

# FAIR Phenotyping with APHRODITE

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# The Need

- The common failure to reproduce published results has created an atmosphere of crisis even in disciplines where precise measurement and tight experimental control are the norm
- There is even more reason for vigilance in disciplines that must manage lower degrees of measurement accuracy and experimental control
- One response to this crisis has been the emergence of open science principles that publicly expose the process of defining hypotheses, data selection and development, study design and analytic choices

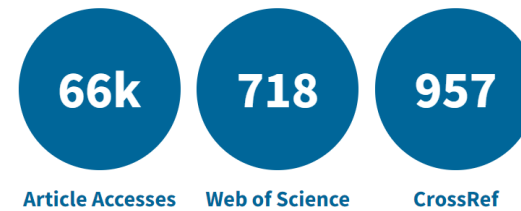
# The FAIR Guiding Principles for scientific data management and stewardship

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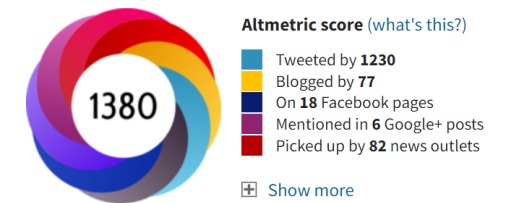
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# Rapid adoption of principles:



European Commission > Press releases database > Press Release details  
European Commission - Statement

## G20 Leaders' Communique Hangzhou Summit

Hangzhou, 5 September 2016

1. We, the Leaders of the G20, met in Hangzhou, China on 4-5 September
2. We met at a time when the global economic recovery is progressing, re are emerging. But growth is still weaker than desirable. Downside risks rei of commodity prices, sluggish trade and investment, and slow productivity from geopolitical developments, increased refugee flows as well as terroris
3. We also met at a time of continued shifts and profound transformations for growth. With these transformations come challenges and uncertainties determine the effectiveness of our response to the challenges of today and

Realising  
the European  
Open Science Cloud

## Progress towards the European Open Science Cloud

News item | 01-12-2017 | 18:00

Germany and the Netherlands establish office for the GO FAIR Initiative / France joins



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## Big Data to Knowledge

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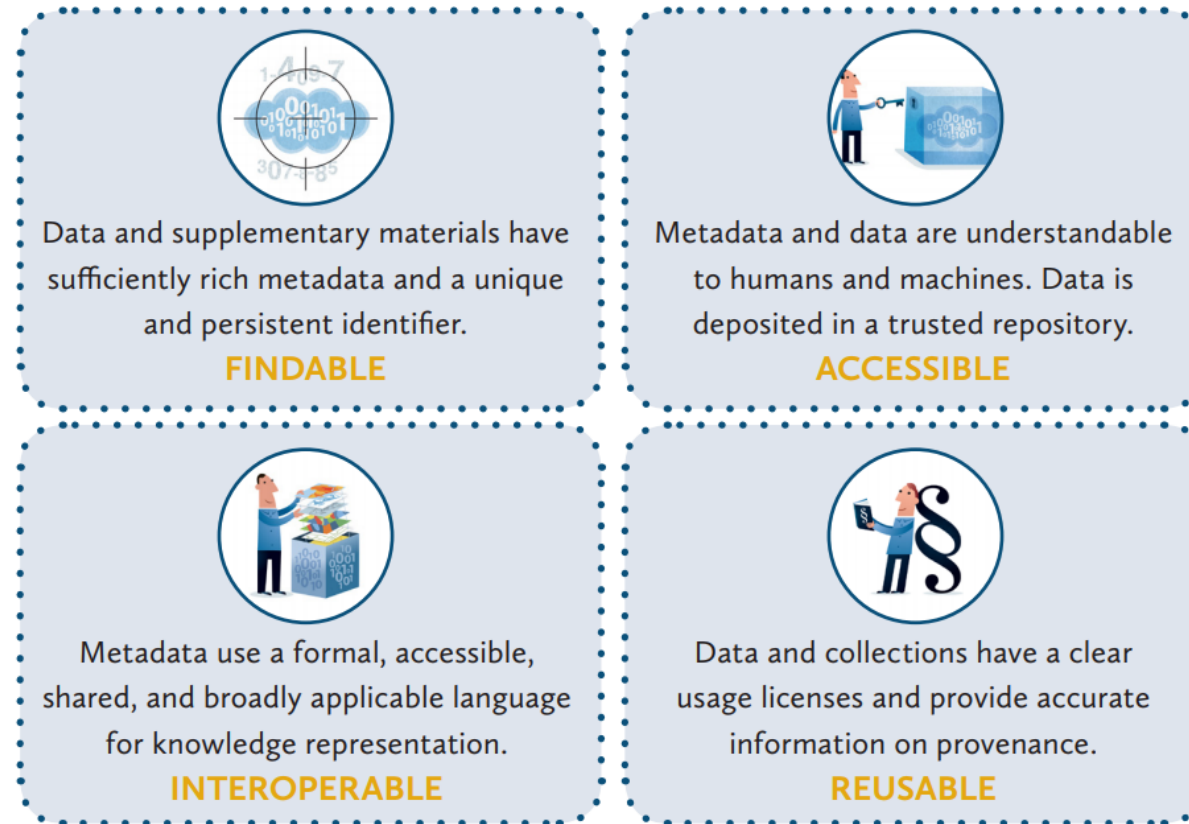


