

Results

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#Results

##Review methods

###Flow diagram

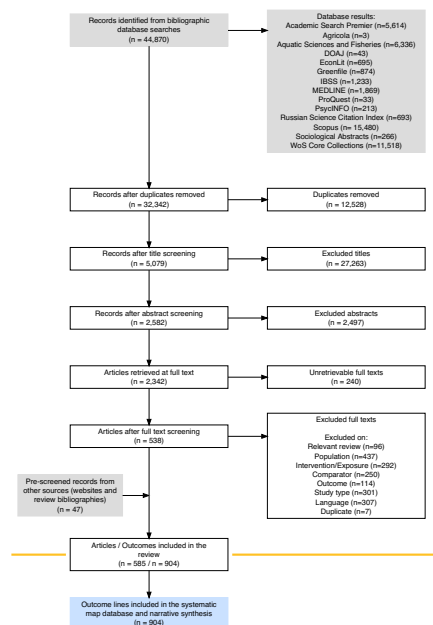


Figure 1: ROSES flow chart for the systematic map, showing the number of records retained at each stage of the review process. Produced using the R package ‘ROSES_flowchart’ (Haddaway 2020).

##Map database

-- Attaching packages ----- tidyverse 1.3.1 --

```
## v ggplot2 3.3.5      v purrr  0.3.4
## v tibble  3.1.6      v dplyr  1.0.8
## v tidyr   1.2.0      v stringr 1.4.0
## v readr   2.1.2      v forcats 0.5.1

## -- Conflicts ----- tidyverse_conflicts() --
## x tidyr::extract()   masks magrittr::extract()
## x dplyr::filter()    masks stats::filter()
## x dplyr::lag()        masks stats::lag()
## x purrr::set_names() masks magrittr::set_names()

##Research interest ###Publications over time
```

