

Self-assembly pathways of intrinsically disordered glycomacropeptide

Andrea Lombardo^{1,2}, Tobias Winckler-Carlsen², Vito Foderà² and Valeria Vetri³

1. Department of Biological, Chemical and Pharmaceutical Sciences and Technologies, University of Palermo, 90133 Palermo, Italy

2. Department of Pharmacy, University of Copenhagen, Universitetsparken 2, 2100 Copenhagen, Denmark

3. Department of Physics and Chemistry, University of Palermo, 90128 Palermo, Italy

INTRODUCTION

Glycomacropeptide (GMP)

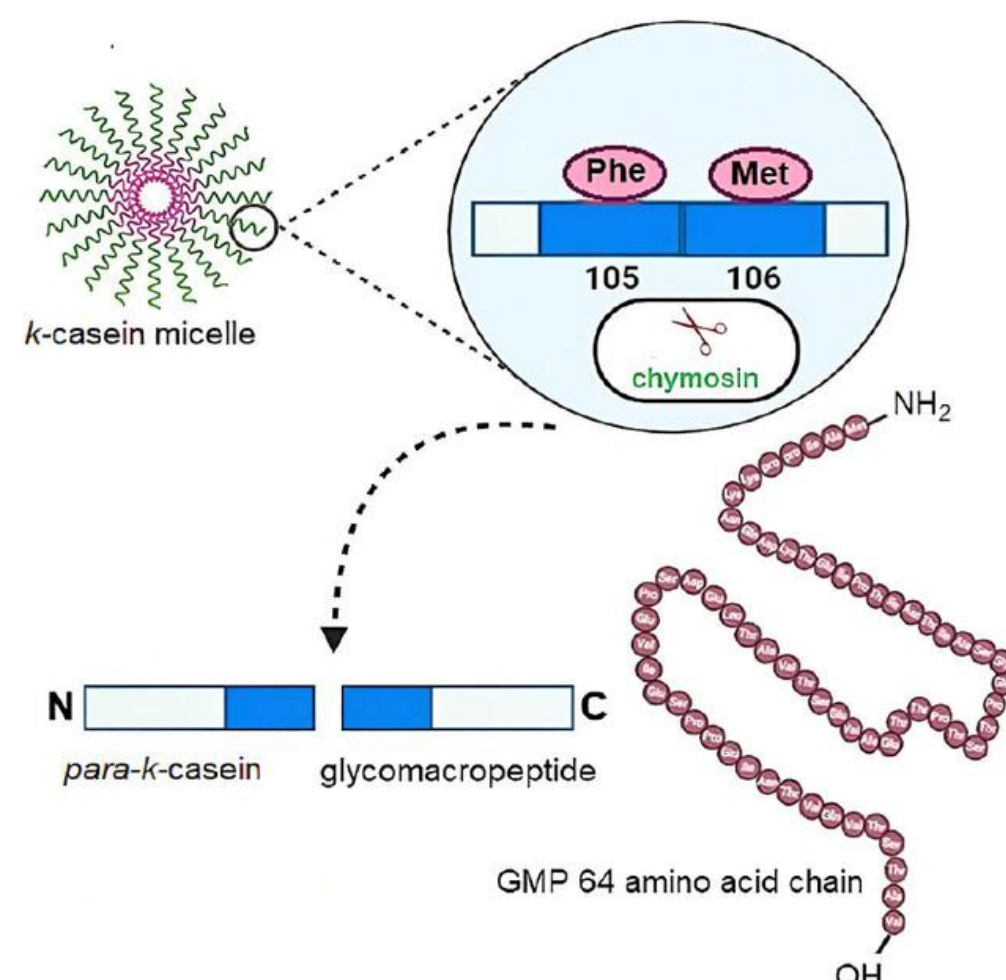
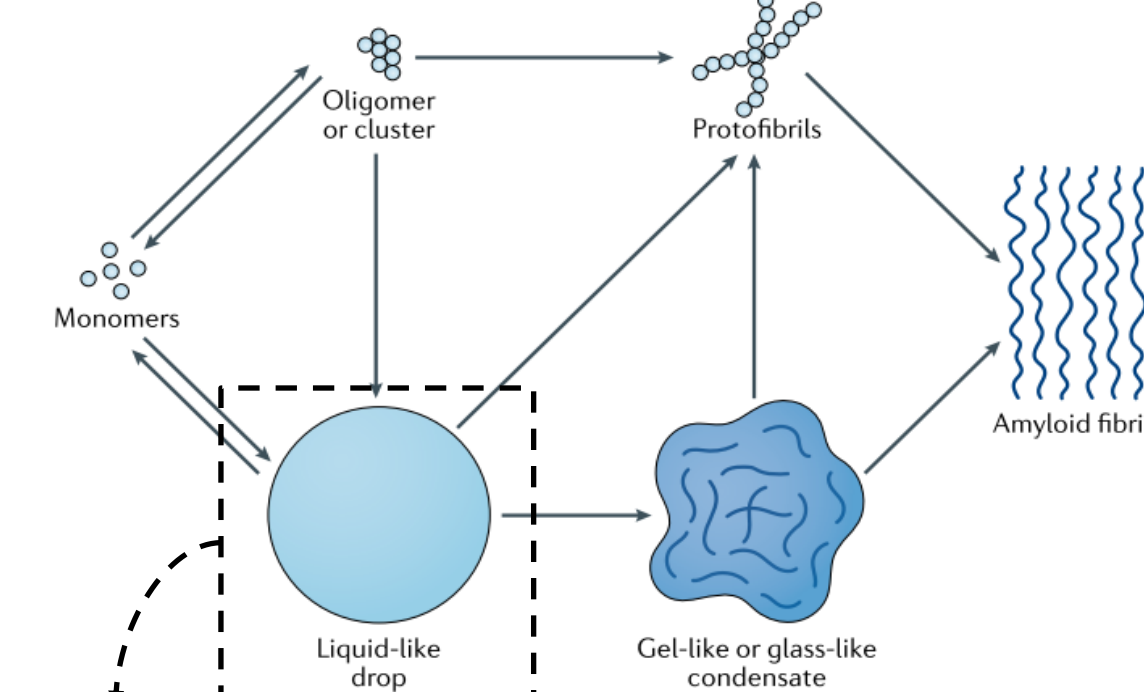


