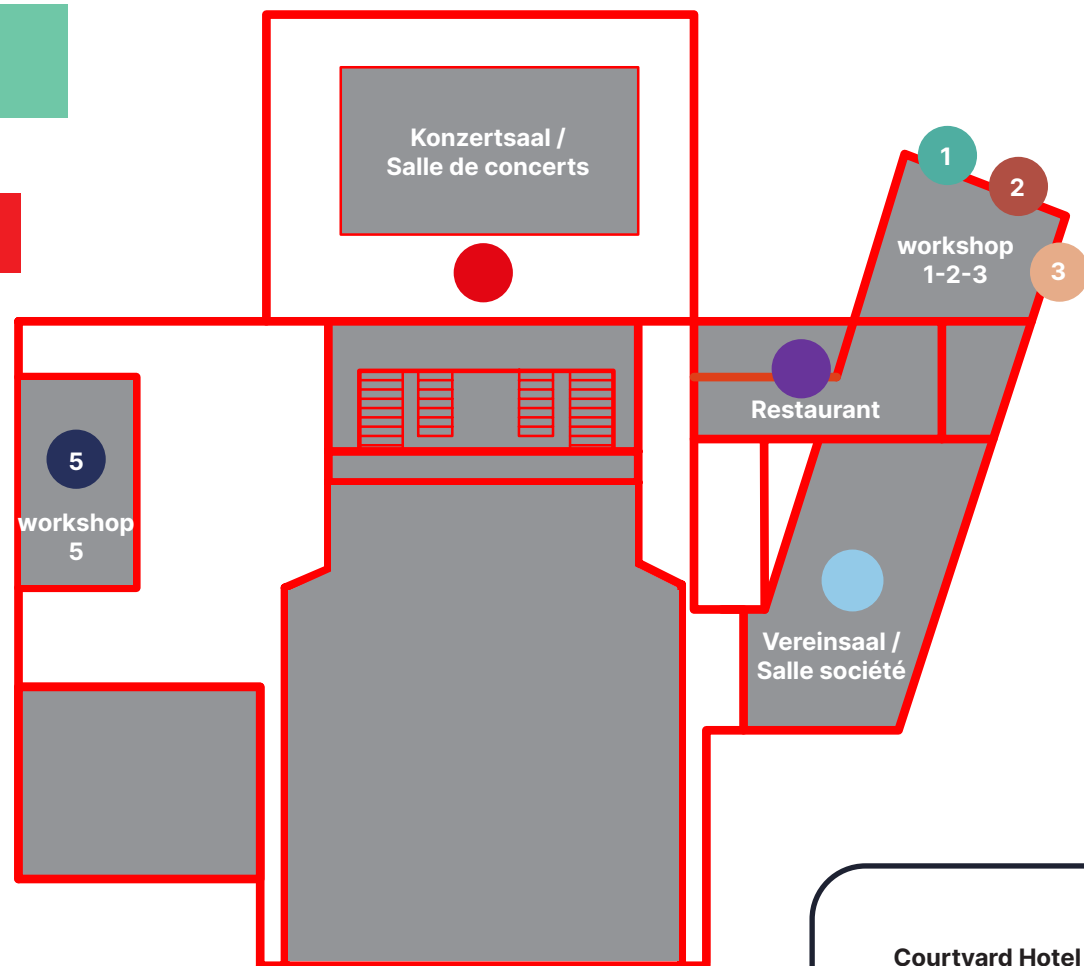













Tutorials & workshops



Congress Center

-  Analysis of spatial transcriptomics data (**Restaurant**)
-  Best practices to support the reproducible research value chain in bioinformatics: from raw data to final publication (**Konzertsaal**)
-  Machine learning in R (**Vereinsaal**)
-  PASTA: Pattern Analysis for Spatial Omics Data (**Workshop 2**)
-  Single cell RNA sequencing data analysis: Requirements for reproducibility and meaningful multi-omics integration (**Workshop 5**)
-  Towards sustainable science: navigating environmental challenges in bioinformatics (**Workshop 1**)
-  Understanding protein and glycoprotein structure – activity relationships with applications in personalized medicine and computer-aided drug design (**Workshop 3**)

Courtyard Hotel

-  How to avoid pitfalls and misinterpretations during the profiling of metagenomes (**Cinnamon**)
-  Incorporating biological sex as a variable in the design and analysis of biomedical research experiments (**Juniper**)
-  Quality Control for Protein 3D Structures and Models (**Ginger 1**)
-  Reproducibility Hackathon (**Ginger 2**)

