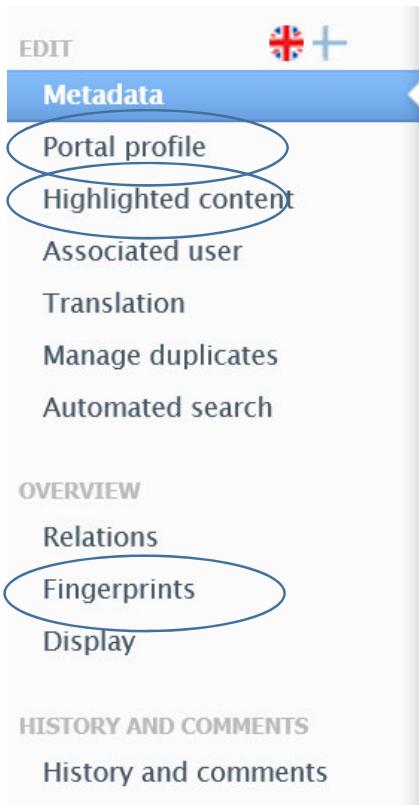


# Functionality of the new portal

Please note that you make all the changes in acris.aalto.fi



## Portal profile

1. You can choose to show or hide your collaboration map.



2. Choose which affiliations you want to show in your personal profile.

Select the organisational unit affiliations to be shown on your profile on the Pure Portal. By default, all your active organisational unit affiliations are shown.

If you hide an affiliation it may still be present on the Pure Portal on other pages, but it is not shown on your personal profile.

### 3. Metrics

We have disabled H-index and aggregated citations in the personal profile. Scopus citations are shown with publications.

## Highlighted content

You have the possibility to highlight your most important publications, activities, prizes, press/media and projects.

If you choose not to highlight content, the newest content is used.



## Research outputs



## Fingerprints

The Fingerprint Engine mines the text of documents, i.e. publication abstracts, project summaries, patents – to create an index of weighted terms which defines the text, known as a Fingerprint visualization. These fingerprints are used to map content. The Elsevier Fingerprint Engine uses a variety of thesauri to support applications pertaining to different subject areas. The total number of concepts is over 700,000 distinct concepts

You can choose which fingerprint concepts to show in your personal profile.

Fingerprints

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Engineering & Materials Science (CpxTree)

Sort by: Rank A-Z

INCLUDED CONCEPTS

Knowledge acquisition

Factorization

Feedback

Visualization

Reinforcement learning

Learning systems

Brain

Gene expression

Experiments

Medicine

Inference engines

Decision making

Search engines

Linear regression

Information retrieval

Display devices

Genes

Simulators

Factor analysis

Eye movements

Biomarkers

Set theory

Drug therapy

Brain computer interface

Sensors

Image classification

Statistics

Bayesian networks

Intelligent systems

Medical problems

Tensors

RNA

Demonstrations

User interfaces

Cells

Electroencephalography

Dynamical systems

Navigation

Differential equations

Oncology

Cytotoxicity

Bioinformatics

Fluxes

Neural networks

Data privacy

Unsupervised learning

Communication

Electronic mail

Biomolecules

Microarrays

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Medicine & Life Sciences (MeSH)

Sort by: Rank A-Z

INCLUDED CONCEPTS

Pharmaceutical Preparations

Computational Biology

Learning

Information Storage and Retrieval

Uncertainty

Genomics

Privacy

Neoplasms

Precision Medicine

Gene Expression

Metabolic Flux Analysis

Toxicogenetics

Genes

Cell Line

Fenofibrate

Chemical and Drug Induced Liver Injury

Magnetoencephalography

Biological Models

Chemical Safety

Biomarkers

Therapeutics

Terminal Repeat Sequences

Genome

Poisons

Pharmacogenetics

Benchmarking

Drug Combinations

HDL Lipoproteins

Type 1 Diabetes Mellitus