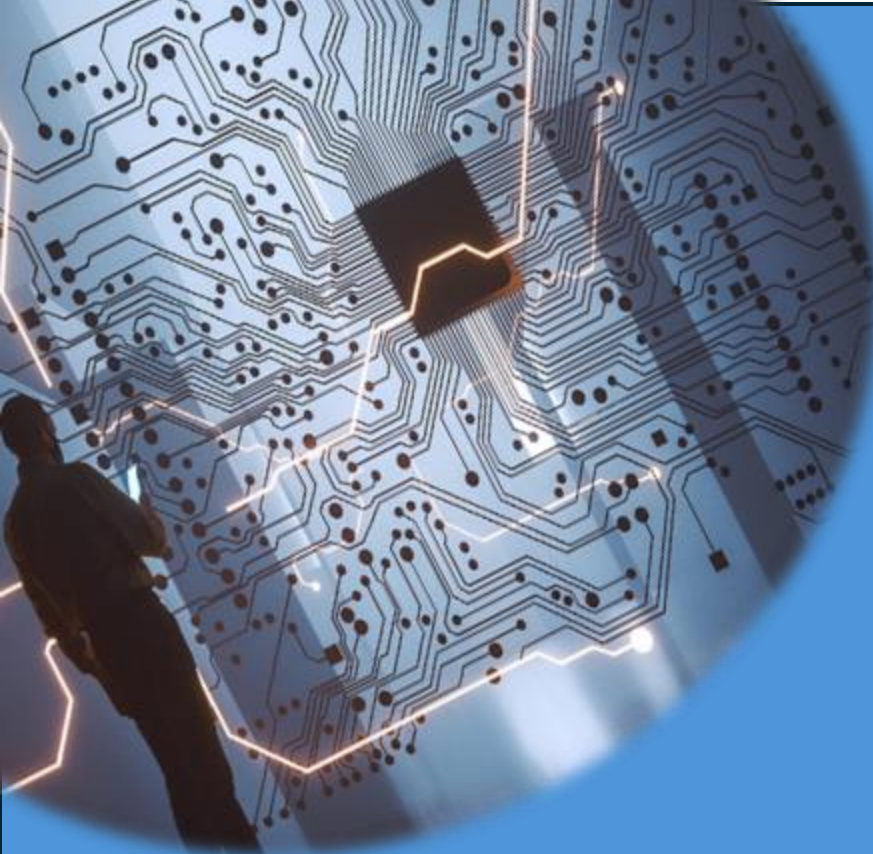




Public data, metadata

FAIR

Re-use, re-integration,
re-investigation

CONFÉRENCE

Public nucleotide sequence
databases: obstacles and
opportunities for data mining

12 octobre
13h

LMGE / UCA

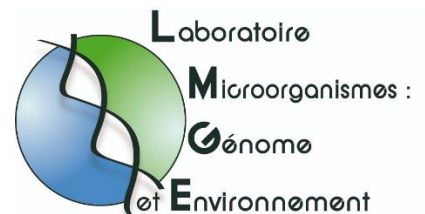
Amphi 1 Cézeaux

Only a fraction of the information contained within a data set is explored by the study in which the data were generated. Public sequencing data therefore holds a tremendous potential for re-use and integration to investigate novel scientific aspects. However, such approaches require the data to be FAIR, i.e. described with rich, standard-compliant and machine-readable metadata

PRÉSENTÉE PAR:

**Christiane
Hassenrück**

**Leibniz Institute for Baltic Sea
Research**



**ÉCOLOGIE &
ENVIRONNEMENT**