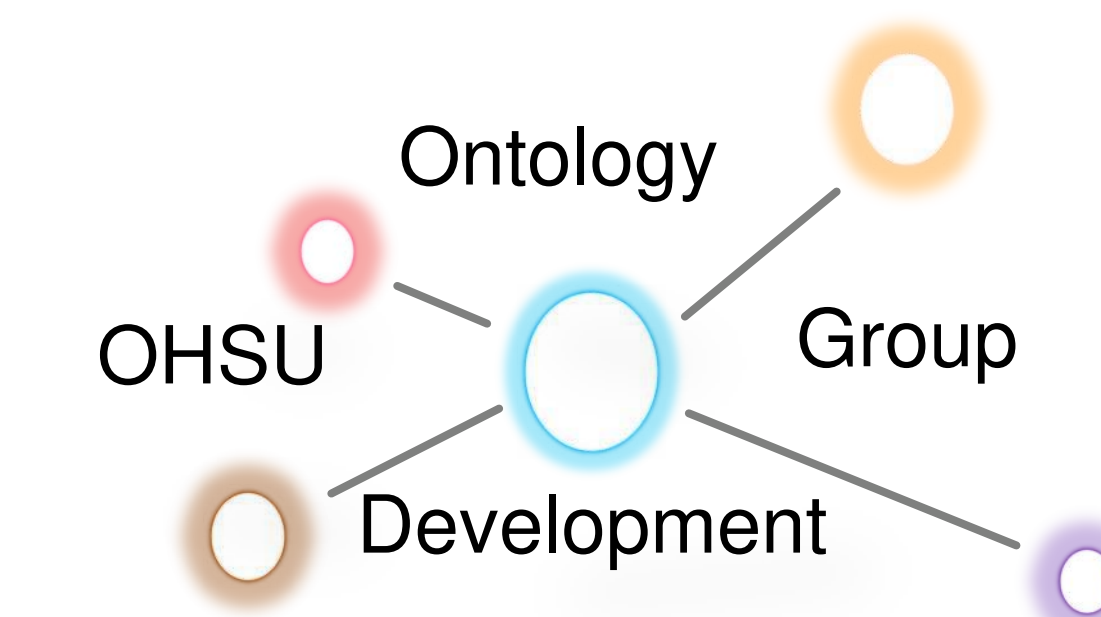


# On the Reproducibility of Science: Unique Identification of Research Resources in the Biomedical Literature



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## Introduction

Despite the proliferation and easy access to scholarly communications, a problem still exists - there is a significant lack of detailed information about the resources reported in publications, which hinders adequate research reproducibility. In cases such as antibodies and model organisms, this lack of unique reference makes it difficult or impossible to reproduce the experiments. In order to better understand the magnitude of this problem, we designed an experiment to evaluate the “identifiability” of research resources in the biomedical literature.

## Methods

### 5 Resource Types:

Model organisms
Antibodies
Cell lines
Knockdown reagents
Constructs

### 5 Domains:

Cell Biology
Developmental Biology
General Biology
Immunology
Neuroscience

### Example criteria for identifiability:

- ☒ Source reported
- ☒ Identifiable in vendor site
- ☒ Identifiable in MOD
- ☒ Catalog number reported

