

Mass spectrometry data  
Mass spectrum  
NMR spectrum  
Mass spectrometry spectra  
Peptide mass fingerprint  
Gene expression matrix  
Protein sequence  
DNA sequence  
Database metadata  
Compound name  
ChemSpider ID  
Atomic occupancy  
Compound identifier  
ID list  
Ecological data  
Gene identifier  
Molecular mass  
Peptide identification  
Small molecule structure  
Pathway or network  
Map set data  
Sequence record  
Version information  
Raw microarray data  
KEGG object identifier  
Expression data  
Molecule name  
Sequence assembly  
Matrix  
Ontology mapping  
Protein identifier

Mass spectrometry data format

Textual format

mzXML

TSV

XML

protXML

FASTA

Graph format

raw

Biodiversity data format

DSV

dat

Binary format

BLOM format

Data index format

Format

GenBank format

imzML metadata file

mzQuantML

R file format

Thermo RAW

netCDF

idXML

GML

ibd

mzIdentML

pepXML

TIDE

TXT

xlsx

MGF

dta

CSV

OBO

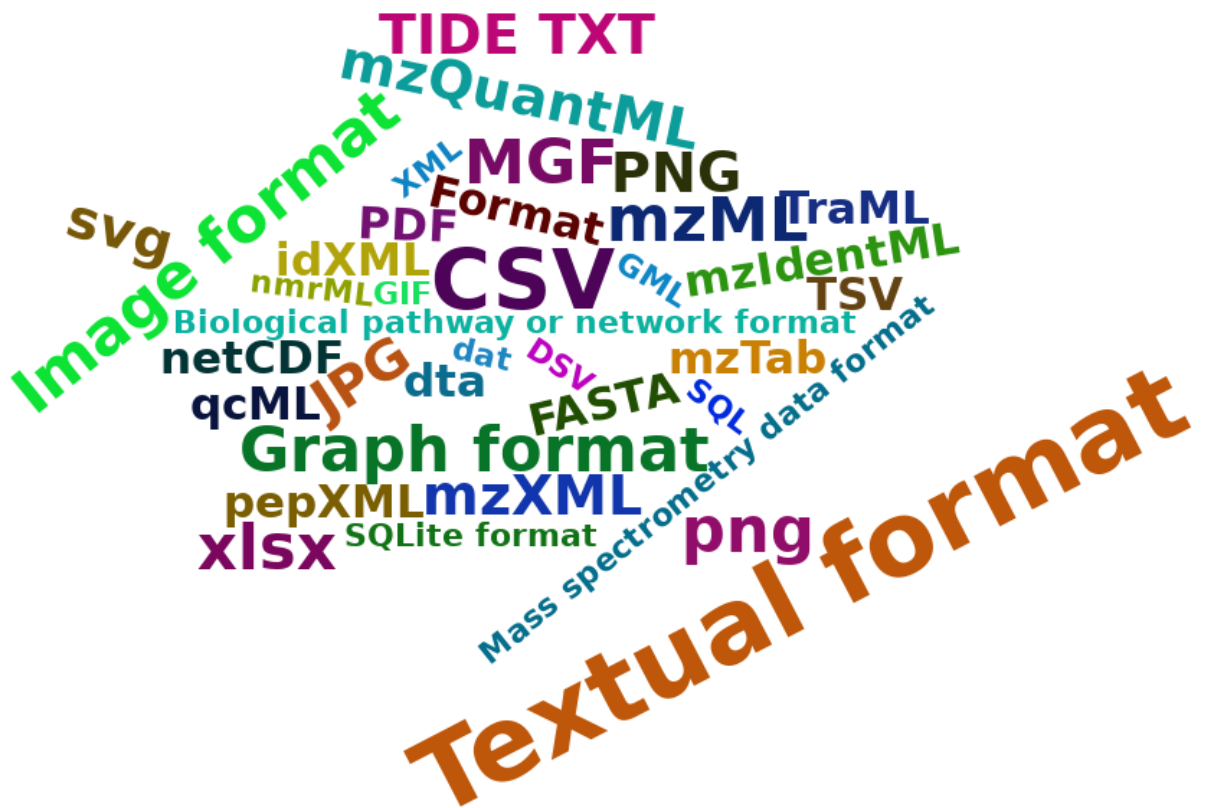
TraML



**Mac**  
**Linux**  
**Windows**



Peptide mass fingerprint  
Mass spectrometry data  
Protein identifier  
Image  
Ontology  
Plot  
Peptide identification  
Expression data  
Gene identifier  
Compound ID (HMDB)  
Chemical formula  
Molecule name  
Q-value  
Alignment  
PubChem ID  
Clustered expression profiles  
Compound name  
Data reference  
Matrix  
Score  
Compound identifier  
NMR spectrum  
Heat map  
Mass spectrum  
Report  
Sequence-structure alignment  
Version information  
Pathway or network

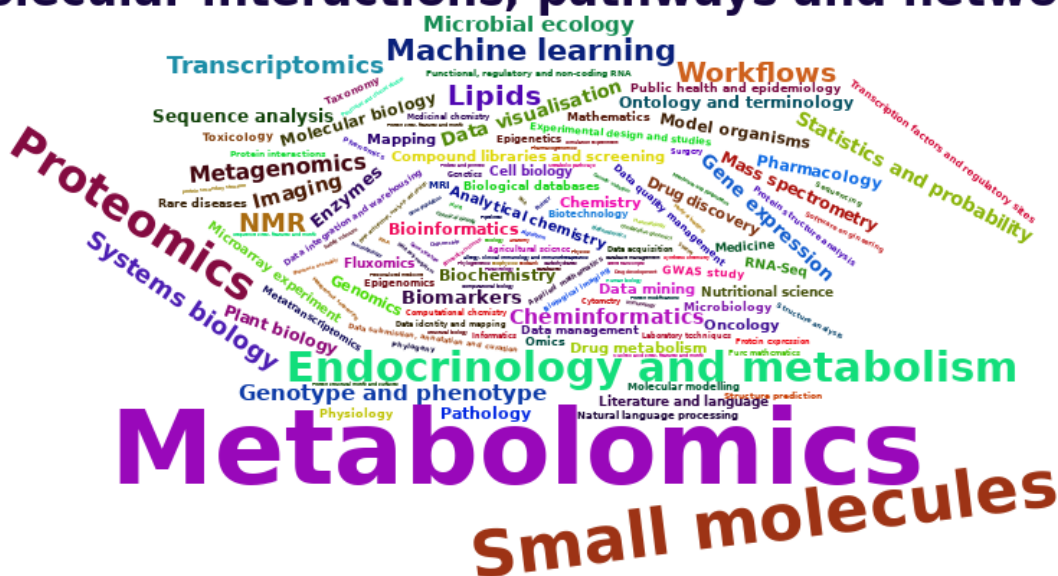


**Web application**  
**Desktop application**  
**Database portal**  
**Workbench**  
**Script**  
**Workflow**  
**Web API**  
**Web service**  
**Library**  
**Command-line tool**  
**Bioinformatics portal**  
**Plug-in**  
**Suite**  
**SQL endpoint**  
**CloudAPI**

