

Training and research in *Listeria*monocytogenes adaptation through proteomic
and transcriptome deep sequencing analysis

#### Workshop 2

#### 09-11 October 2017

# Dyrlaegevej 88, 1870 Frederiksberg, Copenhagen, Denmark

## **Programme**

## Monday, 9th

09:00-1:00 pm

**08:00 am**Bus departure 8:12 Norrepørt station (15E)

- Visit of company ChrHansen, R&D departmentthe importance and use of science
  - discovery of microbial strains for new business
  - microbial concepts for bioprotection and less food waste
  - use of analytical methods and big data on microbes (trials in an e-lab, global phage monitoring program)

#### Lunch

Questions and answers: life of a PhD in microbiology in ChrHansen

1:15 pm Bus/train back to UCPH

2:45 pm Cold stress response mechanisms in *Listeria monocytogenes* 

3:30 pm WGS in Listeria outbreak investigation

4:15 pm *coffee break* 

4:15 pm Group discussion of the topics of the day

5:00 pm End of the day\*

### Tuesday, 10th

9:00 am	System biology part I: can the bacterium be considered as a system?
10:00 am	System biology part II: modelling transcription and translation in bacteria
11:00 am	coffee break
11:30 am	Model vs Data: parameter estimation: Bacillus subtilis as an example
12:30 pm	Lunch pizza



<sup>\*</sup> Optional: joint dinner (https://www.paludan-cafe.dk)



1:30 pm Model vs Data: advanced analysis of absolute protein quantification

2:00 pm Modelling the bacterial cell as a system: the resource allocation as a simple principle

3:30 pm coffee break

4:00 pm Practice: resource allocation predication as a predictive tool (Matlab)

**5:00 pm** End of the day

# Wednesday, 11th

**9:00 am** Listeria monocytogenes growth and survival, discussion of pitfalls

09:45 am coffee break

10:00 am Interactive session-modelling of the ESRs data

12:00 pm Lunch (sandwiches)

**1:30** End of the workshop\*





<sup>\*</sup> Optional: meet outside the building to follow the locals to the auditorium where the annual meeting takes place