

University of Stuttgart

Stuttgart Research Center Systems Biology (SRCSB)

Systems Biology Seminar Talk



"Regulatory networks involving small non coding RNAs connect stress responses, metabolism adaptation and virulence in *Staphylococcus aureus* "

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Abstract:

It is now well recognized that RNAs are key intracellular effectors in fast adaptive processes. In many cases, they co-regulate the expression of functionally related genes at the post-transcriptional level. Today, more than 200 potentially regulatory RNAs have been identified so far in S. aureus. In order to define their direct targets and the regulatory networks in which they belong, we have recently applied the MS2-affinity purification approach coupled with RNA sequencing to several non coding RNAs in S. aureus (1,2). Using a combination of in vivo and in vitro approaches, numerous of the co-purified and enriched mRNAs have been validated as direct targets. Several non coding RNAs will be used to illustrate the intricate interactions between RNAs, two-component systems and transcriptional regulatory proteins and their implication in stress responses (including oxidative and NO), sugar metabolism and virulence..





<u>CV:</u>

- Education:

1986 Habilitation, Doctorat d'Etat ès sciences, University of Strasbourg

1983 PhD Molecular Biology, thèse 3ème cycle, IBMC, University of Strasbourg

- Scientific Career:

2016 Director of the Unit UPR 9002 CNRS 2004 Team leader « Bacterial regulatory RNAs and mRNAs », IBMC Strasbourg

- Awards:

Lecture Hall 0.106 Allmandring 31 Stuttgart 2018 Pasteur Medal, French Academy of Sciences
2010 Langevin Prize, French Academy of Sciences
Administrative Responsabilities:

2016 - 2020 Nominated member of the section CSS1 of INSERM
2014 - 2015 Nominated member of ATIP/AVENIR committee
2008 - 2012 Nominated member of the section 21 CNRS