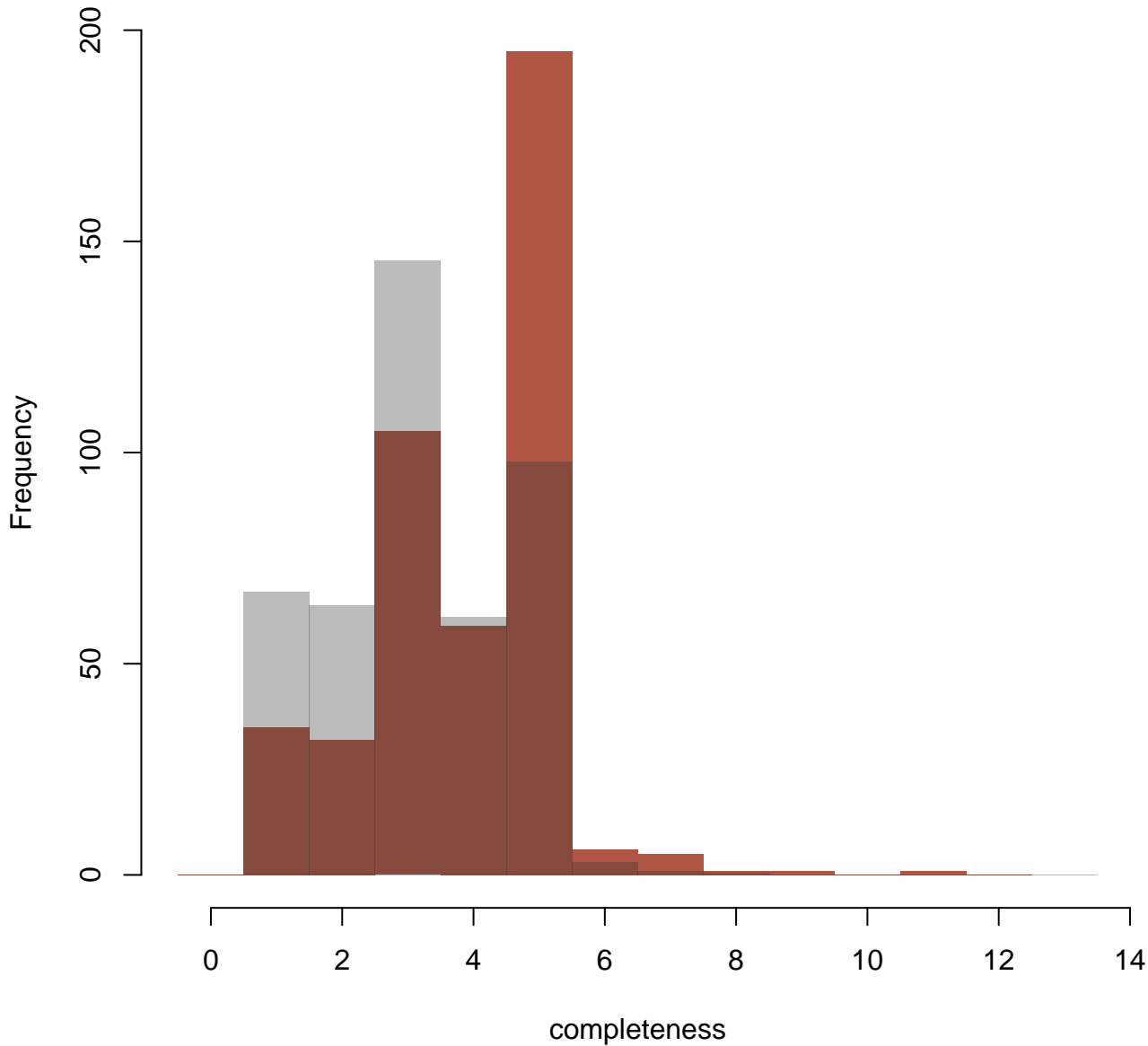


# Proteomics experiment

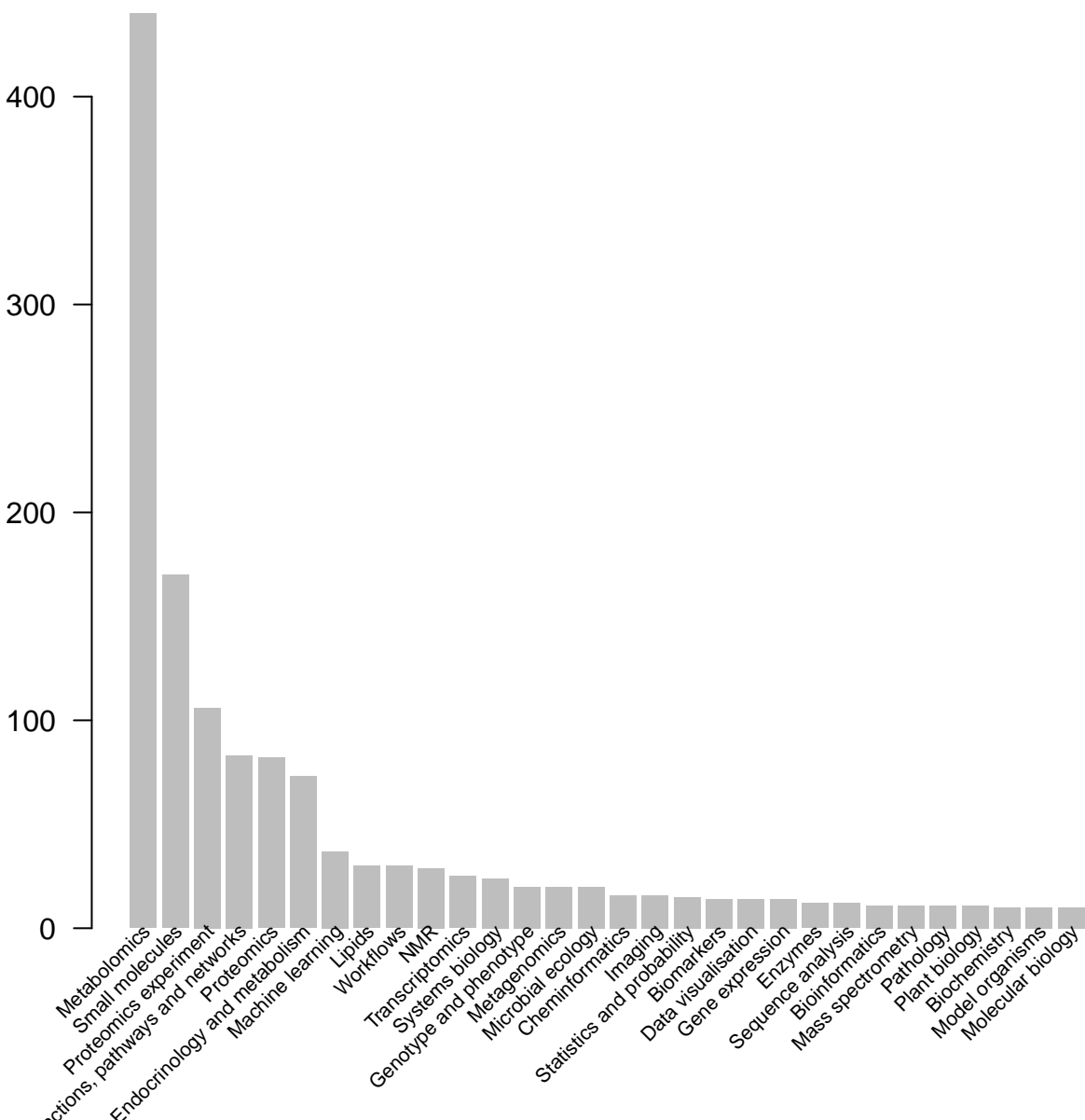
## Molecular interactions, pathways and networks



