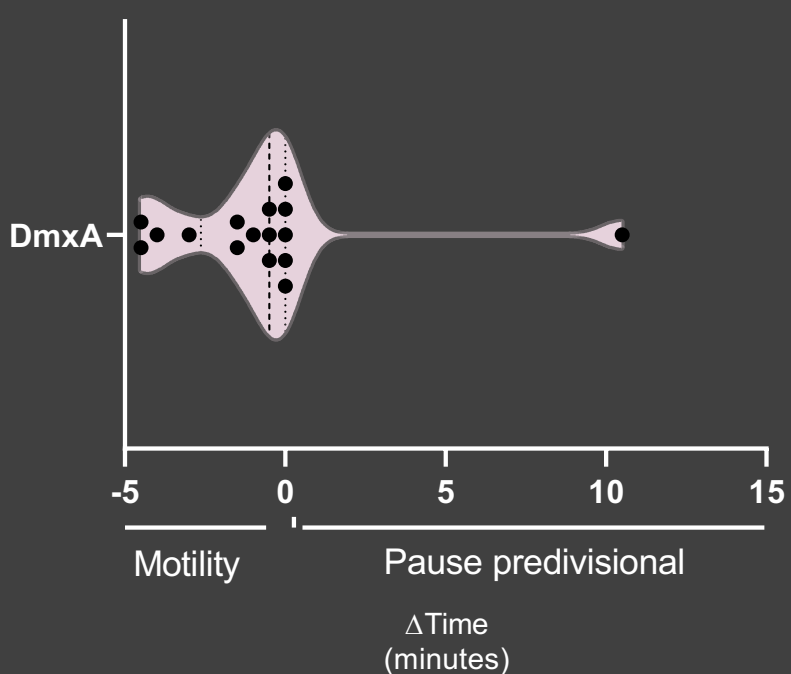
Kaczmarczyk et al. 2024

3 A DIGUANYLATE CYCLASE IS RESPONSIBLE FOR PREDIVISIONAL PAUSES



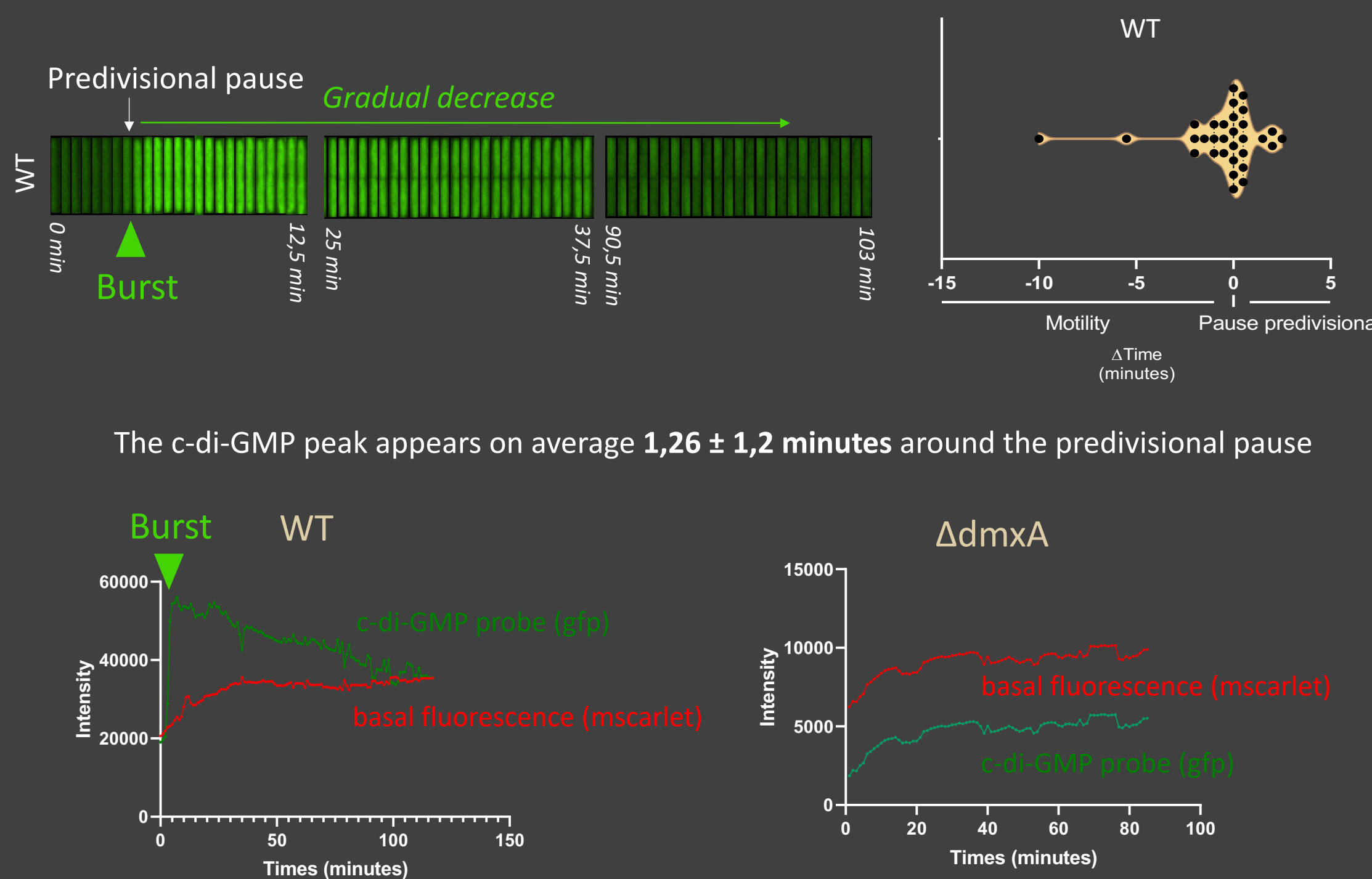
DmxA localizes at the septum on 2 ± 2.1 minutes around the predivisional pause

