

BioSharing is a web-based, searchable portal of 3 interlinked registries, containing both in-house and crowd-sourced curated descriptions of **standards**, **databases** and **data policies**. Indicators of readiness and use highlight mature standards and databases endorsed by data policies from journals and funders.

It's so simple and quick to register your resource.

Go to biosharing.org/new to get started.

A curated, informative and educational resource on inter-related data standards, databases, and policies in the life, environmental and biomedical sciences

Find

Recommendations

Standards and/or databases recommended by journal or funder data policies.

Discover

Collections

Standards and/or databases grouped by domain, species or organization.

Learn

Educational

About standards, their use in databases and policies, and how we can help you.

Search

☒ Standards
 ☒ Databases
 ☒ Policies
 ☒ Collections/Recommendations

Advanced Search

Fine grained control over your search.

Search Wizard

ask biosharing

Let us guide you to your results.

670 Standards

Terminology Artifact	346
Model/Format	211
Reporting Guideline	113

View all

827 Databases

Protein	162
Genome	127
DNA	83

View all

85 Policies

Funder	20
Journal	59
Society	2

View all

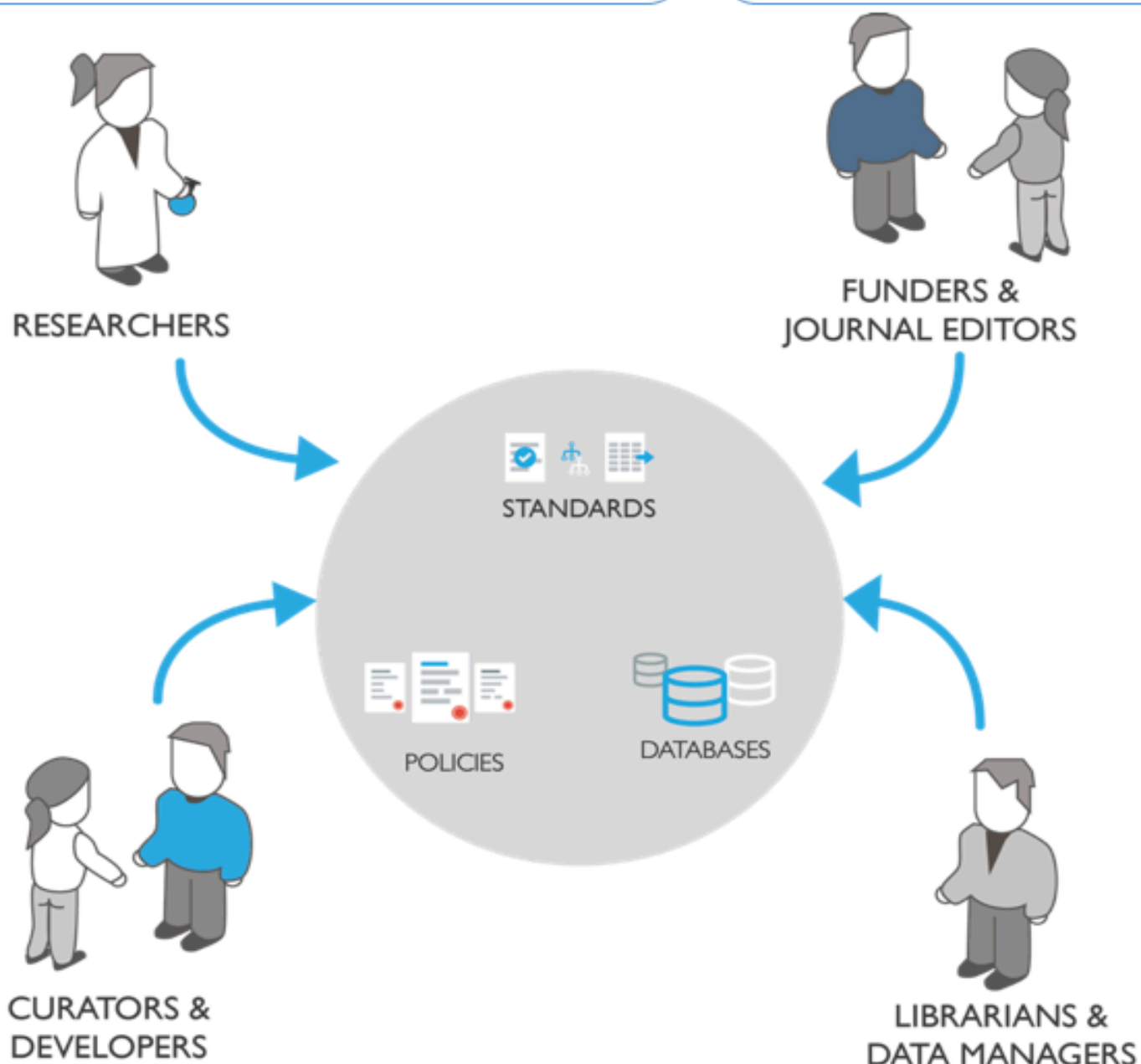
Our Advisory Board, collaborators and adopters include:



BioSharing enables you to make an informed decision as to which standard or database is appropriate to your use case, by mapping the relationships between standards and the databases that implement them, or the policies that recommend them.

My funder's data policy recommends the use of **established standards**, but which ones are widely **endorsed** and applicable to my **toxicological and clinical** data?

What are the **mature standards** and **standards-compliant databases** we should recommend to our authors?



Am I using the **most up-to-date version** of this terminology to annotate **cell-based assays**?

I understand this format has been **deprecated**; what has it been **replaced** by and is it appropriate for this data?