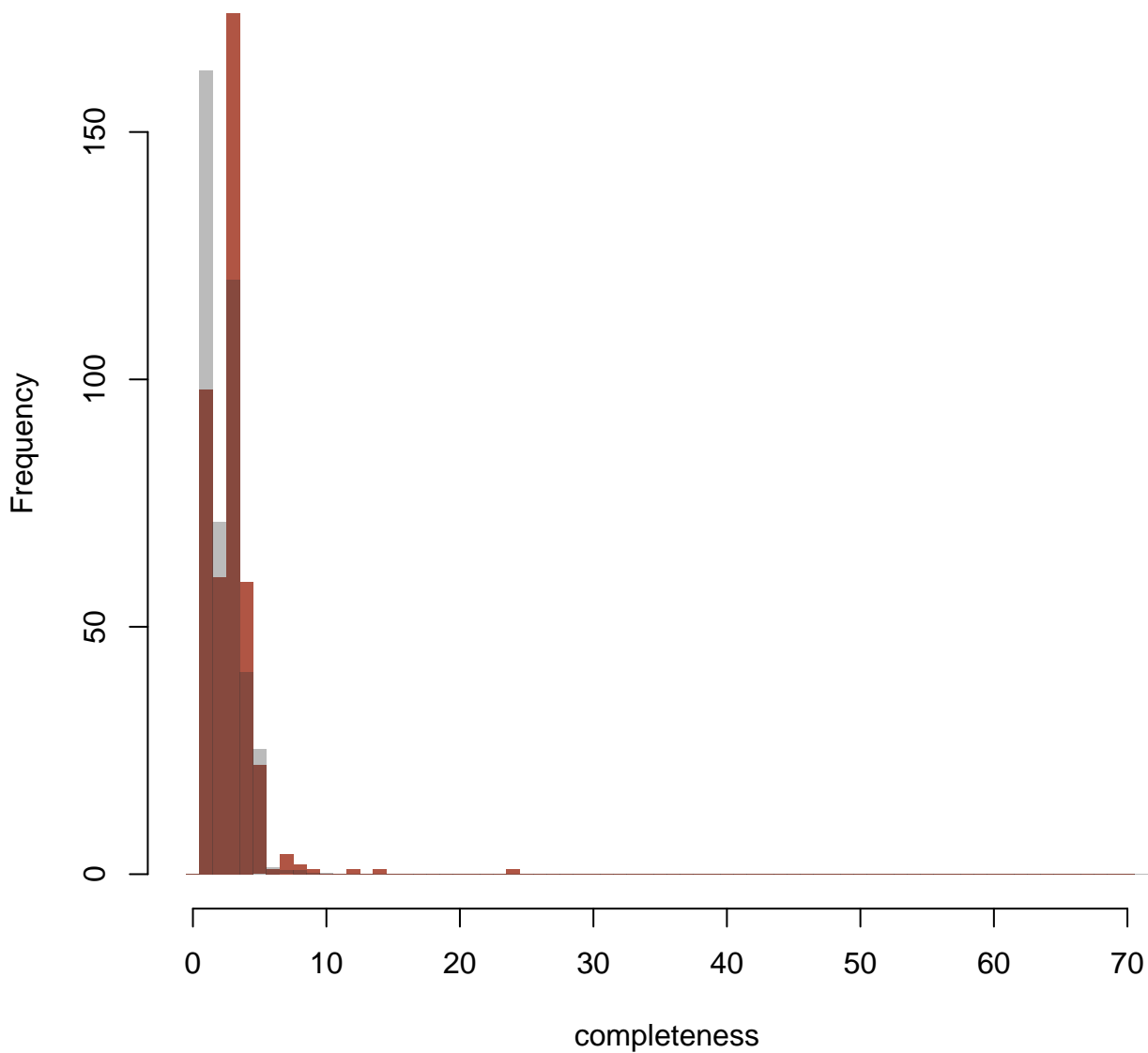
**Topic Number of terms per tool**



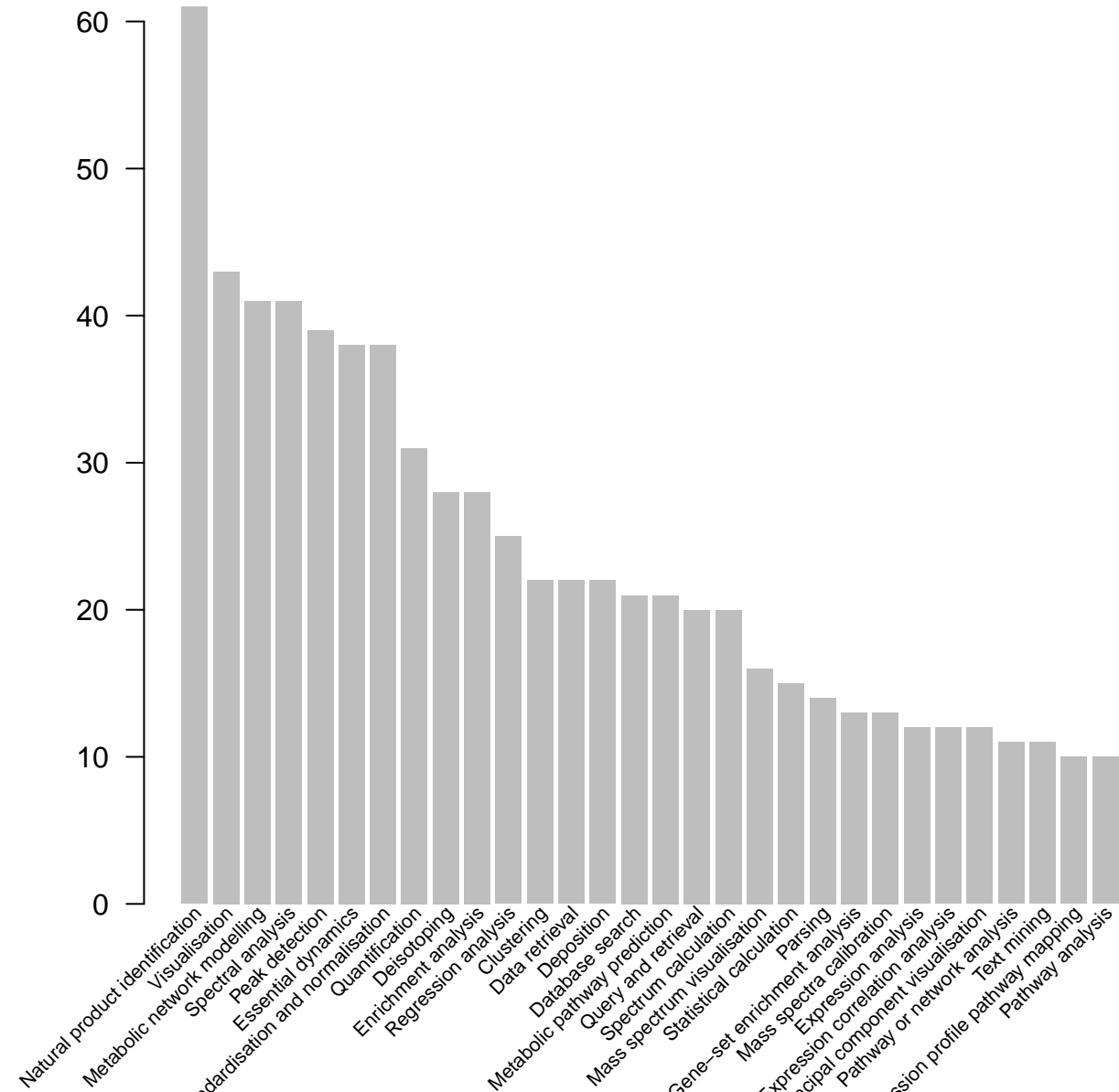
## Topic Top 30 terms



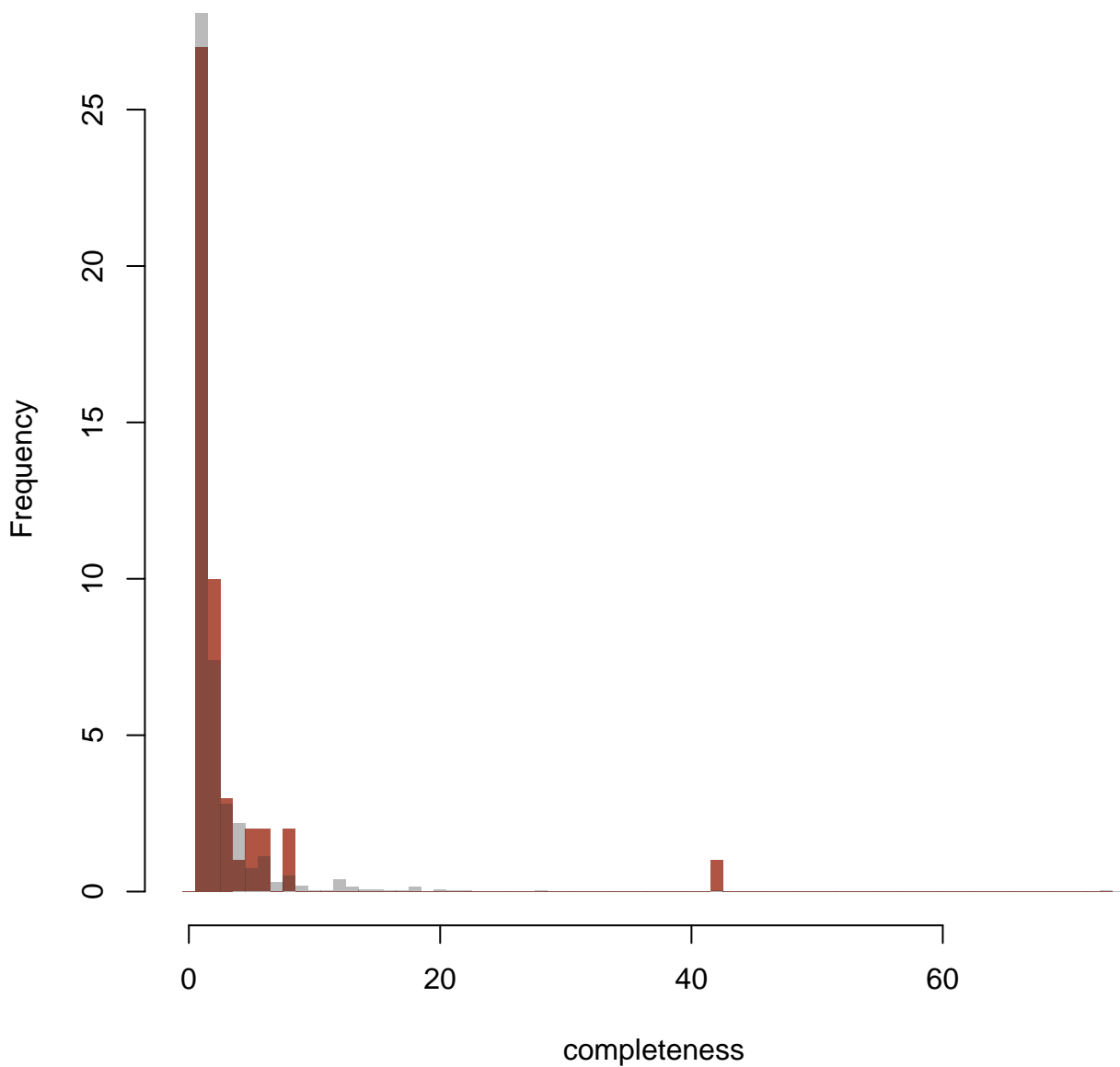
## Operation Number of terms per tool



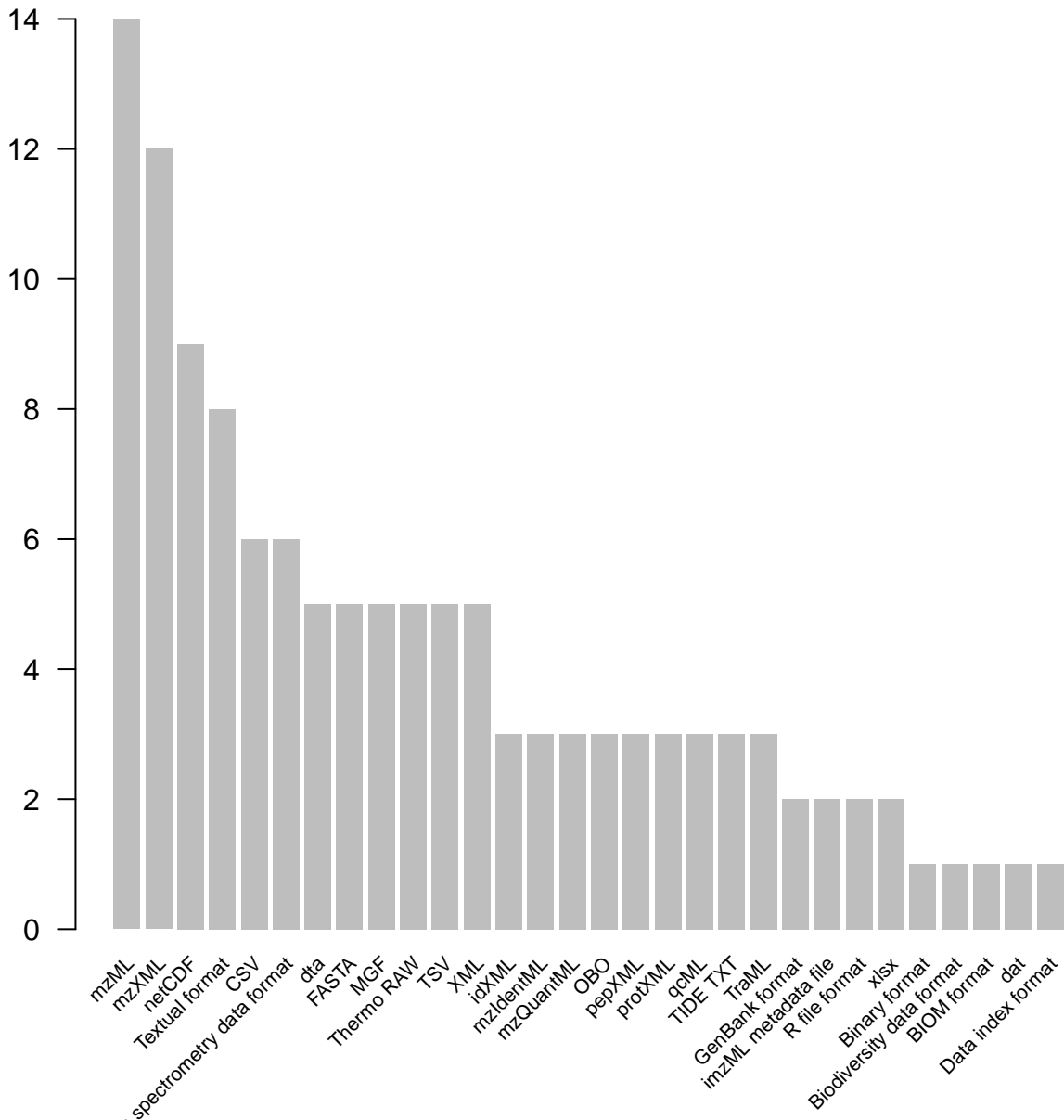
