- You have identified a new probiotics strain that harbors a gene encoding for a key enzyme that could provide beneficial effects on digestion. You plan to study if the probiotic could indeed be beneficial in order to propose it as a treatment for this digestive disorder.
- How can you evaluate if the bacterial strain could be an efficient probiotic? How can you evaluate if this strain is safe? What analytical strategy and study design can you propose to address these questions?

- You would like to characterize as completely as possible the genome, transcriptome and proteome of a new model organism (mammalian) which has tremendous advantages for the study of metabolic disorders.
- How would you proceed steps by steps?
- Which approach could you envisage to create a population of this organism with genetic diversity to identify genes involved in metabolic disorders?

- You are collaborating with your colleagues from Case Study 2! You are highly interested to benefit from their new model organism to evaluate how it could help you in assessing a library of 100 natural compounds for muscle fitness.
- What type of experiment could you propose to identify active compounds, and to identify the mechanism of action of the compound?

- Proteomics is more challenging than genomics, for which reasons?
- In particular what are the challenges of proteomics in analyzing complex protein mixtures?
- Could you give an example of how you would discover/develop a proteomics-based biomarker?

- How would you evaluate the potential of a library of short peptides on cell ageing?
- Once you identified candidates, how would you test them on animal models?