DmxA has an average cluster residence time of 24.57 ± 6.1 minutes



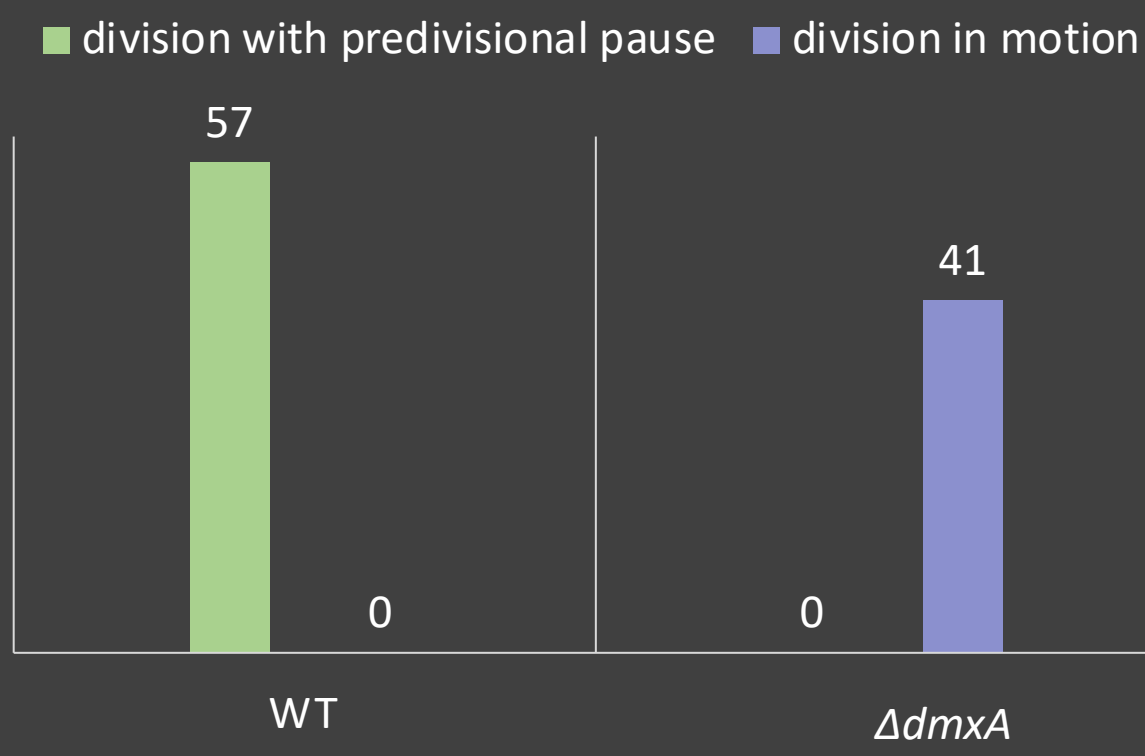
DmxA localizes to the division site prior to cytokinesis, suggesting a role in the early stages of cell division

A transient increase in c-di-GMP coincides with the division pause and is lost in $\Delta dmxA$ cells, indicating a role for DmxA in c-di-GMP signaling during division

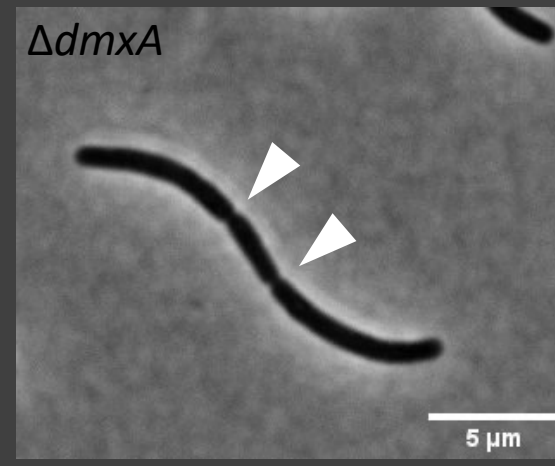


The c-di-GMP peak appears on average 1.26 ± 1.2 minutes around the predivisional pause

$dmxA$ deletion results in the absence of predivisional pause



Defects in cell division are also observed including asymmetric divisions and faulty septation



DmxA, a diguanylate cyclase, localizes to the septum before division and induces a transient increase in c-di-GMP levels, promoting a predivisional pause in *M. xanthus*. Without DmxA, this pause does not occur, leading to division defects.

Conclusions & perspectives

For the first time, we link predivisional pause of *Myxococcus xanthus* to a burst of c-di-GMP triggered by the diguanylate cyclase DmxA.

This transient signal precedes the disassembly of both motility systems, suggesting that c-di-GMP acts as a checkpoint linking motility shutdown to cell division. The relocation of key motility regulators to the septum further suggests a shared regulatory mechanism for both motility systems (A-motility and S-motility), although the downstream effectors remain to be identified.