## Operation Top 30 terms



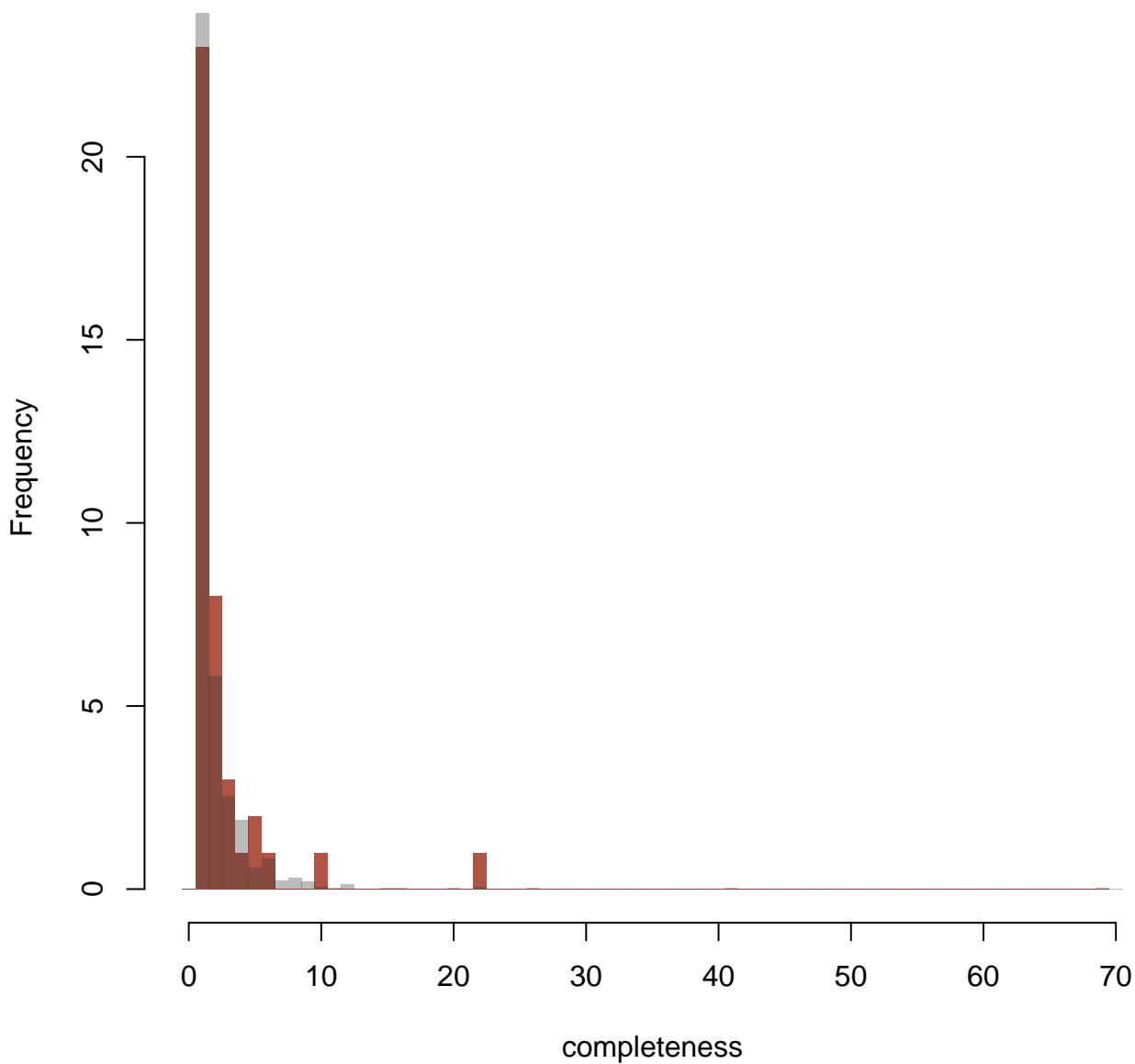
# Input format Number of terms per tool



## Input format Top 30 terms

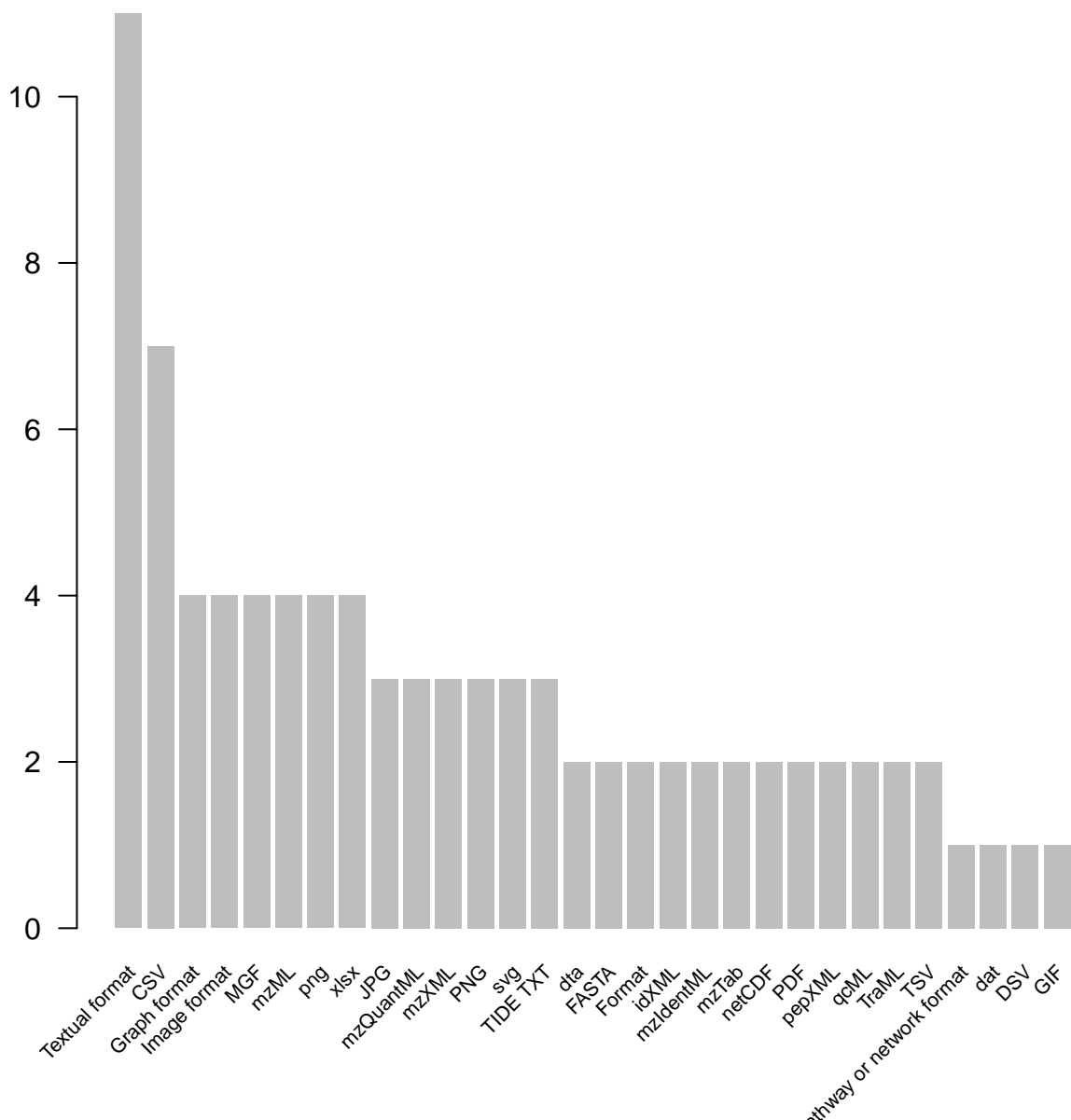


**Output format Number of terms per tool**

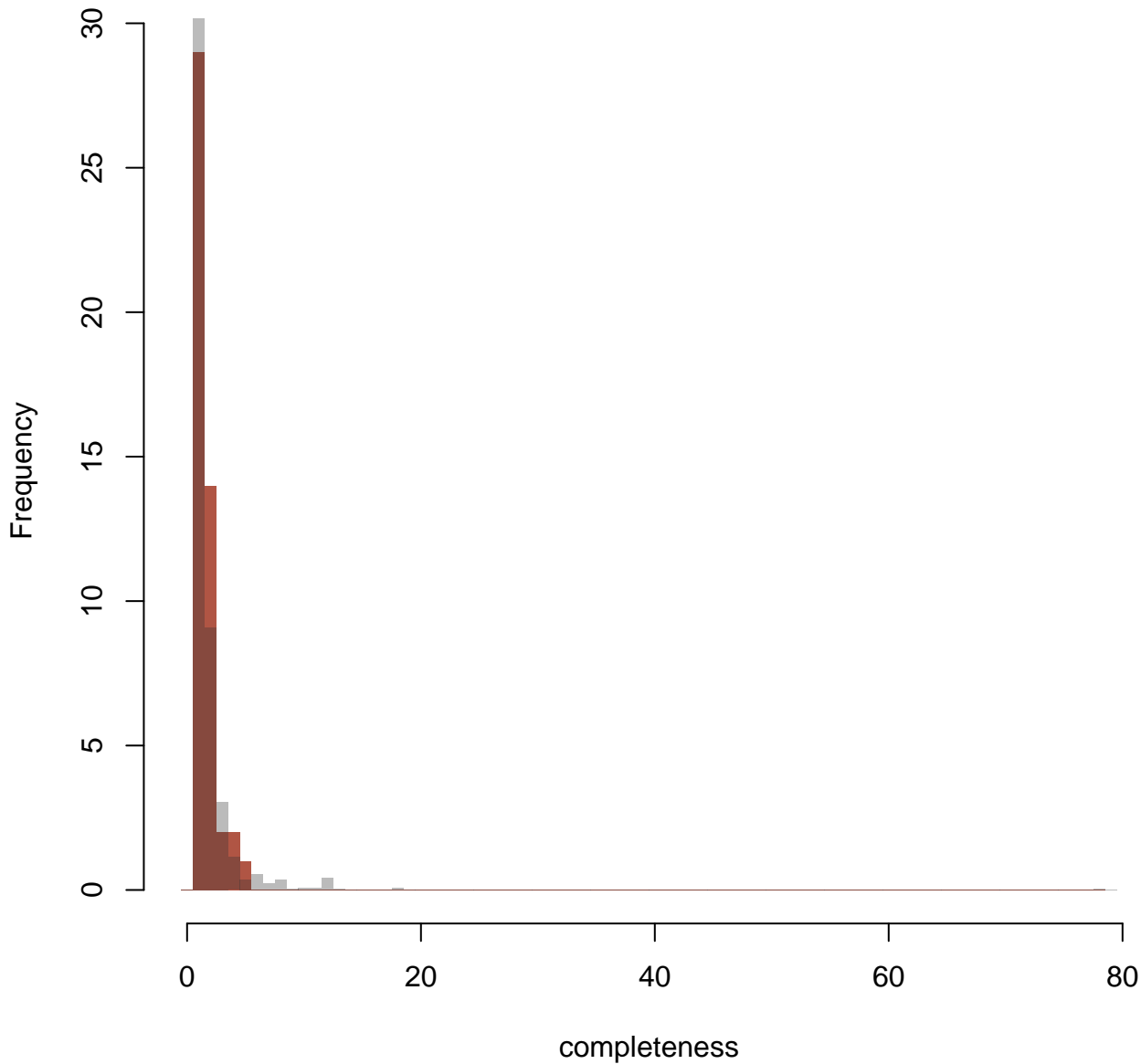




