

# Introduction and Theory

## 1 What is Phase Separation and Intrinsically Disordered Regions (IDRs)?

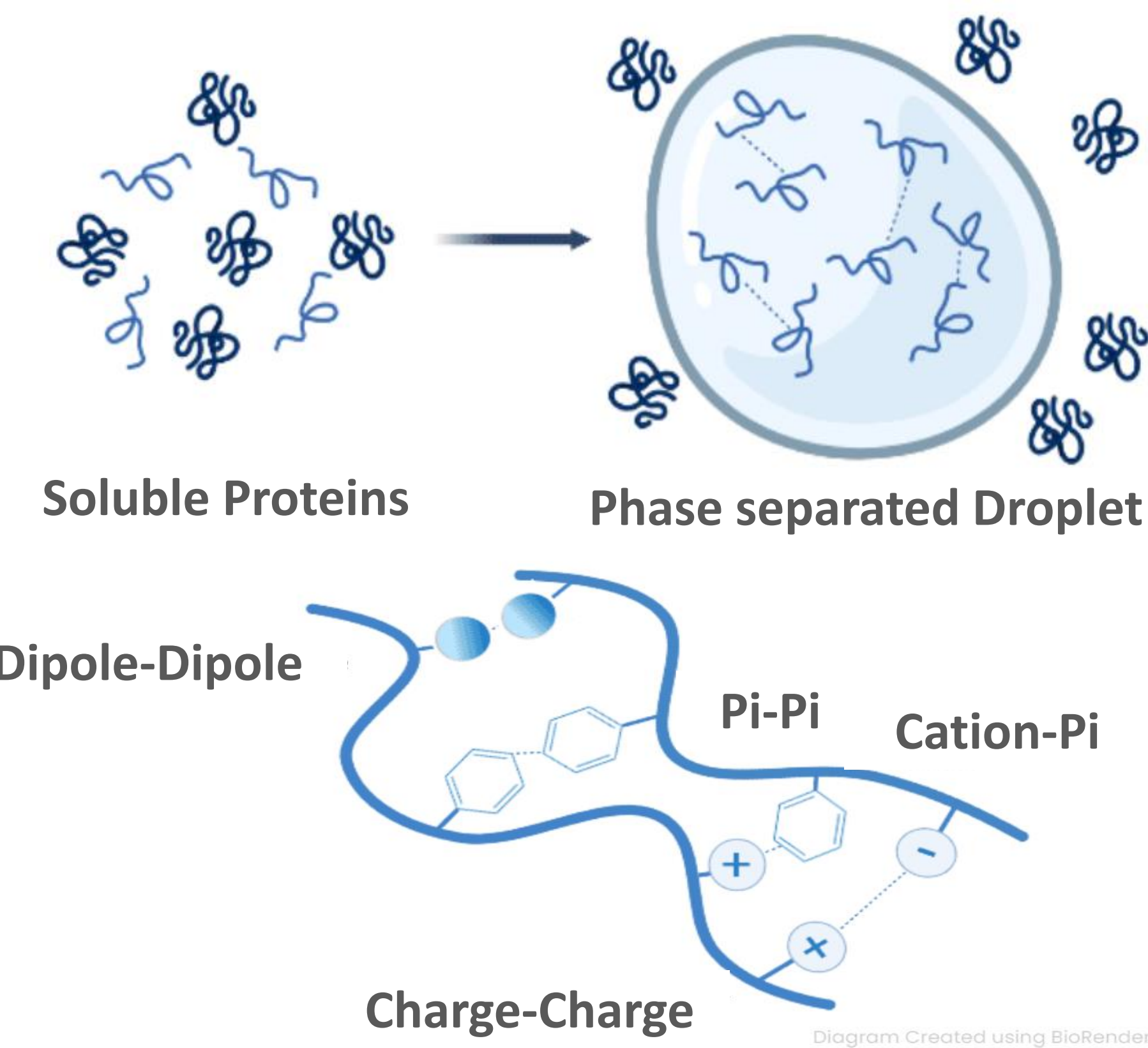


Figure 1. Figure 1a represents the phase separation of soluble proteins into their own “droplet”. Figure 1b shows the type of interactions that causes a protein to phase separate, and these are the type of interactions that occur between intrinsically disordered regions.