Figure 1: Chymosin action in the production of bovine glycomacropeptide (GMP). Adapted from Ebrahimi, 2024, Compr Rev Food Sci Food Saf.

Glycomacropeptide (GMP) is an **intrinsically disordered** casein-derived peptide composed of 64 amino acid residues and characterized by a low isoelectric point ($pI < 3.8$). It is rich in polar amino acids and presents up to 4 carbohydrate chains.

Self-assembly pathways

Figure 2: Adapted from Alberti, 2021, Nature Reviews.



Why self-assembly?

- Drug delivery
- Studying biological colloid stability
- Model system

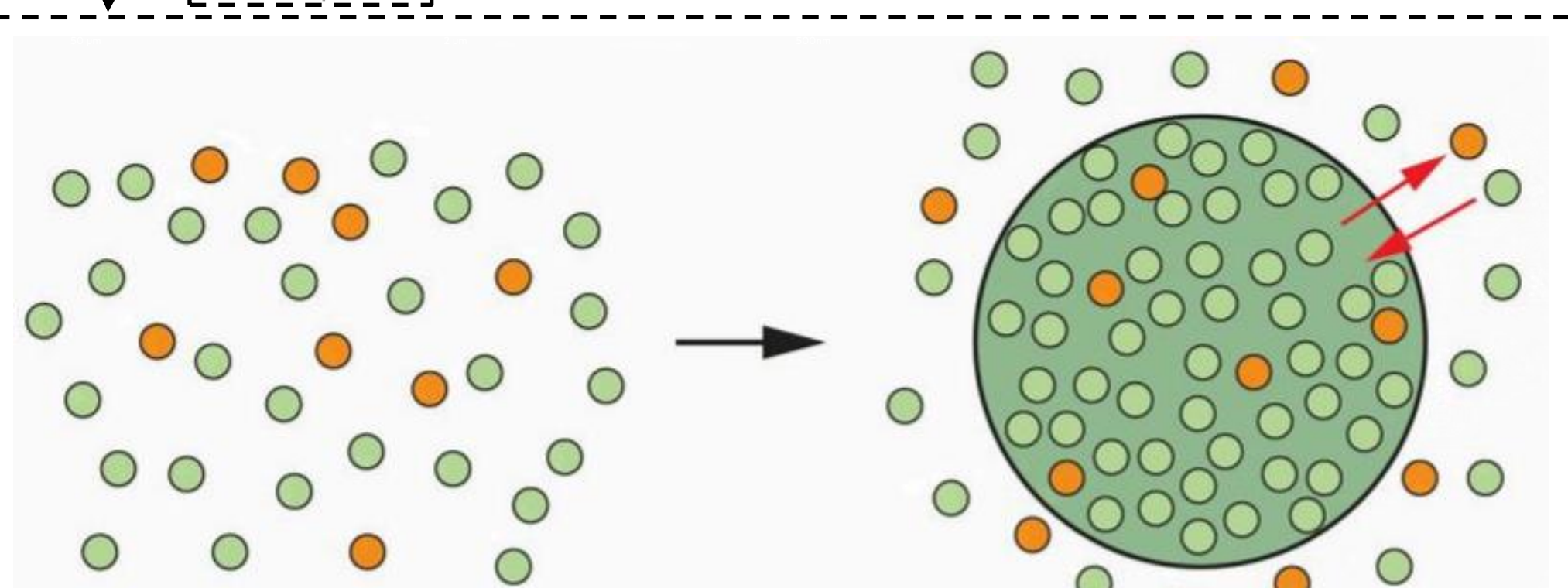
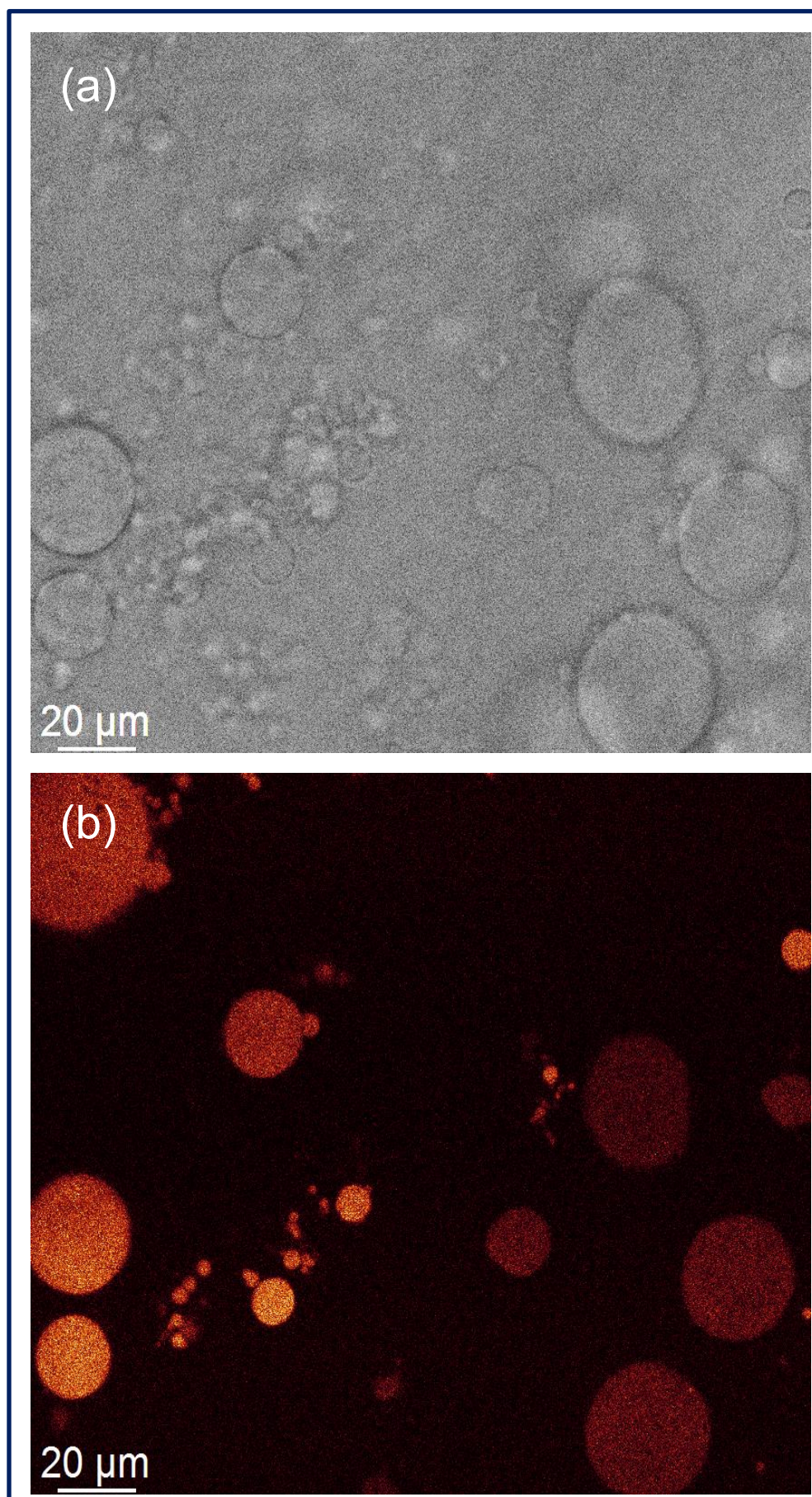