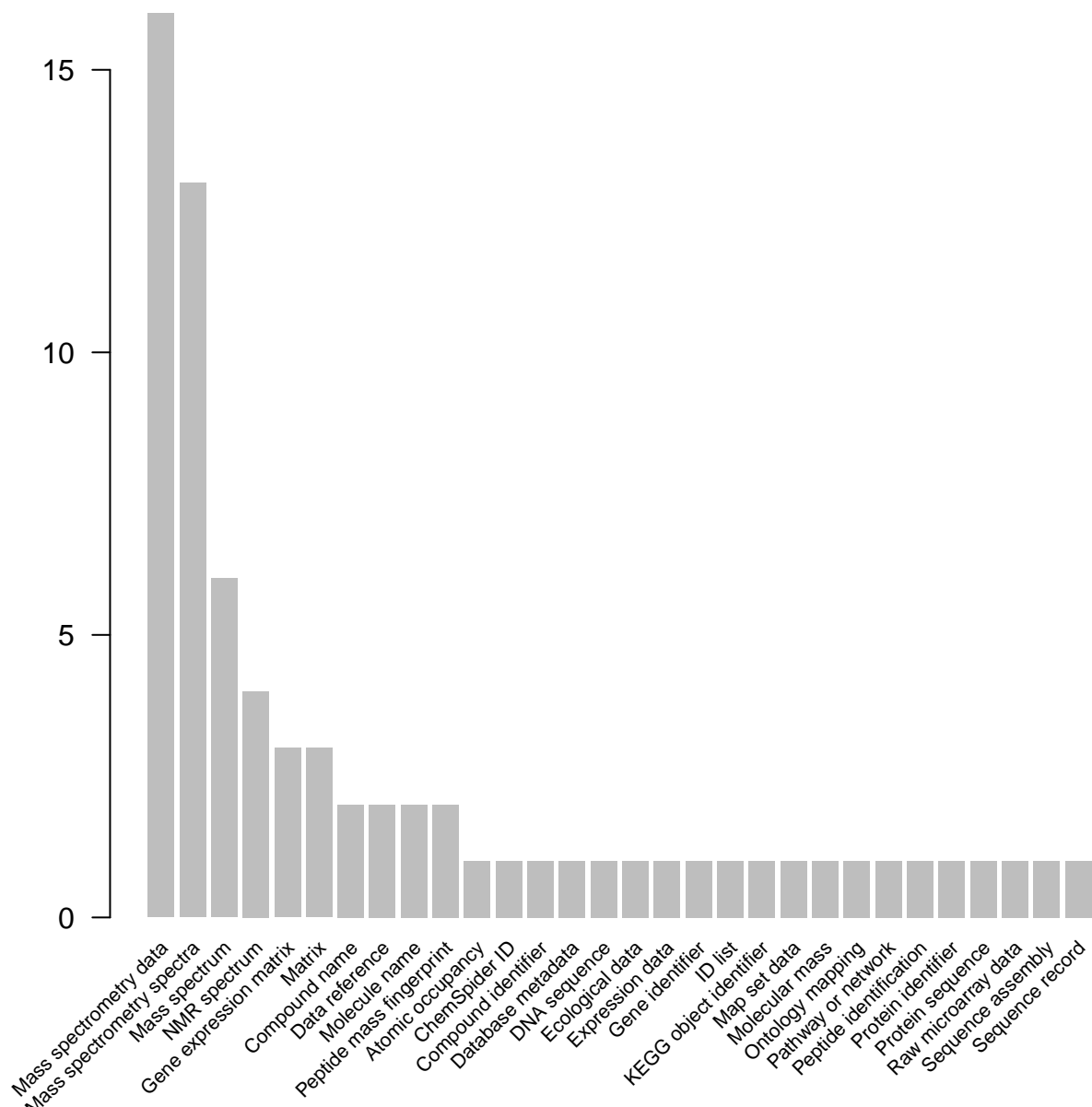
## Output format Top 30 terms



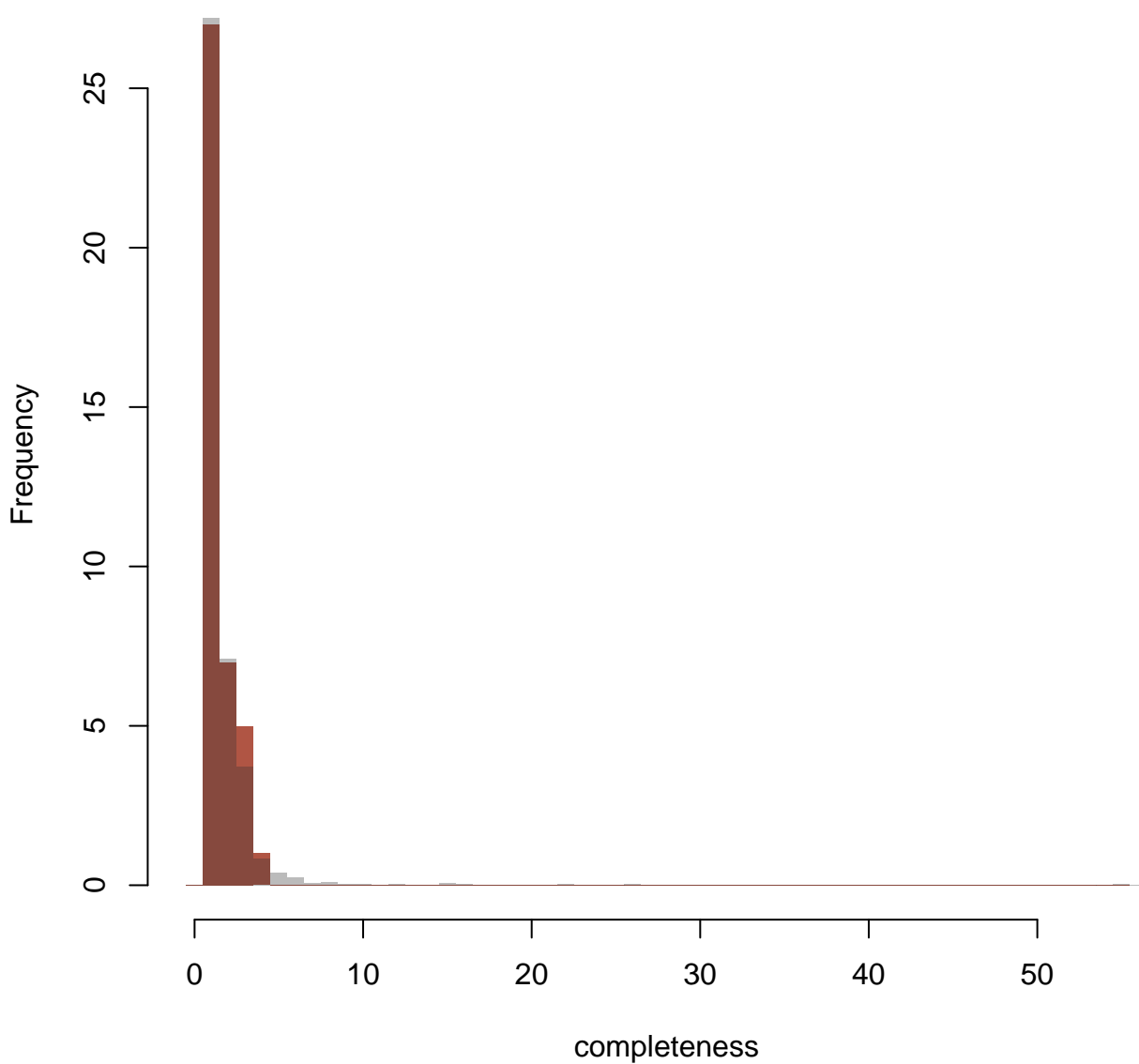
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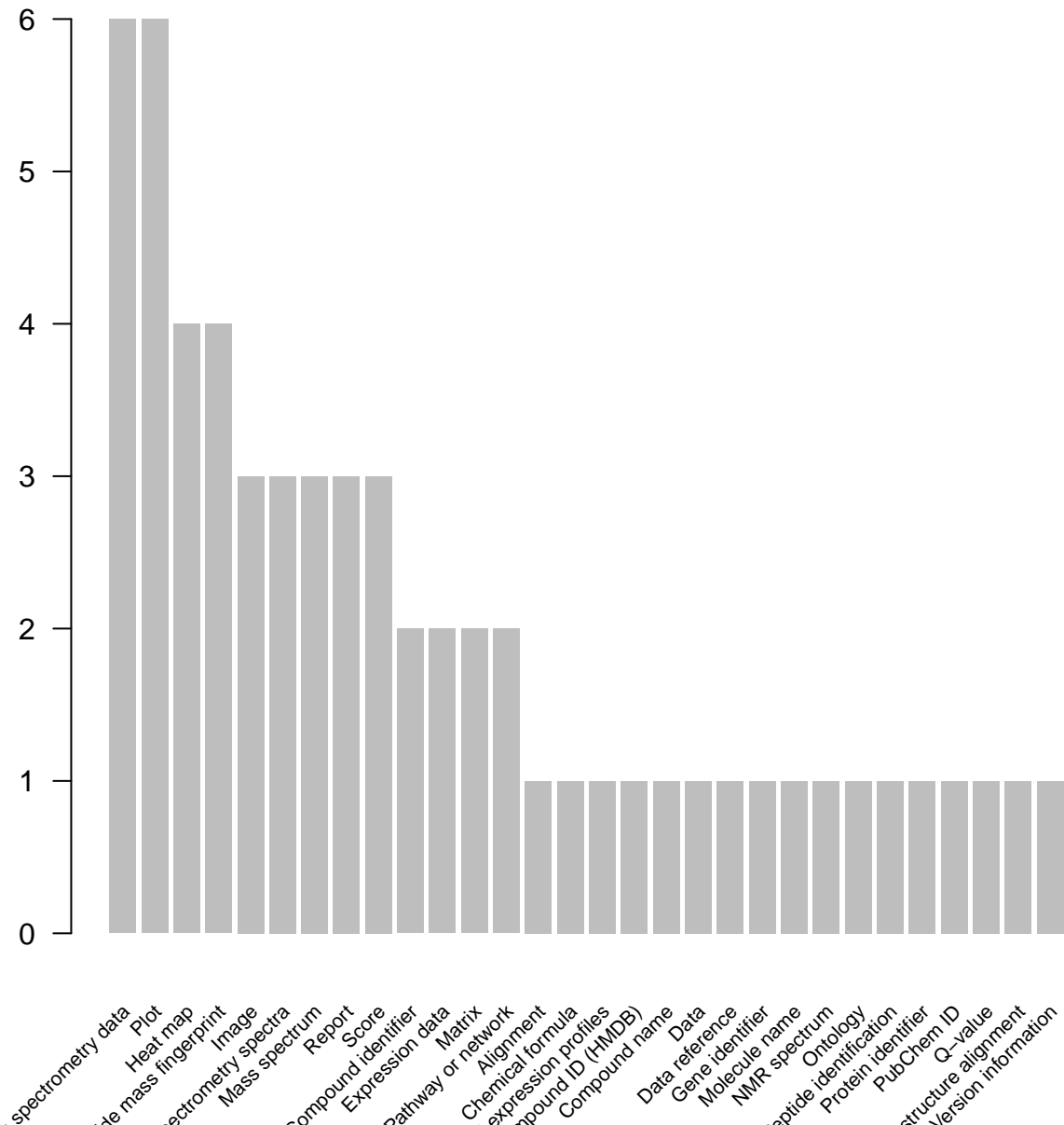
## Input data Top 30 terms