## 2 What are Min Waves?

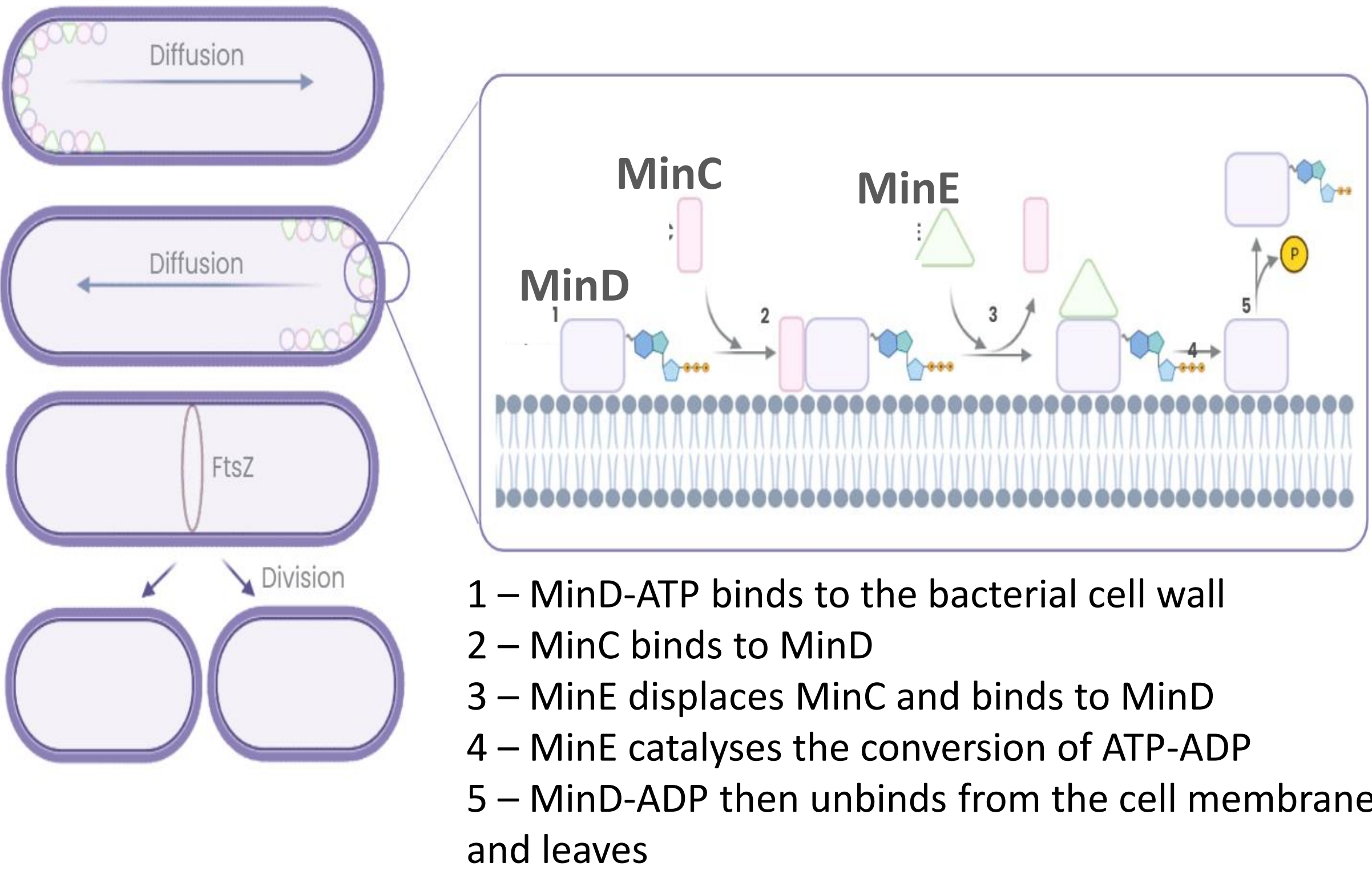


Figure 2. Diagrams representing the formation of Min waves.

