

1 cBioPortal new study report

Report for study: ./acyc_mda_2015/raw_data_clinical.txt

1.1 Attribute matches

Below are the possible matches between attributes from existing data on the cBioPortal and the new study. The metric used to detect each match is denoted by the symbols following the attribute name of the match. Additionally, the number of studies in which the matching attribute occurs is given to indicate how popular the attribute is among existing studies.

New study attribute	Possible matches
AGE	AGE^ (found in 117 other studies)
FISH	No matches found
FUSION_OTHER	No matches found
HISTOLOGY	HISTOLOGY^ (found in 12 other studies)
HISTOLOGY	KATABI_PATH* (found in 1 other studies)
HISTOLOGY	OFFICIAL_PATH* (found in 1 other studies)
MYB/MYBL1-FUSION	No matches found
MYB_MYBL1_FUSION_METHOD	No matches found
MYB_NFIB_REARRANGEMENT	MYB_NFIB_REARRANGEMENT^ (found in 1 other studies)
OTHER NFIB FUSION	No matches found
PNI	No matches found
SEX	SEX^* (found in 32 other studies)
SEX	GENDER* (found in 81 other studies)
SITE	No matches found
SIZE(CM)	No matches found

^ represents matches found based on the attribute names

* represents matches found based on clustering of the attribute values

1.2 Number of attribute distributions

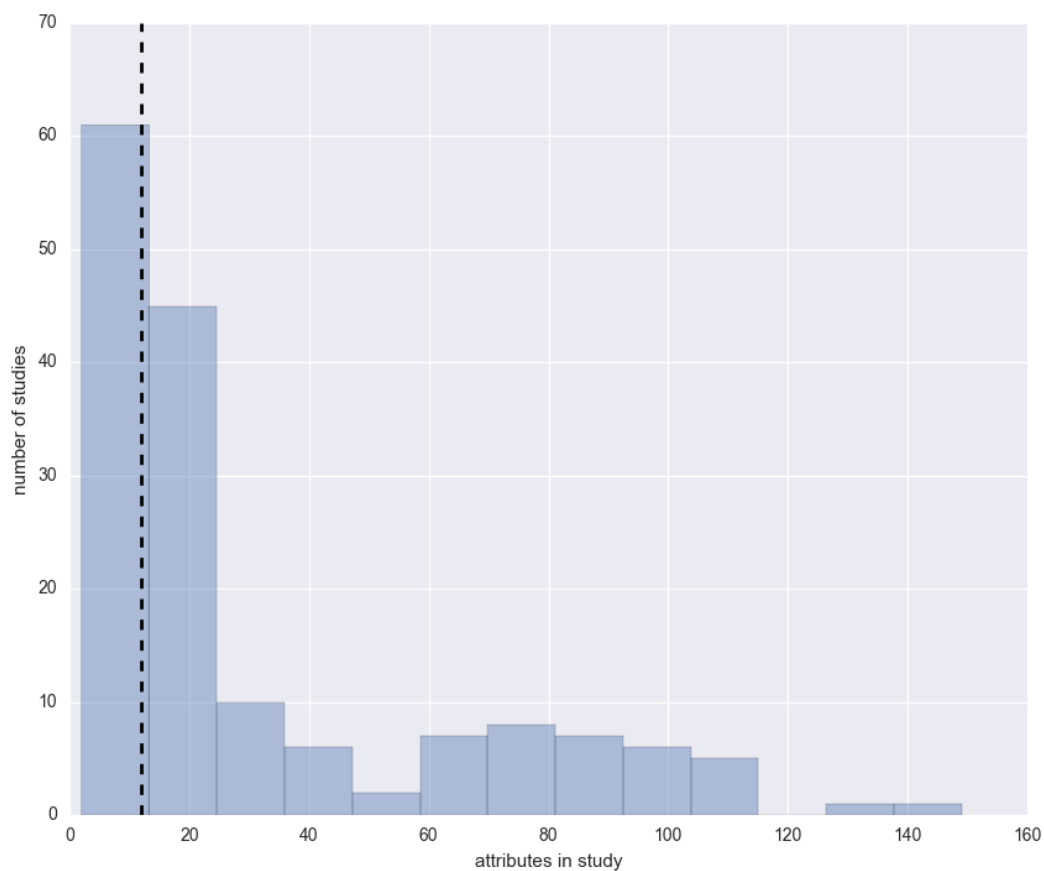


Figure 1: Comparison between the number of attributes in the new study and the number of attributes in each existing study on cBioPortal. The dashed black line indicates the number of attributes in the new study, while the histogram shows the data for existing cBioPortal studies.