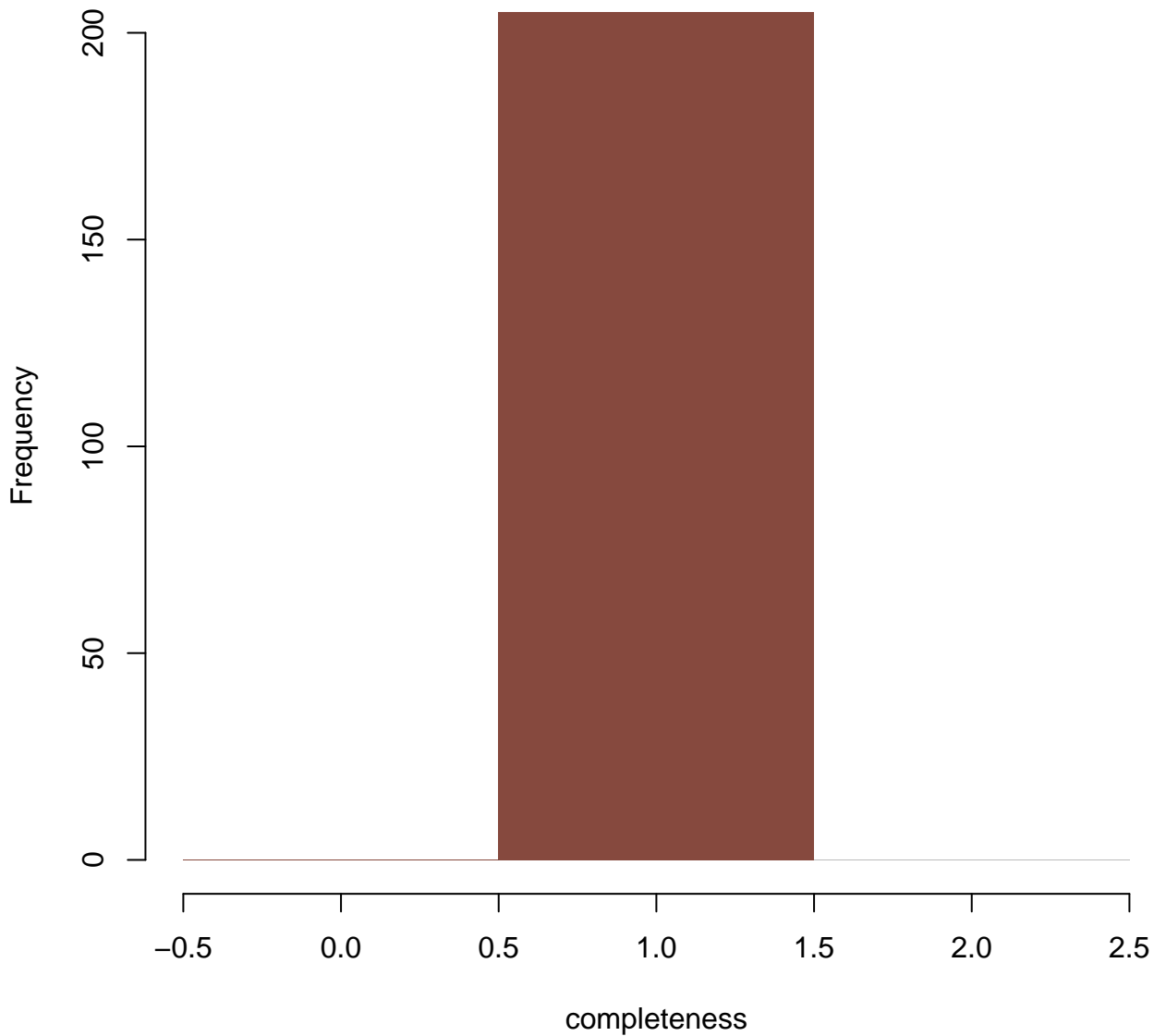
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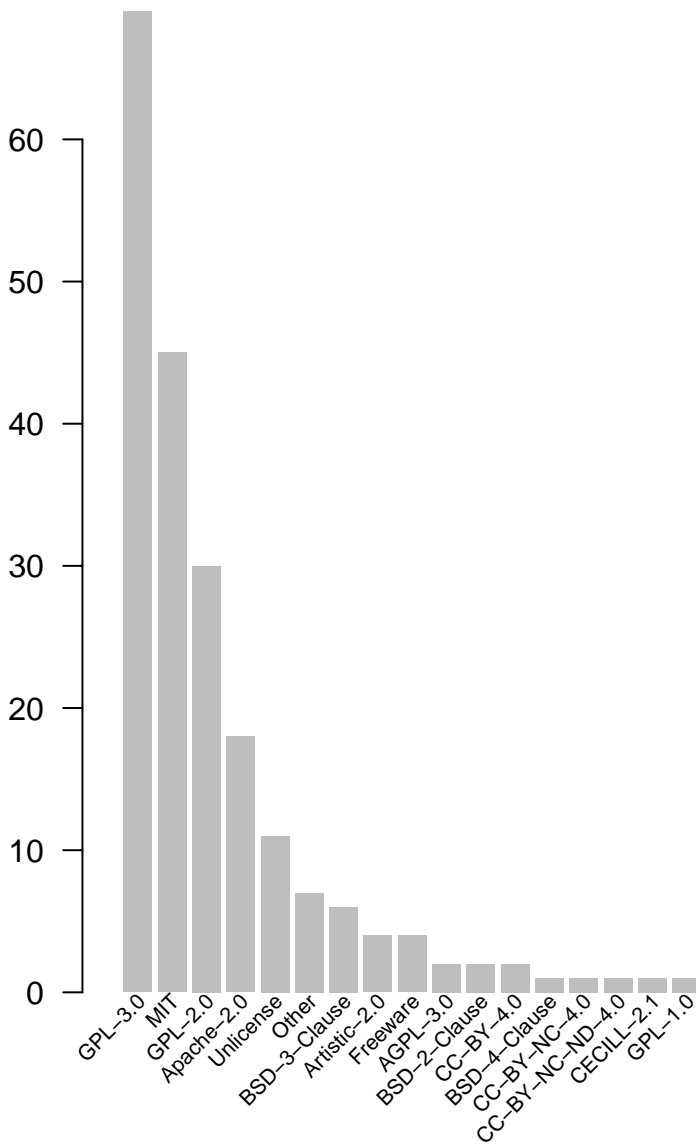
## Output data Top 30 terms