Figure 3: **Liquid-liquid phase separation (LLPS)** mechanism. Adapted from Shin, 2017, Science

RESULTS

Lower GMP Concentration (37.5 mg/mL)



Setting conditions for phase separation

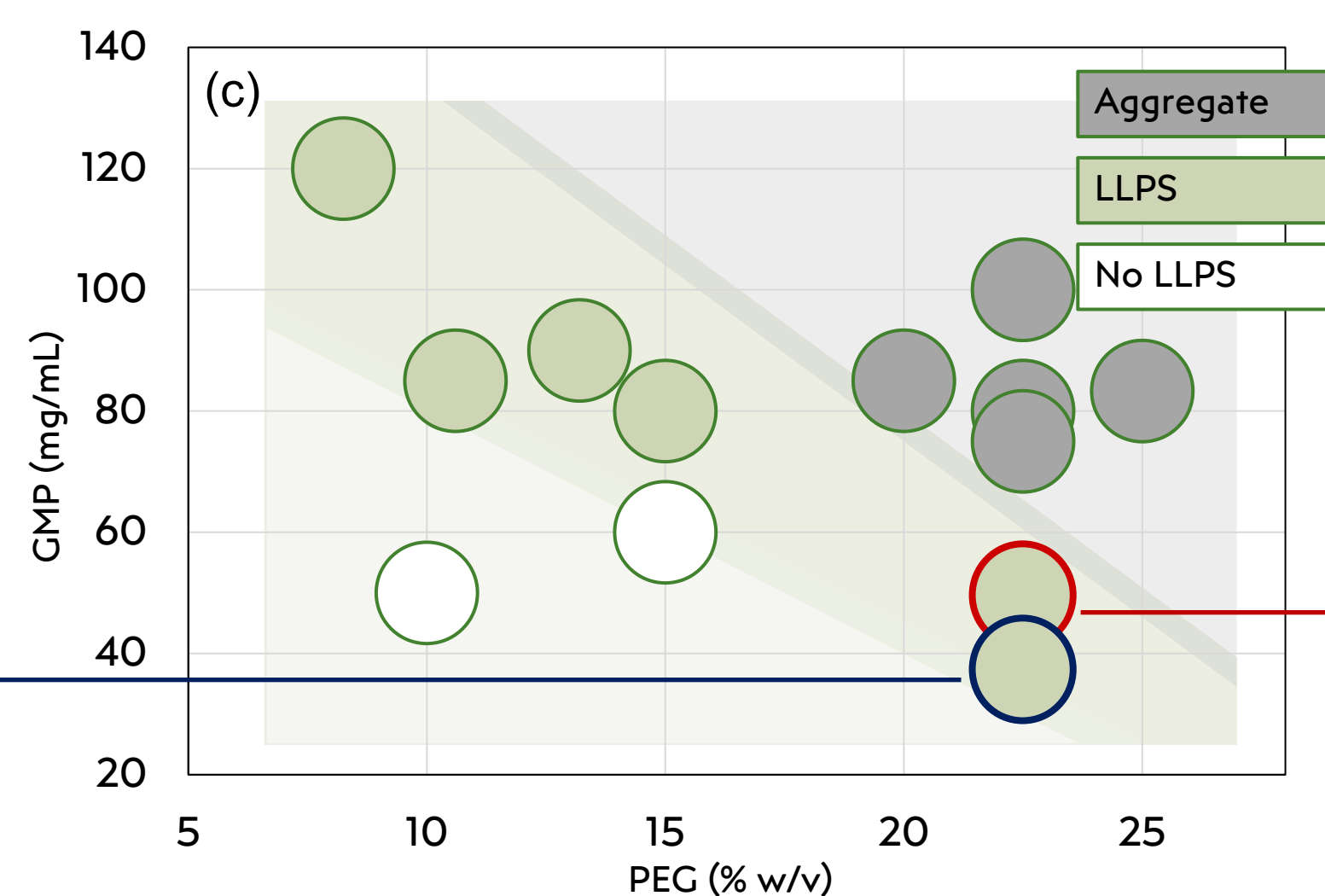
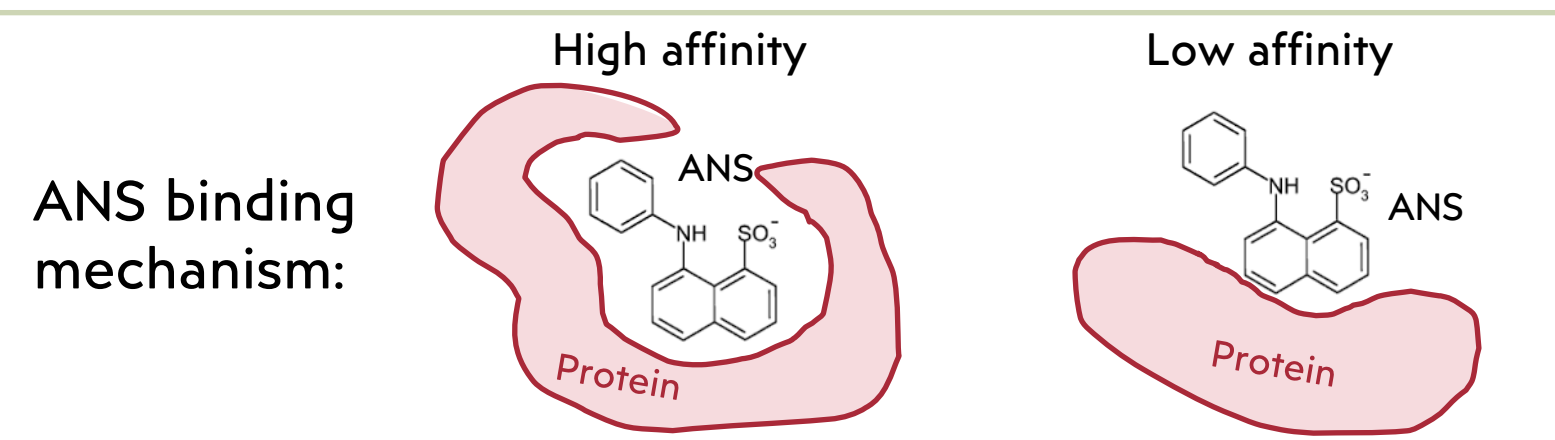
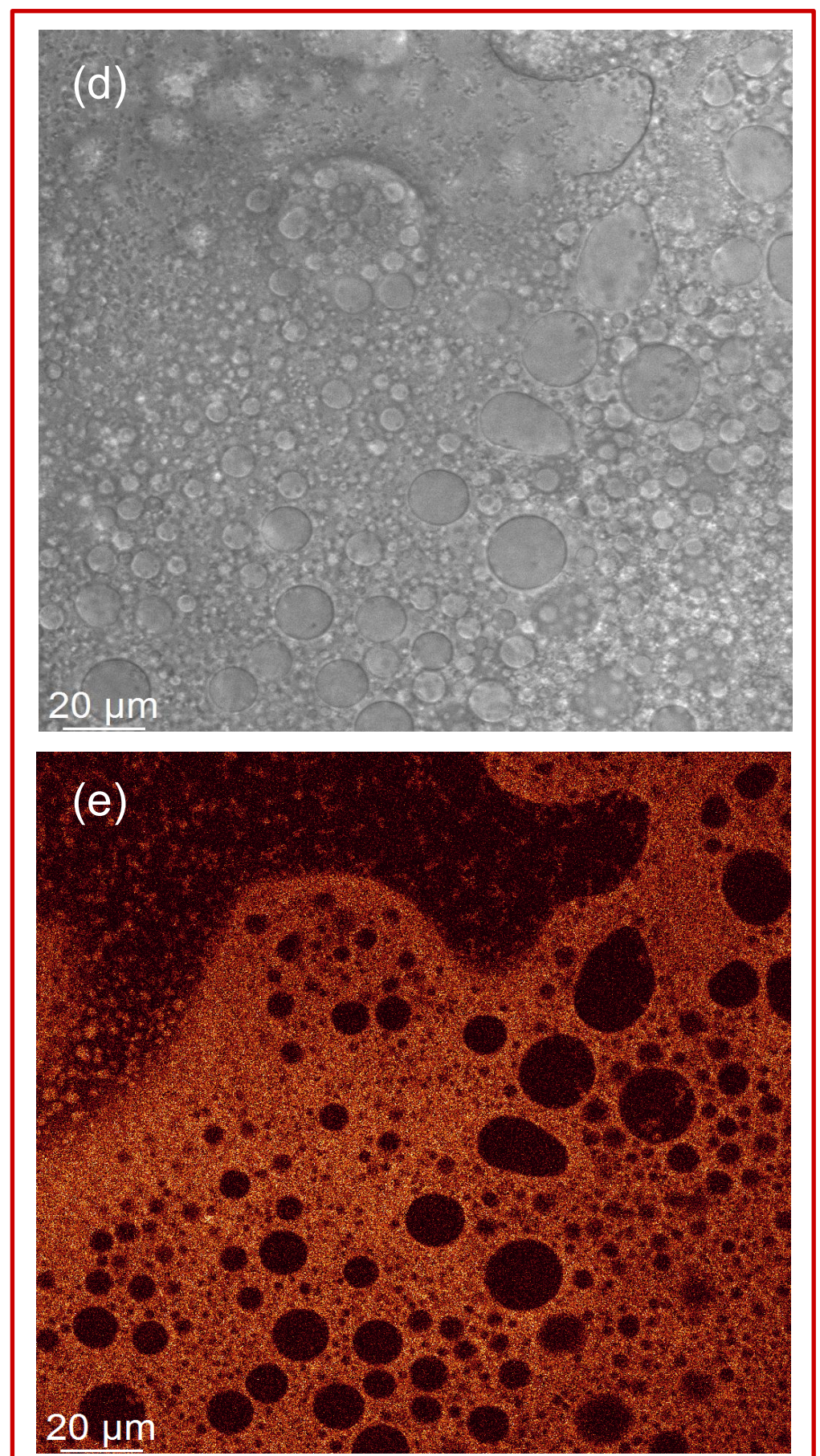