## 3 Reproducing Min Waves + The Use of IDRs

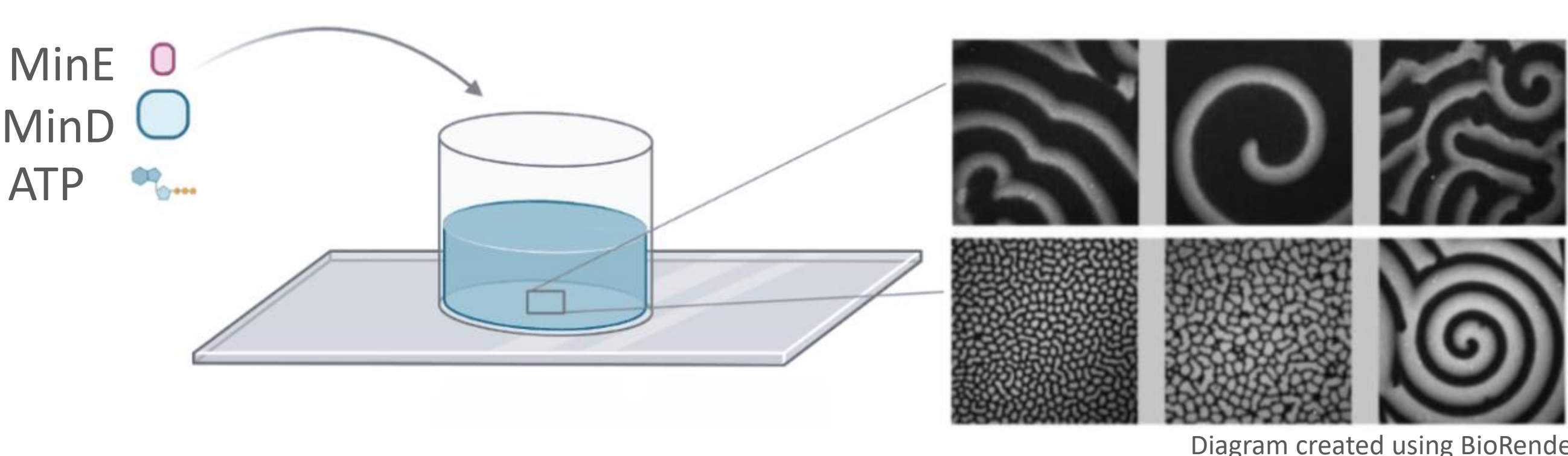


Figure 3. Diagram representing the reproduction of Min waves outside of bacterial cells. Min waves should give a consistent pattern. Therefore, this project aims to add IDR sequences to MinD and assess the pattern changes using mathematical models. These changes in the patterns should reflect the differing interaction strengths from the IDRs. Diagram adapted from Loose *et al.*

## Aims

- New method to look at IDR interactions
- Create a dynamic system that incorporates concentration gradients
- Create a system whereby the interactions can be clearly mapped

# The Use of Min Waves to Study Intrinsically Disordered Regions Involved in Phase Separation

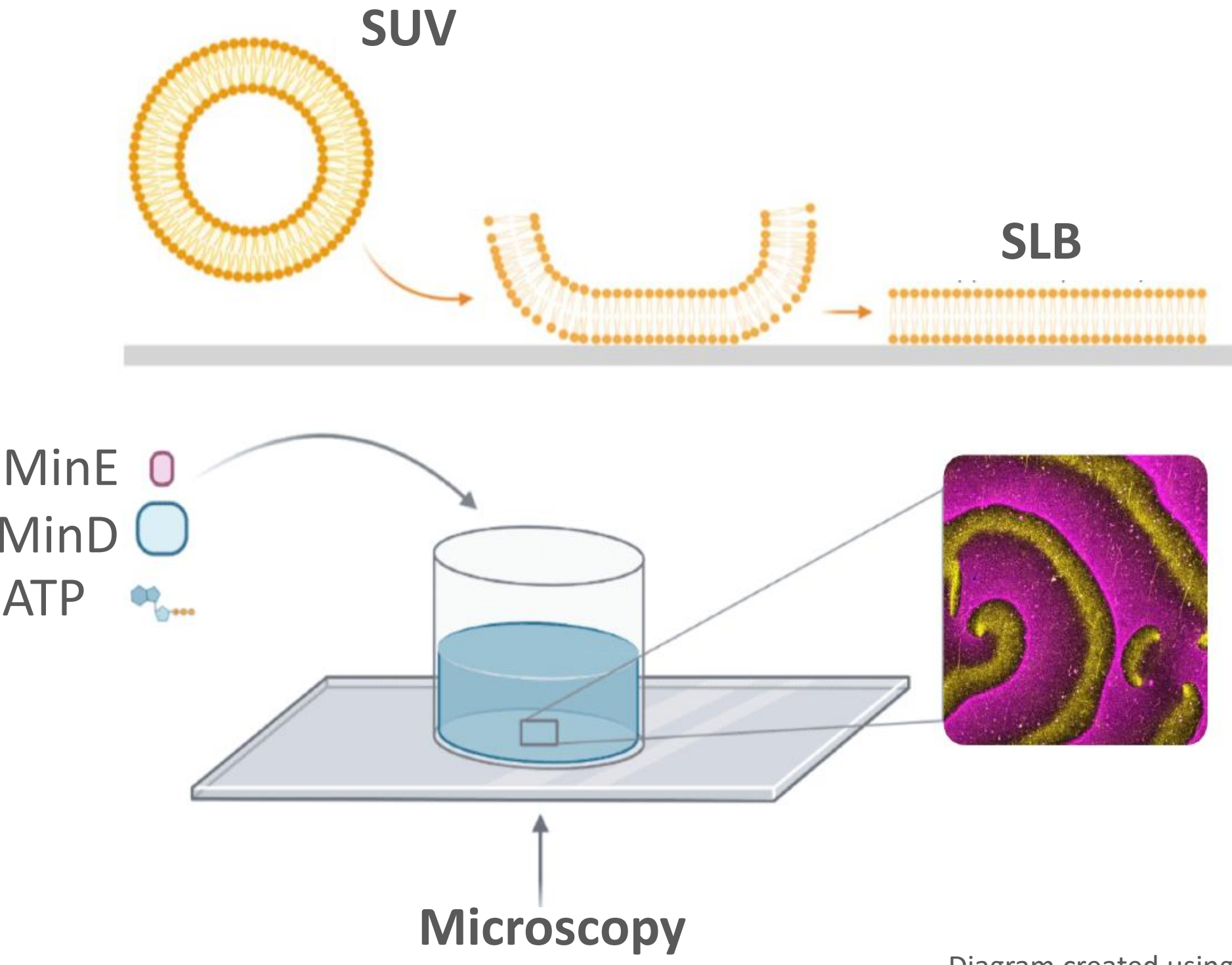
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## Methods and Results