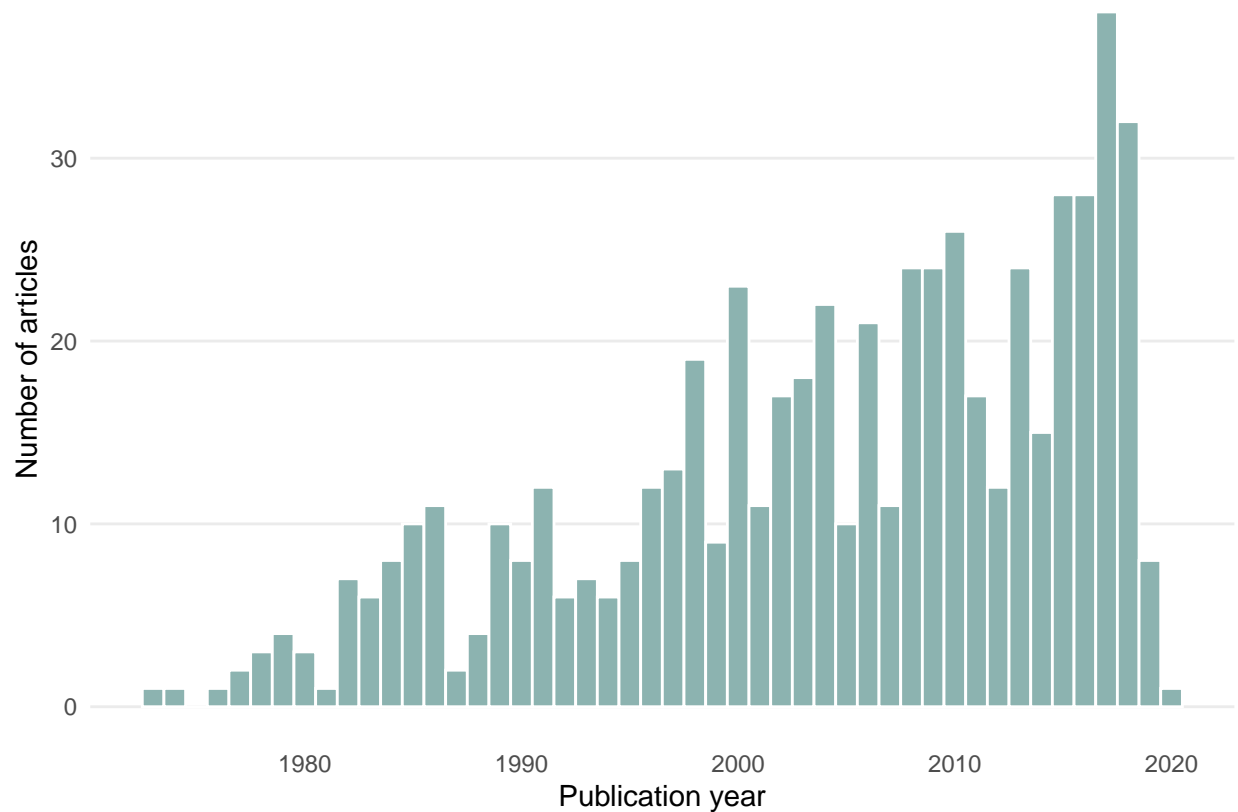


Figure 2: Plot showing the final number of articles included in the systematic map by publication year.

```
###Authors
```