Figure 4: Transmittance (a-d) and ANS fluorescence (b-e) of GMP in the presence of PEG. (c) Phase diagram illustrating the experimental conditions for aggregation and liquid-liquid phase separation (LLPS). The blue and red markers highlight the specific condition selected for the adjacent images.



Higher GMP Concentration (50 mg/mL)

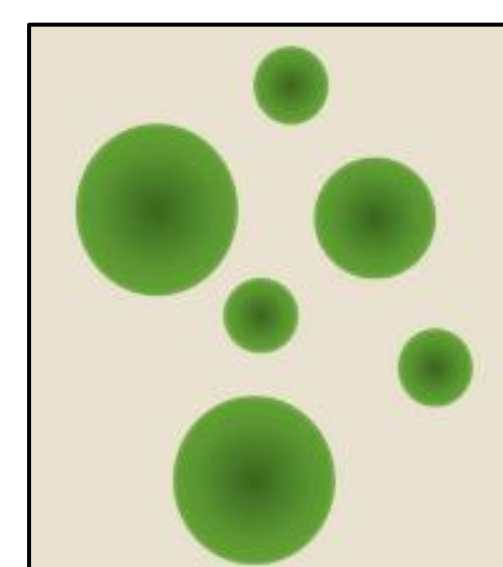


CONCLUSION AND FUTURE DIRECTIONS

Conclusions

- Conditions leading to GMP **phase separation** were identified using PEG as a crowding agent.
- Different GMP concentrations lead to droplets enriched in **different components**.
- ANS fluorescence reveals heterogeneous protein environments, supporting the coexistence of **different protein states** under phase-separation conditions.

Next steps



FRAP
SpIDA
ThT-phasor FLIM

- Revealing internal dynamics and diffusion rates.
- Mapping oligomerization within coacervates.
- Protein conformational changes.