



CiQUS

Centro Singular de Investigación
en Química Biolóxica e
Materiais Moleculares

Conferencia:

Systems biochemistry of bacterial division: Reconstructing minimal divisomes in the test tube

Q

Germán Rivas

CIB-CSIC, Madrid

03/07/15

.....
Salón de Graos
Fac. Bioloxía

12:15 h

Más información:
www.usc.es/ciqus



XUNTA DE GALICIA

CONSELLERÍA DE CULTURA, EDUCACIÓN
E ORDENACIÓN UNIVERSITARIA

USC
UNIVERSIDADE
DE SANTIAGO
DE COMPOSTELA

CAMPUS VIDA
CAMPUS DE EXCELENCIA INTERNACIONAL

Systems biochemistry of bacterial division: Reconstructing minimal divisomes in the test tube

Germán Rivas. CIB-CSIC, Madrid <grivas@cib.csic.es>

The research in our laboratory aims at understanding how the elements of the bacterial division machinery - the divisome - work together as an integrated system to fulfil its essential function. To address these questions we develop and applied novel biochemical reconstitution approaches to assemble the minimal set of proteins needed to initiate division (the proto-ring complex) in systems that reproduce the spatio-temporal organization of the divisome at the cellular membrane and the crowded/confined intracellular space.

Using physical biochemistry and synthetic approaches, we study the activities, interactions and assembly properties of minimal reconstructions of the proto-ring structured in membrane-like systems, such as nanodiscs, micro-beads, bilayers, vesicles and micro-droplets. We also investigate the action of Min proteins and nucleoid-like structures (to reproduce Z-ring positioning mechanisms) on the properties of minimal divisomes.

The assembly of the divisome in the cell takes place in microenvironments characterized by the presence of high concentrations of unrelated macromolecules, often structured as soluble and/or membrane-bound dynamic networks. We apply and design synthetic reconstructions of these microenvironments to investigate the impact of the physicochemical properties of facsimile cell media on the reactivity and organization (in time and space) of minimal divisome assemblies.

These studies will contribute to define the precise conditions to build, with a minimum set of elements, functional division assemblies in the absence of cells. This integrated approach will help to complete our knowledge of how bacterial division works and will open new horizons to synthetic and biotechnological applications.

SELECTED REFERENCES

- Cabré EJ, Sánchez-Gorostiaga A, Carrara P, Roperio N, Casanova M, Palacios P, Stano P, Jiménez M, Rivas G*, Vicente M*. 2013. Bacterial division proteins FtsZ and ZipA induce vesicle shrinkage and cell membrane invagination. *J. Biol. Chem.* 288:26625-26634.
- Hernández-Rocamora VM, Reija B, García-Montañés C, Natale P, Alfonso C, Minton AP, Zorrilla S, Rivas G*, Vicente M*. 2012. Dynamic interaction of the *Escherichia coli* cell division ZipA and FtsZ proteins evidenced in nanodiscs. *J. Biol. Chem.* 287:30097-30104
- Hernández-Rocamora VM, García-Montañés C, Rivas G*. 2015. Phospholipid bilayer nanodiscs: A powerful tool to study the structural organization and biochemical activity of proteins in membrane-like environments. *Curr. Top. Med. Chem.* 14:2637-2646
- Martos A, Jiménez M, Rivas G*, Schwille P*. 2012. Towards a bottom-up reconstitution of bacterial cell division. *Trends Cell Biol.* 22:634-643.
- Martos A, Raso A, Jiménez M, Petrášek Z, Rivas G*, Schwille P*. 2015. FtsZ polymers tethered to the membrane by ZipA are susceptible to spatial regulation by Min waves. *Biophys J.* 108:2371-2383.
- Monterroso B, Alfonso C, Zorrilla S, Rivas G* (2013) Combined light scattering, ultracentrifugation and fluorescence correlation spectroscopy studies on the associations and assembly of the *Escherichia coli* cell division FtsZ protein. *Methods* 59:349-362.
- Rivas G*, Alfonso C, Jiménez M, Monterroso B, Zorrilla S. 2013. Macromolecular interactions of bacterial cell division FtsZ protein: From quantitative biochemistry and crowding to reconstructing minimal divisomes in the test tube. *Biophys. Rev.* 5:63-77
- Rivas G*, Vogel SK, Schwille P. 2014. Reconstitution of cytoskeletal protein assemblies for large-scale membrane transformation. *Curr. Opin. Chem. Biol.* 22:18-26