### 3 impact factors

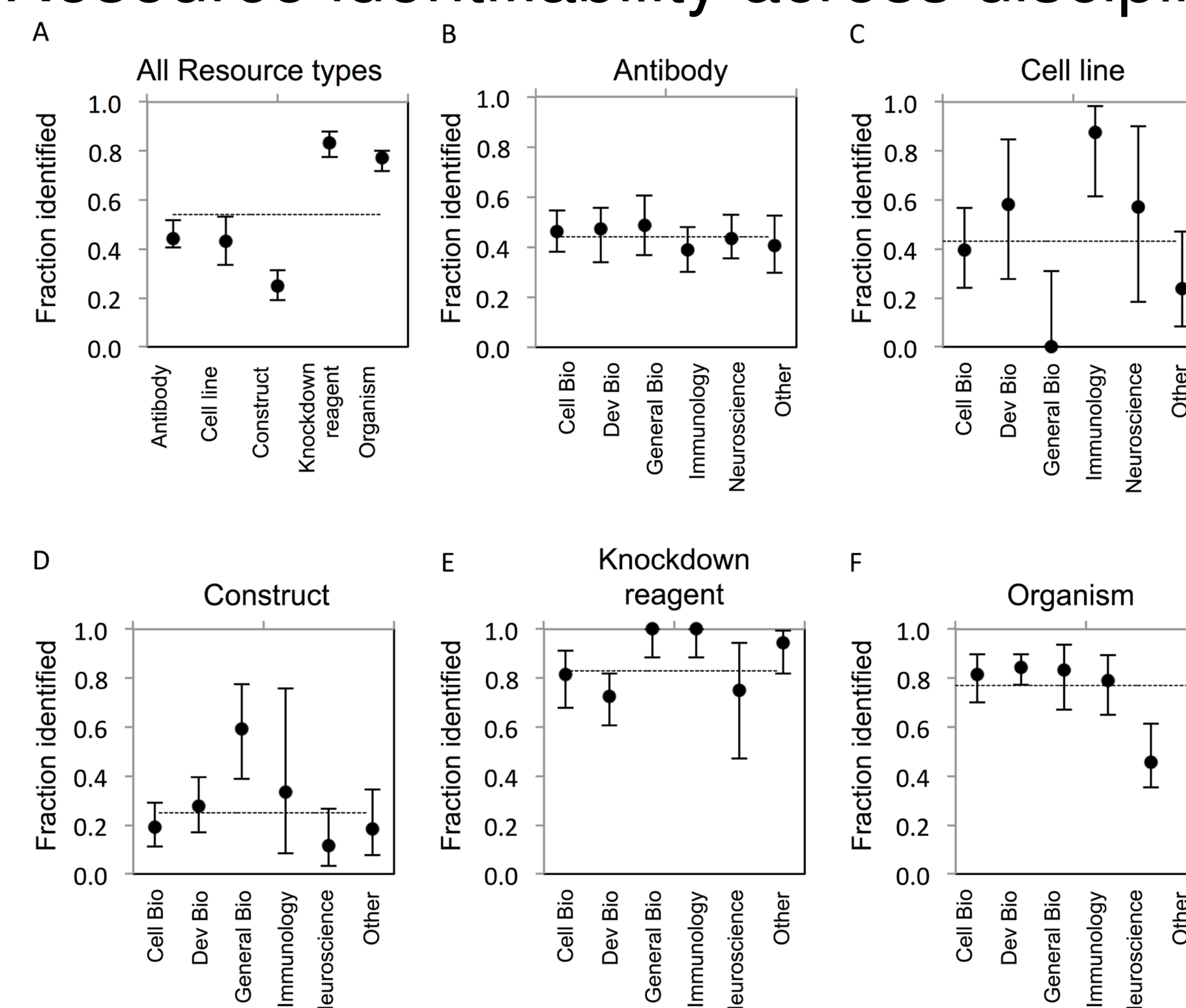
High impact
Mid impact
Low impact

### 3 Reporting guidelines

Stringent
Satisfactory
Loose

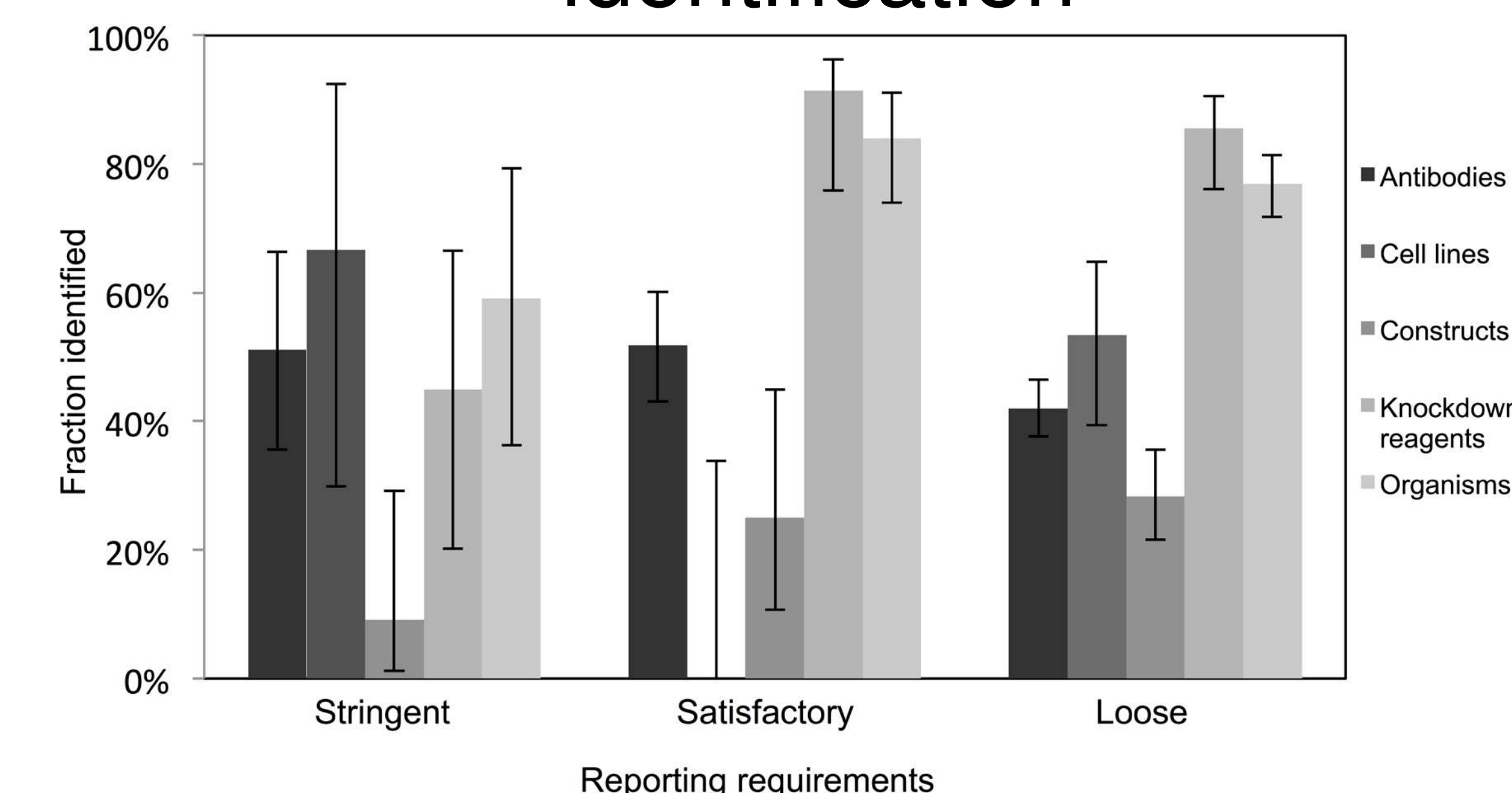
**238 papers were curated**

## Resource identifiability across disciplines



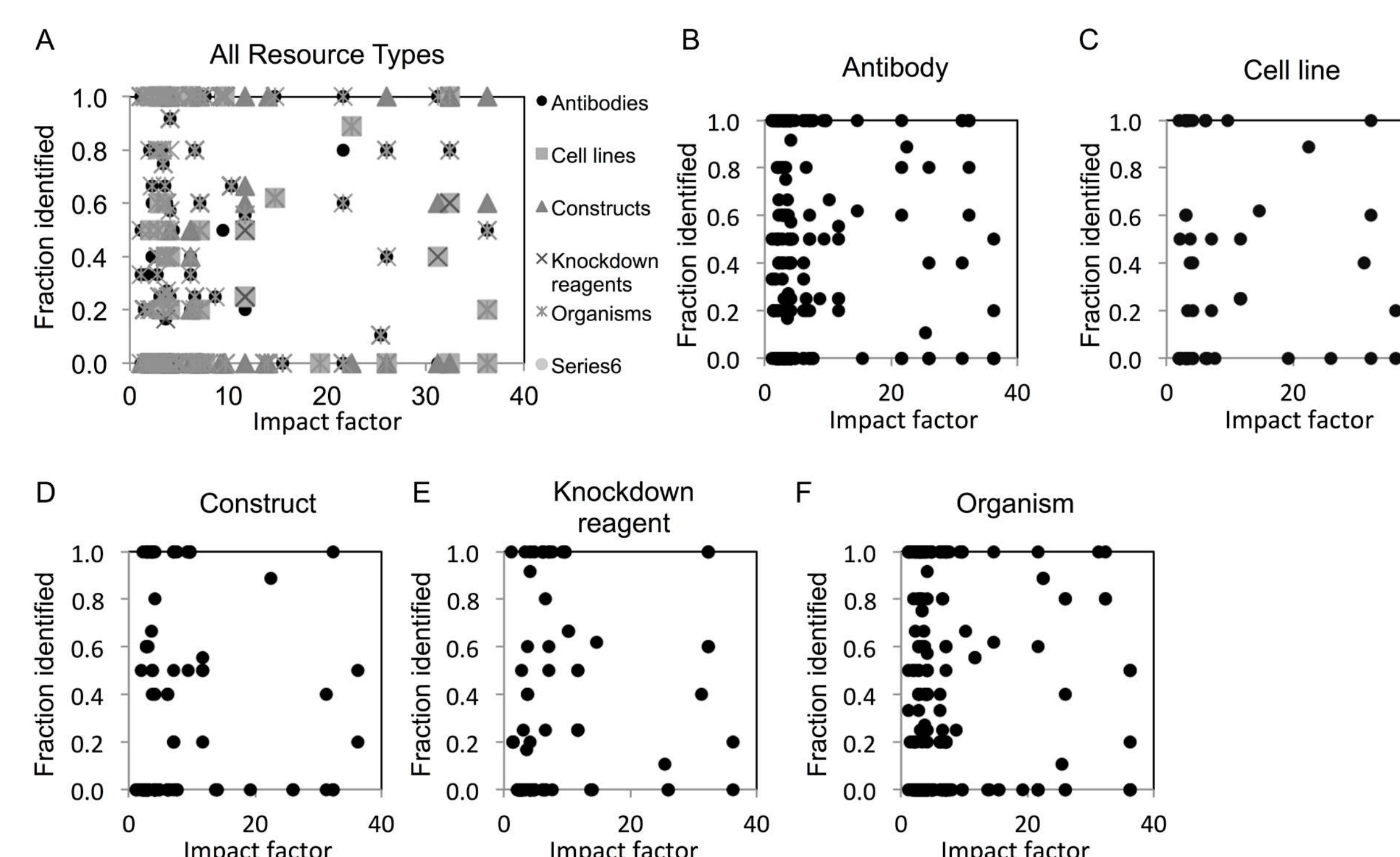
(A) Summary of average fraction identified for each resource type. (B–F) Identifiability of each resource type by discipline.

## Stringent resource reporting requirements does not improve resource identification



The reporting requirements for each journal were classified as stringent, satisfactory or loose. A total of 53 out of 118 resources were identifiable in the stringent reporting guidelines category, 201 resources were identifiable out of 329 resources for the satisfactory category and 662 out of 1,217 resources were identifiable in the loose category.

## Resource identification rates across journals of varying impact factors



(A) An overview of fraction identified by impact factor for all resource types. (B–F) Fraction identified by impact factor for each individual resource type. Increasing height on the x-axis corresponds with a higher impact factor for each journal.

## Recommended reporting guidelines for life science resources

**FORCE11** <http://www.force11.org/node/4433> **biosharing.org** <http://biosharing.org/bsg-000532>



## Conclusions:

- ☐ Inability to identify resources hinders reproducibility
  - ☒ Improve metadata standards for tracking resources, authors should provide unique IDs in publications
- ☐ Current reporting standards are insufficient to uniquely identify resources
  - ☒ Publishers, editors, and reviewers should work together to increase reporting requirements

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