### 1 Min Wave Formation - Theory

- Step 1**
- Create small unilamellar vesicles (SUVs) with DOPC and DOPG lipids
  - Using the sonication method.
  - SUVs will collapse onto a hydrophilic glass surface to create supported lipid bilayers (SLBs).
- Step 2**
- Purify MinE and MinD
  - Add them to a reaction chamber with lipid bilayers on the bottom
  - Add adenosine triphosphase (ATP)
  - Observe with confocal or total internal reflection (TIRF) microscopy



### 2 Supported Lipid Bilayer (SLB) Optimisation

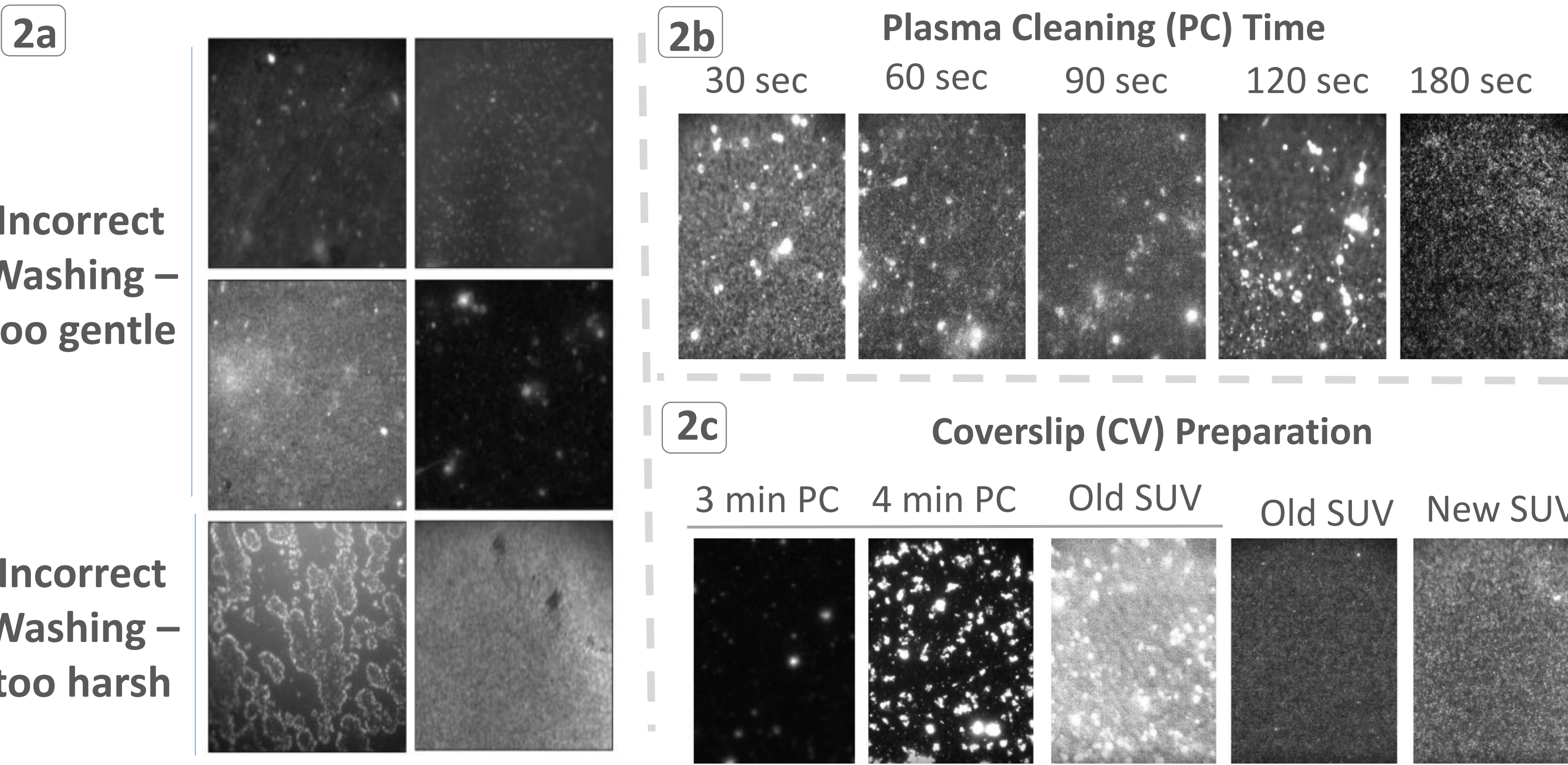


Figure 2a to 2c. Diagrams showing the optimisation of SLBs. Figure 2a depicts issues such as incorrect washing steps. Figure 2b shows the importance of plasma cleaning times to create the correct hydrophilic surface. Figure 2c shows the importance of using freshly prepared coverslips.

### 3 Protein Addition to Bilayers

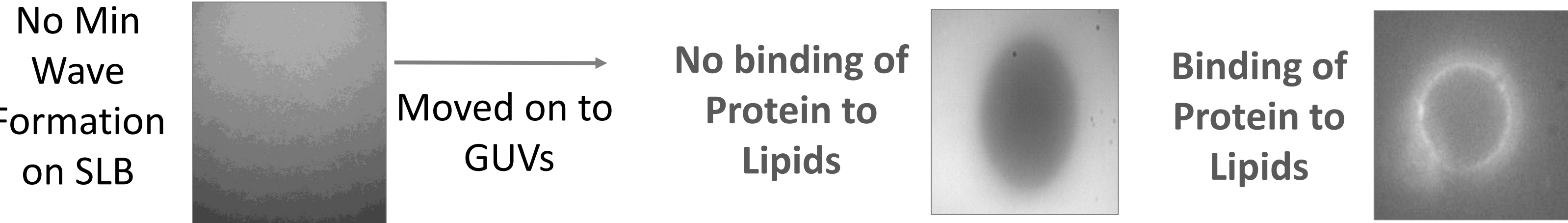


Figure 3. Addition of proteins to my lipid bilayers led to no change. Giant unilamellar vesicles (GUVs) were used as a diagnostic for lipid binding.

### 4 Protein Purification Optimisation

#### 4a Protein Purification Optimisation Steps

- Purify MinE, MinD GFP + MinD with Nickel
- Check IPTG concentration and Growth Times
- Check ADP Concentrations
- Check the Use of MinD-GFP
- Change to using Chitin bead/Intein Purifications
- Check different broths and different bacterial cells
- Swap from BL21 to Lemo(DE3) bacterial cells
- Active proteins, but impure
- Use SUMO tag – double purification step + soluble protein

#### 4b Bacterial Expression Plasmids Tried

Plasmid	Incubation Time	Incubation Temp.	Worked	Issue
MinE-His	3 hrs	37	Yes	No issues
MinD-His	18 hrs	20	No	Present in pellet, not active
MinD-His/MinE	18 hrs	20	No	Present in pellet
eGFP-MinD-His	18 hrs	20	Yes	Only use 20%
MinE-Intein	3 hrs	37	Yes	No issues
MinD-Intein	18 hrs	20	Yes	Active, but low yield
SUMO-MinE	3 hrs	37	?	?
SUMO-MinD	18 hrs	20	?	?