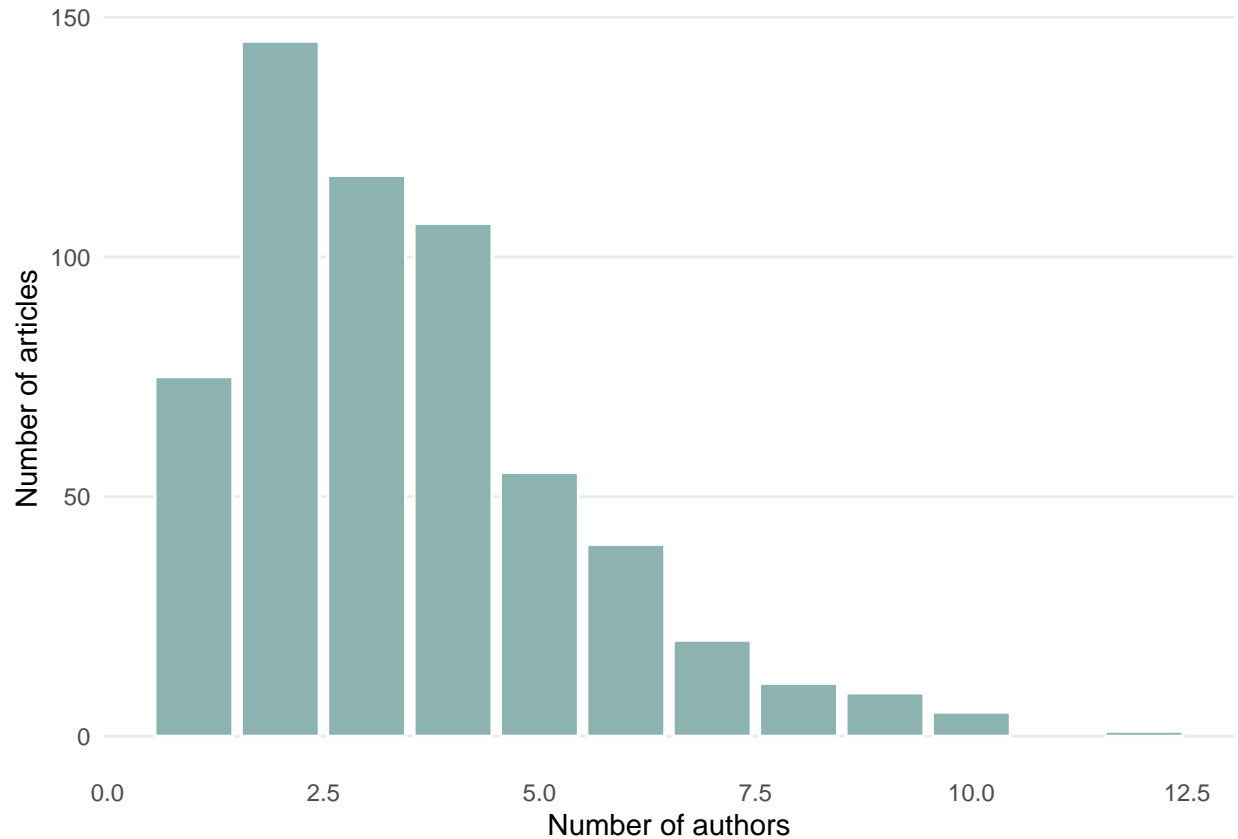


Figure 3: Plot showing the final number of articles included in the systematic map by publication year.

###Evidence atlas [Insert screenshots here]

###Research across mines [mines_articles]

###Study country

##

Attaching package: 'plotly'

The following object is masked from 'package:ggplot2':

##

last_plot

The following object is masked from 'package:stats':

##

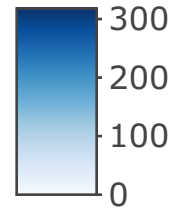
filter

The following object is masked from 'package:graphics':