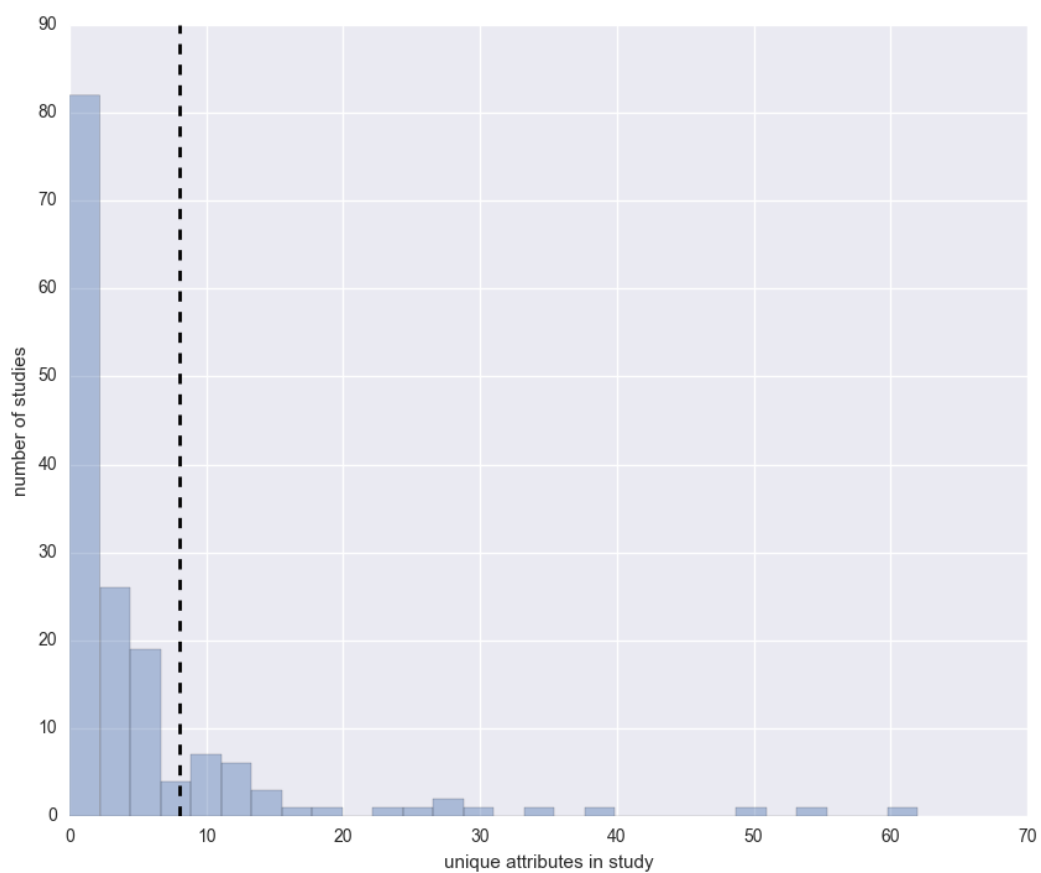


Figure 2: Comparison between the number of unique attributes in the new study and the number of unique attributes in each existing study on cBioPortal. The dashed black line indicates the number of unique attributes in the new study, while the histogram shows the data for existing cBioPortal studies.

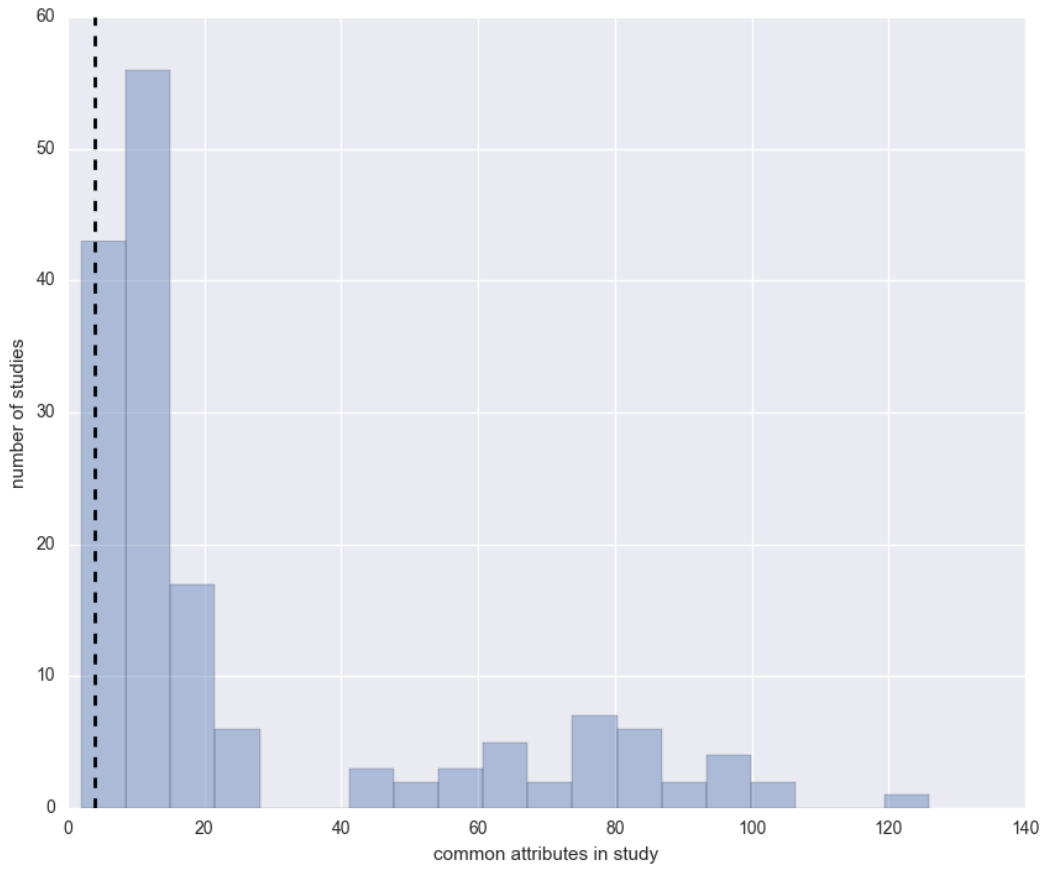


Figure 3: Comparison between the number of common attributes in the new study and the number of common attributes in each existing study on cBioPortal. The dashed black line indicates the number of common attributes in the new study, while the histogram shows the data for existing cBioPortal studies.