Announcements

# What does it mean to be FAIR?

- **FAIR**
- Findable,
- Accessible,
- Interoperable,
- Reusable.

## What is FAIR DATA?



# What are we proposing?

## **Anatomy of an APHRODITE FAIR phenotype definition**



# A phenotype definition will be Findable

- To address the need to have a persistent global unique resource identifier (URI) for each phenotype definition version, we have utilized GitHub unique commit hash value to identify each individual phenotype definition version
- The OHDSI Gold Standard Phenotype Library workgroup has defined and created an additional abstraction layer over the phenotype definitions available as a R Shiny App

# A phenotype definition will be Accessible

- The phenotype definition, generation script, and metadata will be retrievable by their identifier using any regular web browser or the application layer of the phenotype library
- By using a publicly and freely available resource such as GitHub, we offer better accessibility than placing the definitions on an institutional server



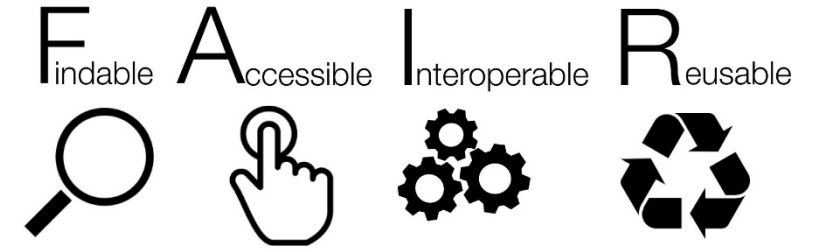
# A phenotype definition will be Interoperable

- We will leverage the OMOP CDM and associated vocabularies to solve the major obstacle to interoperability across sites. Our phenotype definitions' metadata will use JSON for knowledge representation and ease of machine readability
- When developing phenotyping definitions based on prior publications, or when a publication is generated from a definition generated from our pipeline, we will include all proper URI's to the publications in question

# A phenotype definition will be **Re-usable**

- Currently APHRODITE definitions are easily shareable and re-usable for other sites. We have added meta-data elements related to software, CDM, and vocabulary versions, as well as a plurality of accurate and relevant attributes to guarantee re-usability
- All the publicly available phenotypes will be released under relevant open source licenses, details of which will be attached to the definition's meta-data
- Site and researcher information will be recorded as well as relevant publications in allowing fully traceable provenance for each definition

# Questions?



Improving the FAIRness of digital resources will increase their quality and their potential for reuse

@micheldumontier::RDA:2018-01-31

Want to help? reach out: @drjmbanda or jbanda@gsu.edu