Figure 4a to 4b. Figure 4a represents some of the optimisation steps followed or being followed to get pure, high yield and active protein. Figure 4b represents the plasmids that have been tried. His represents the use of a nickel tag, intein represents the use of intein tag and chitin bead column and SUMO tag represents the use of a SUMO tag to keep proteins soluble.

### 5 Current Wave Patterns

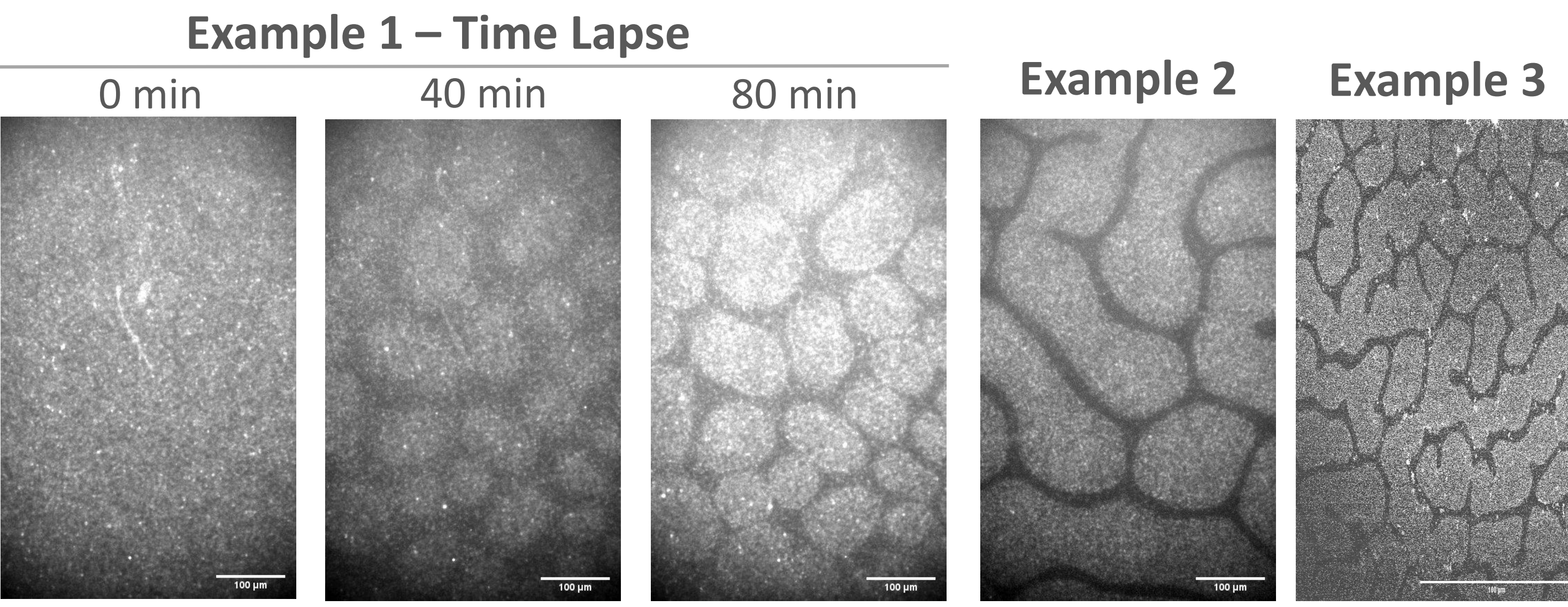


Figure 5. Figure representing the different wave patterns obtained so far. Example 1 shows a time lapse of how the waves form. Example 2 and 3 show the end of the wave formation for two other examples – at this point the waves remain static. Min waves are expected to regenerate in the presence of an excess amount of ATP (which was added in excess). However, these patterns remained static.

## Future Prospects and Ongoing Projects

- Purify MinD/MinE SUMO
- Purify MinD with an IDR attached
- Code for the different interactions to assess the IDR parameters of importance

## Conclusion

- Min waves are specific, concentration dependent reactions
- Optimisation of purifications needs to be performed with a SUMO tag
- Code for the different interactions will assess the IDR parameters of importance

### References

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