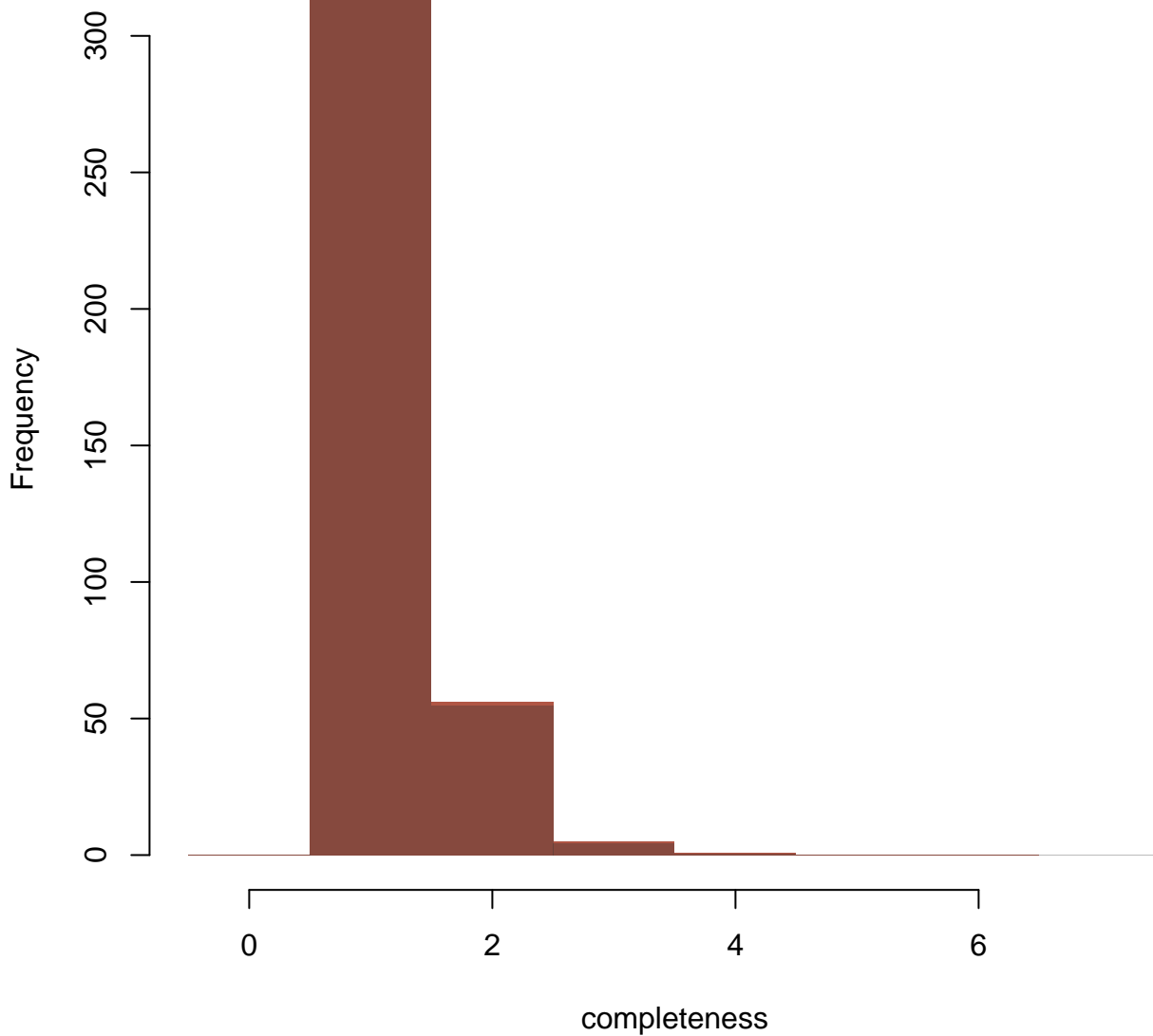
# license Number of terms per tool



## license Top 30 terms

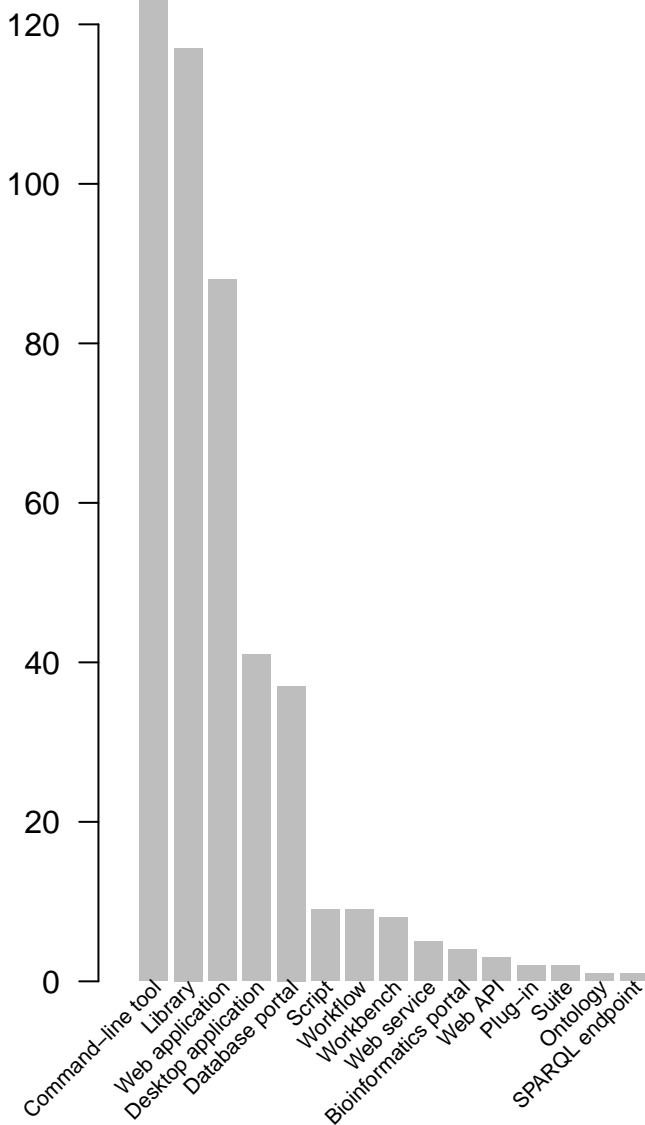


**toolType Number of terms per tool**

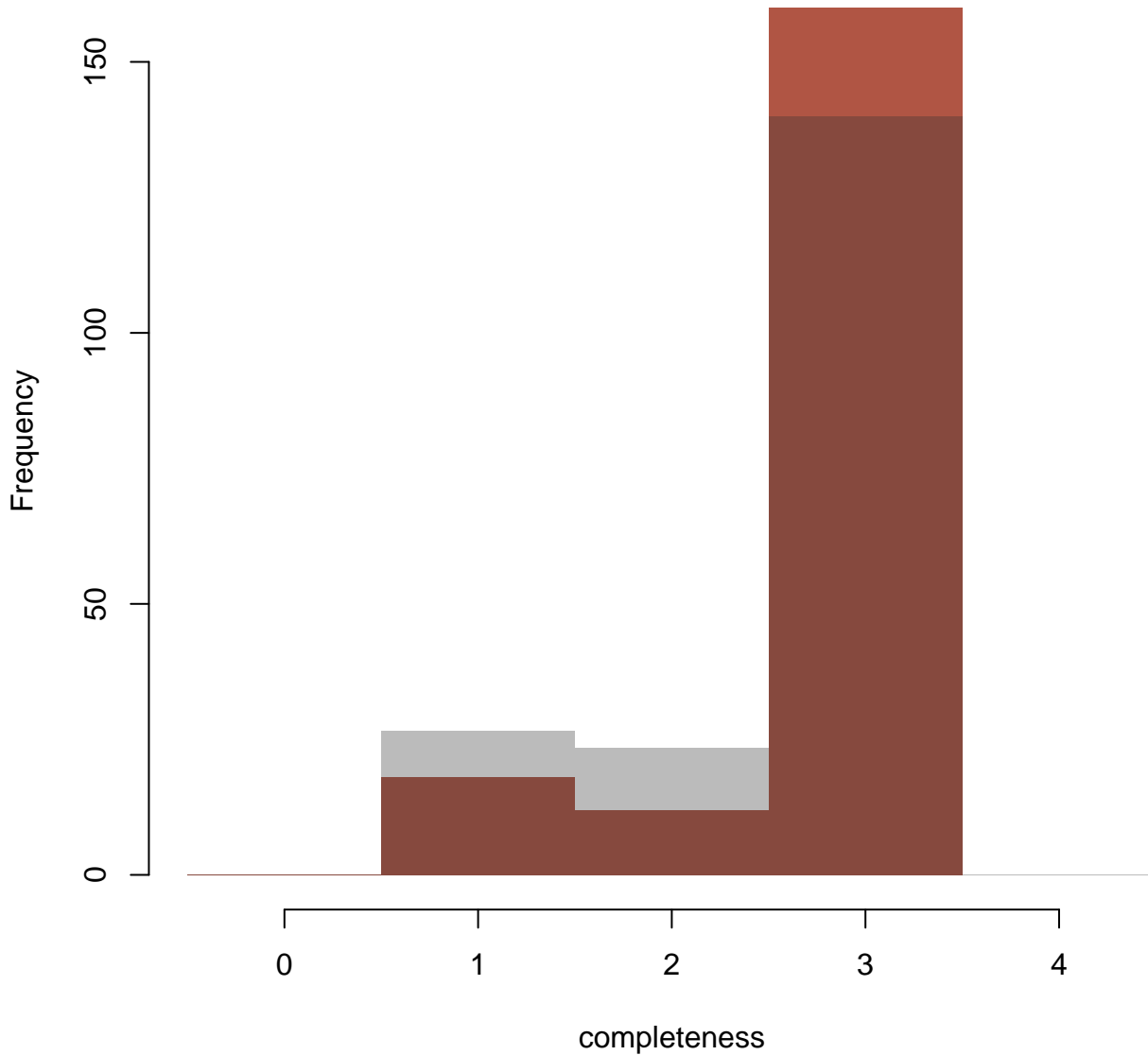




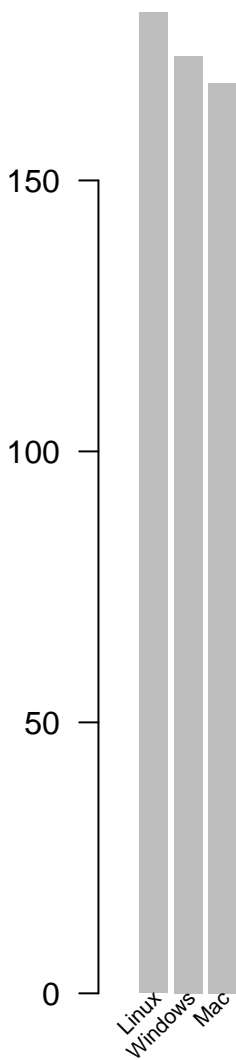
## toolType Top 30 terms



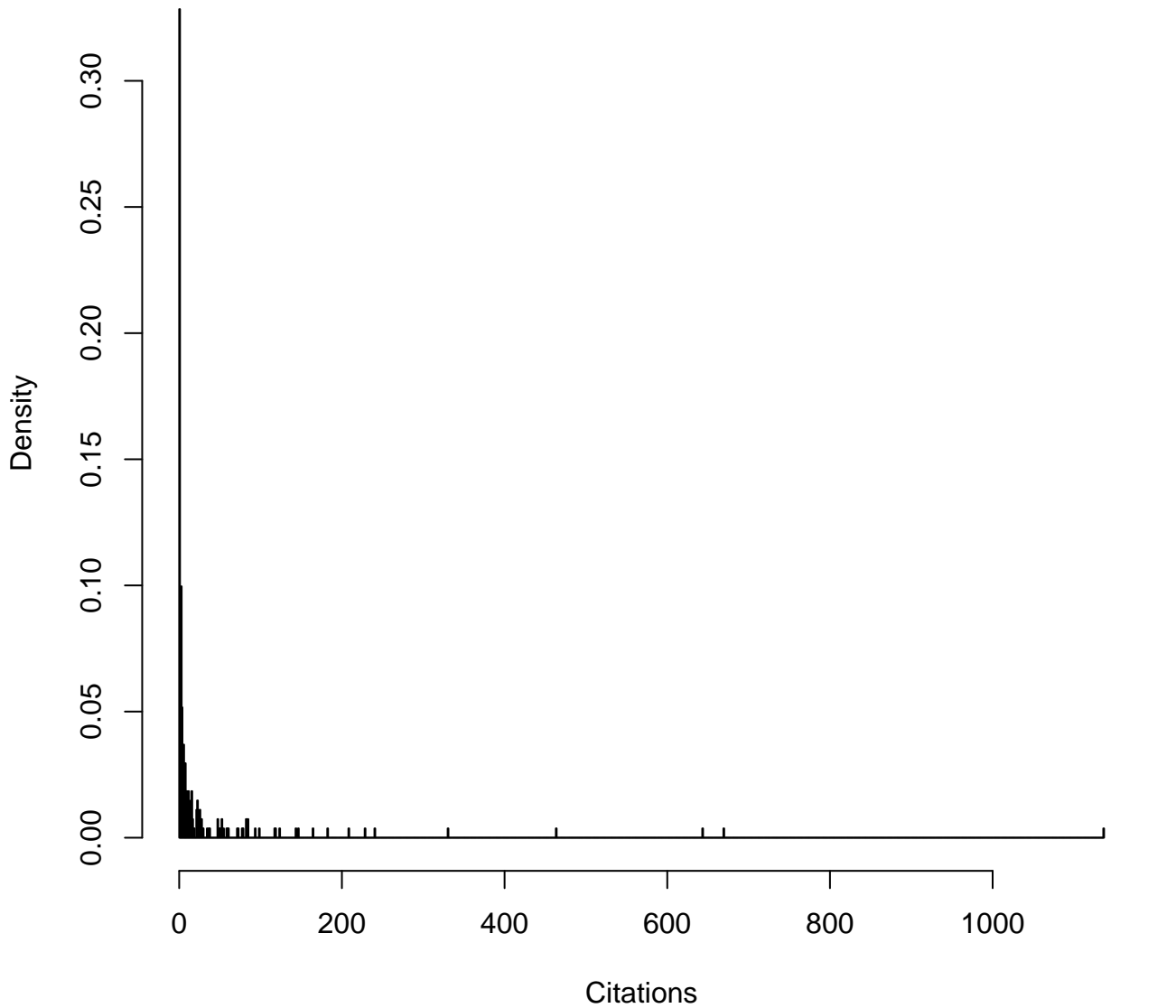
# operatingSystem Number of terms per tool



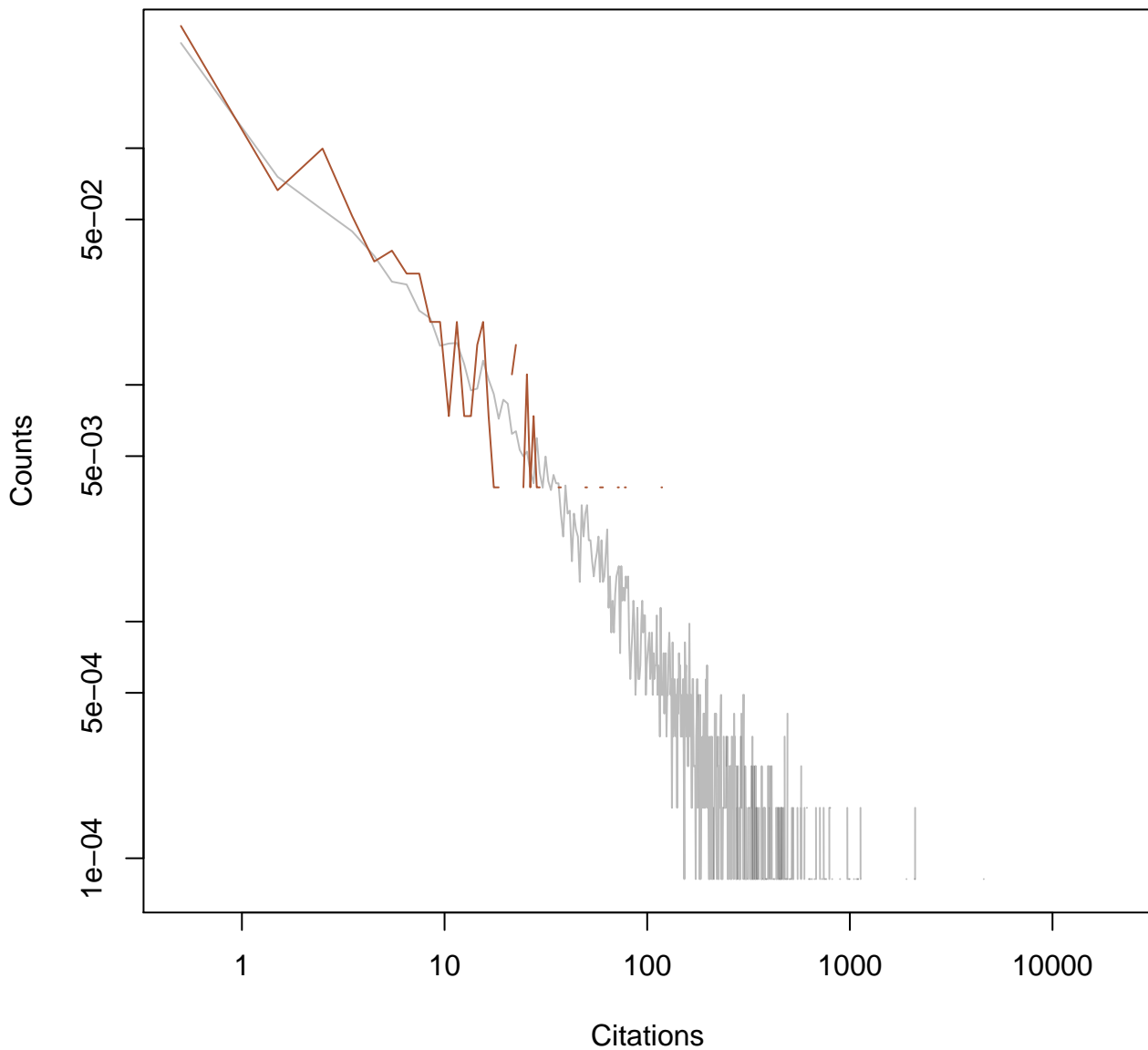
## operatingSystem Top 30 terms



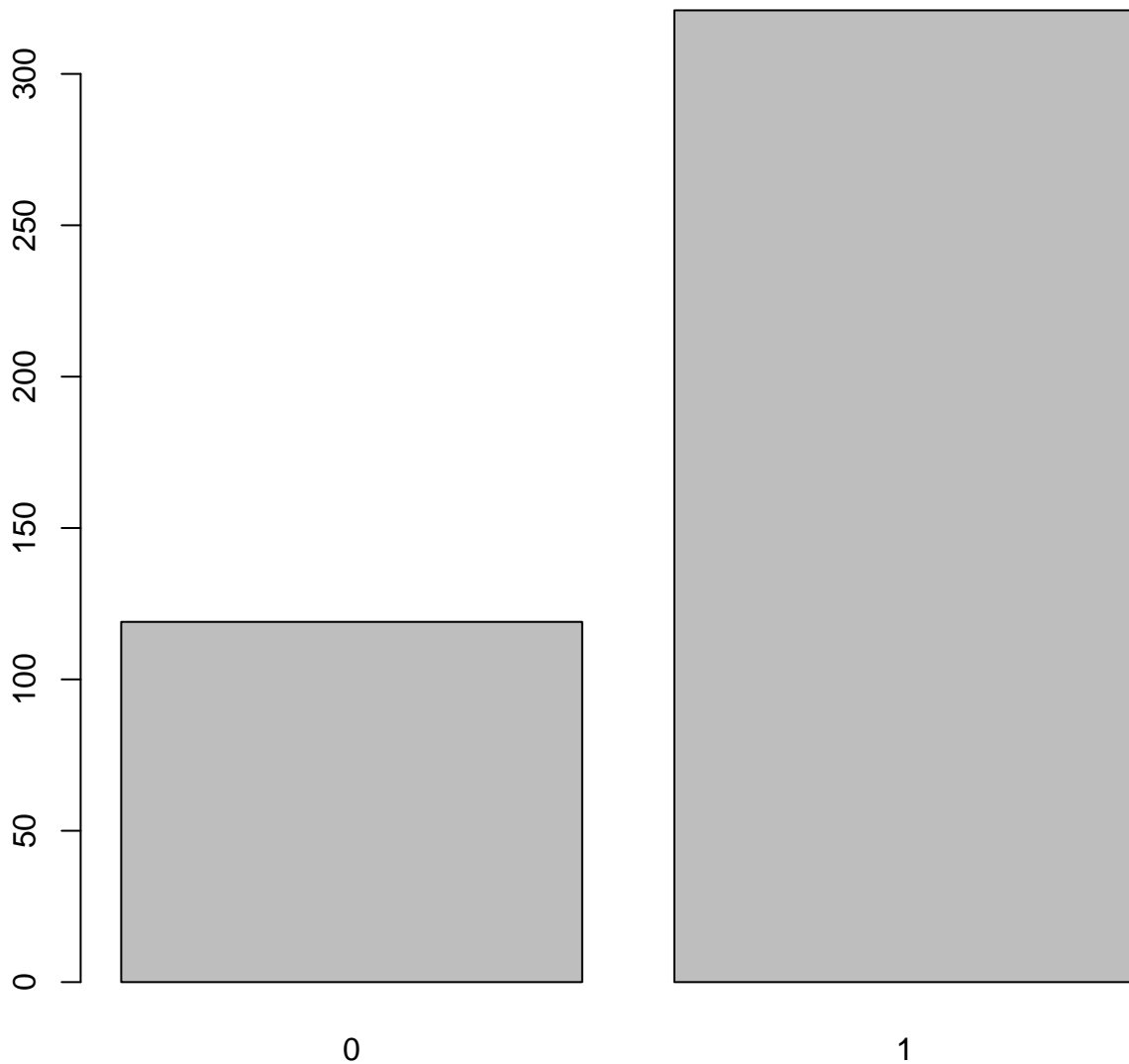
# Citations



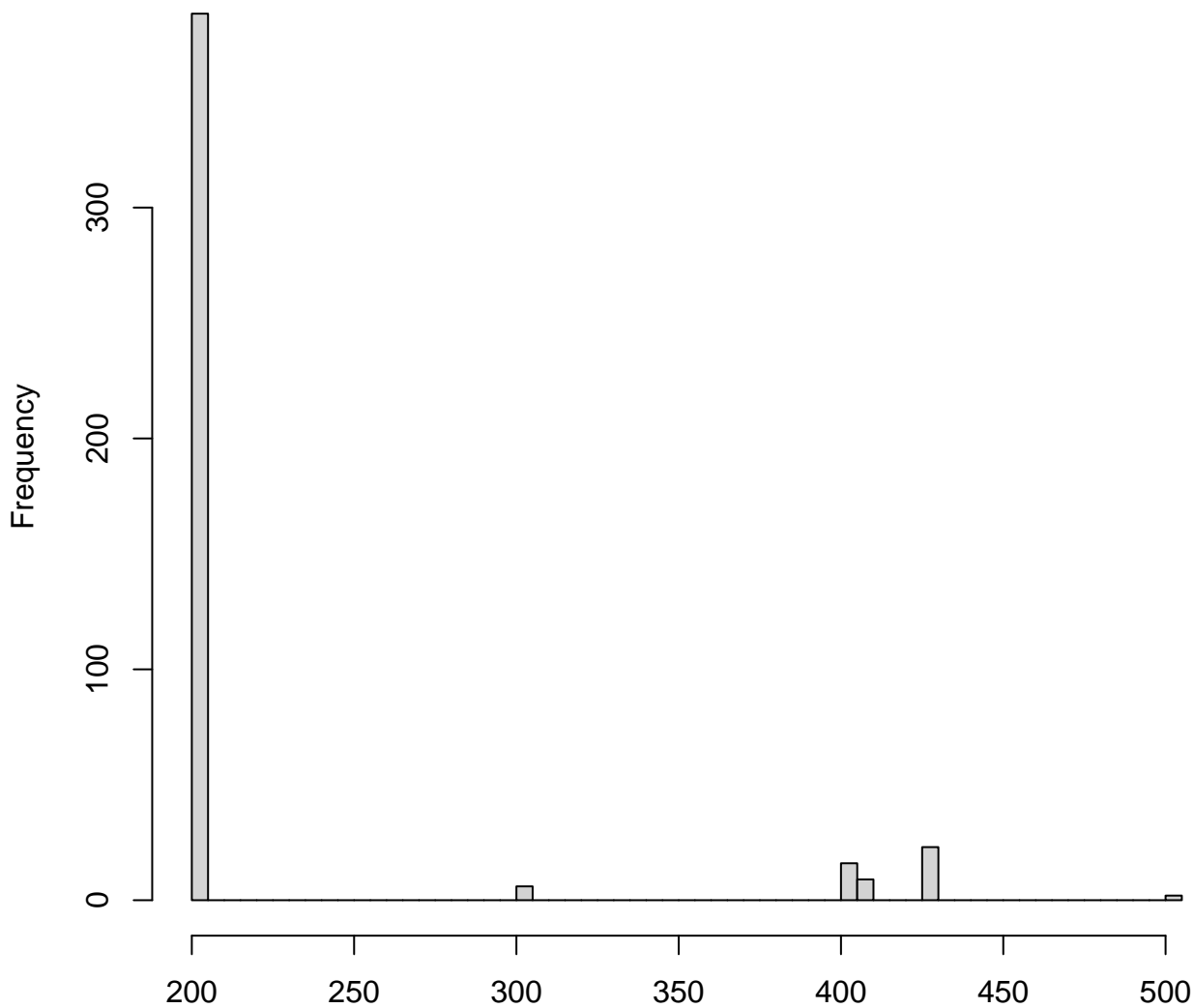
log-log plot of citation count



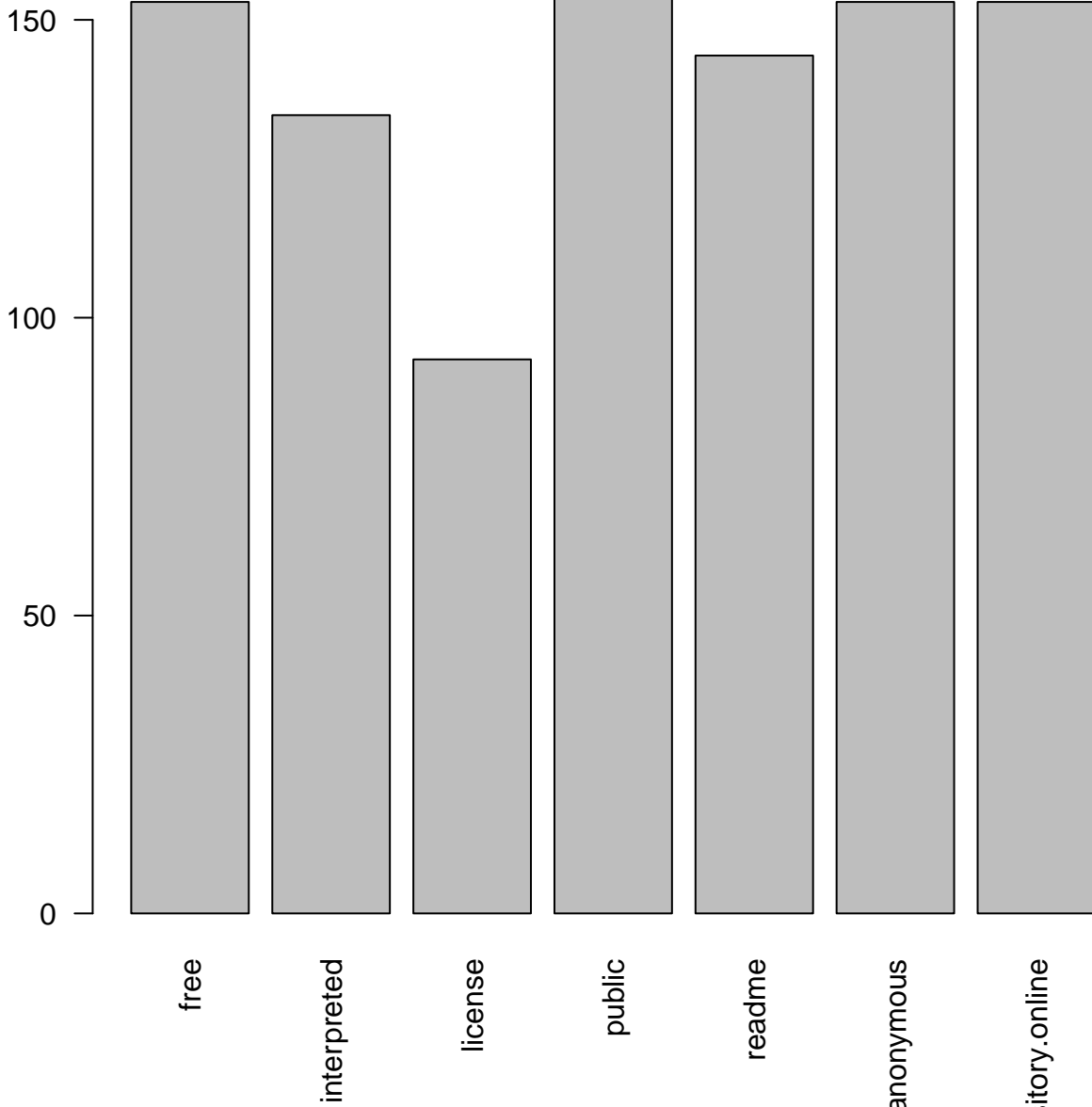
# Is https page



## Operational web page



## Source code





## Issue tracker

60  
50  
40  
30  
20  
10  
0

olic\_tracker