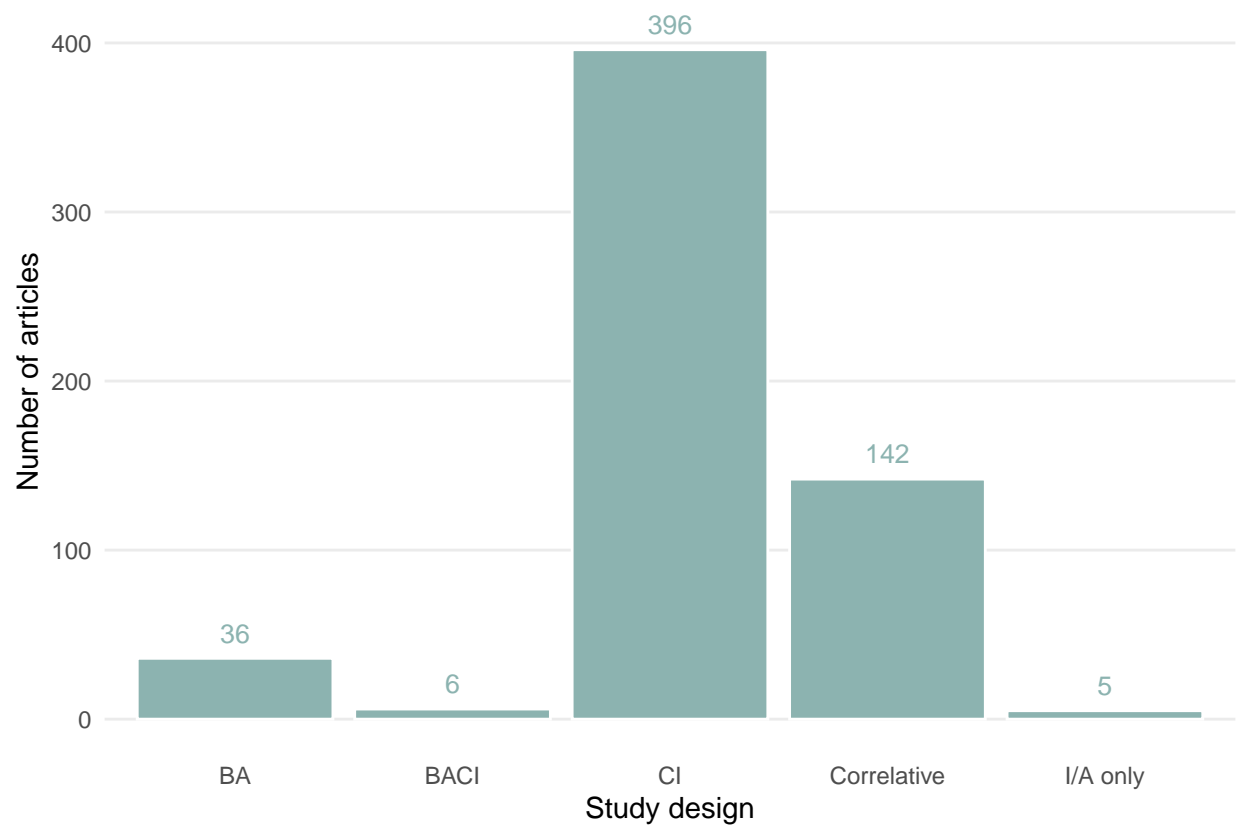
##

layout

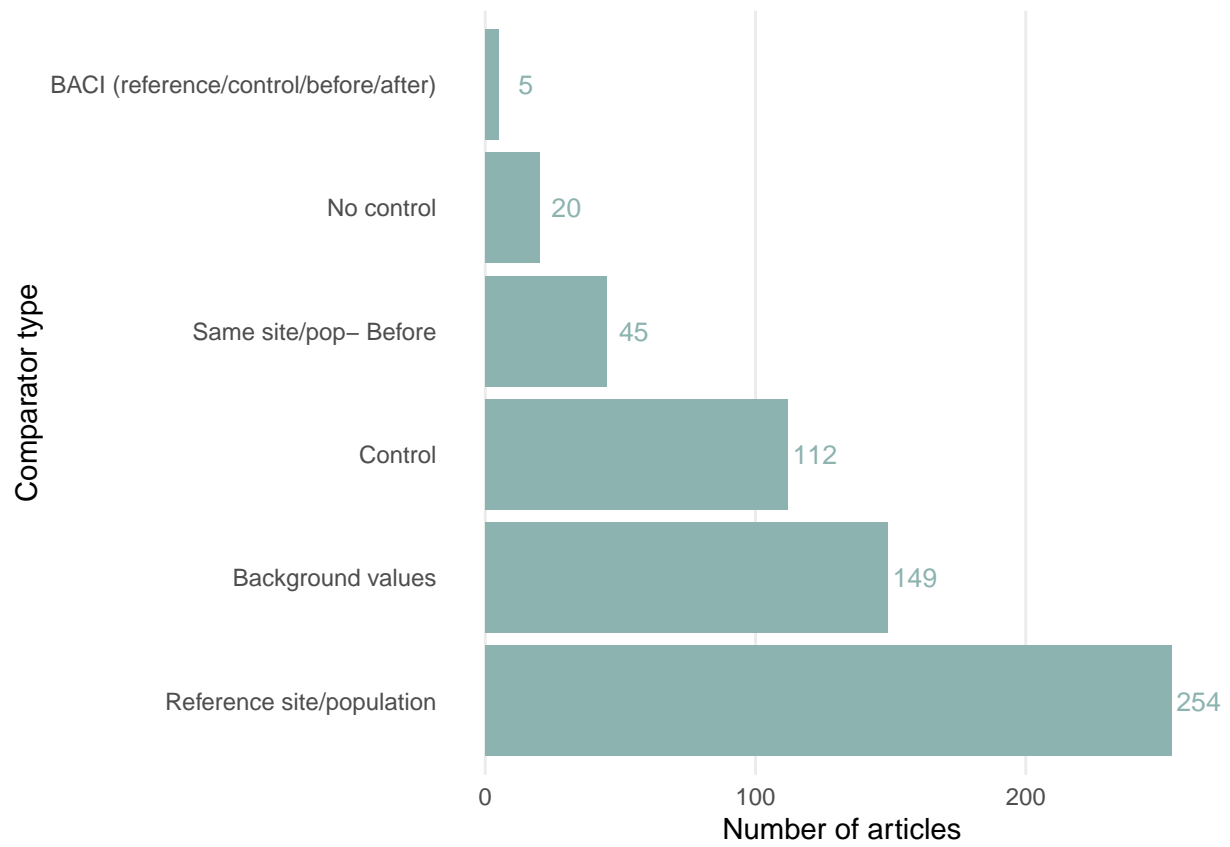
Number of articles
per country



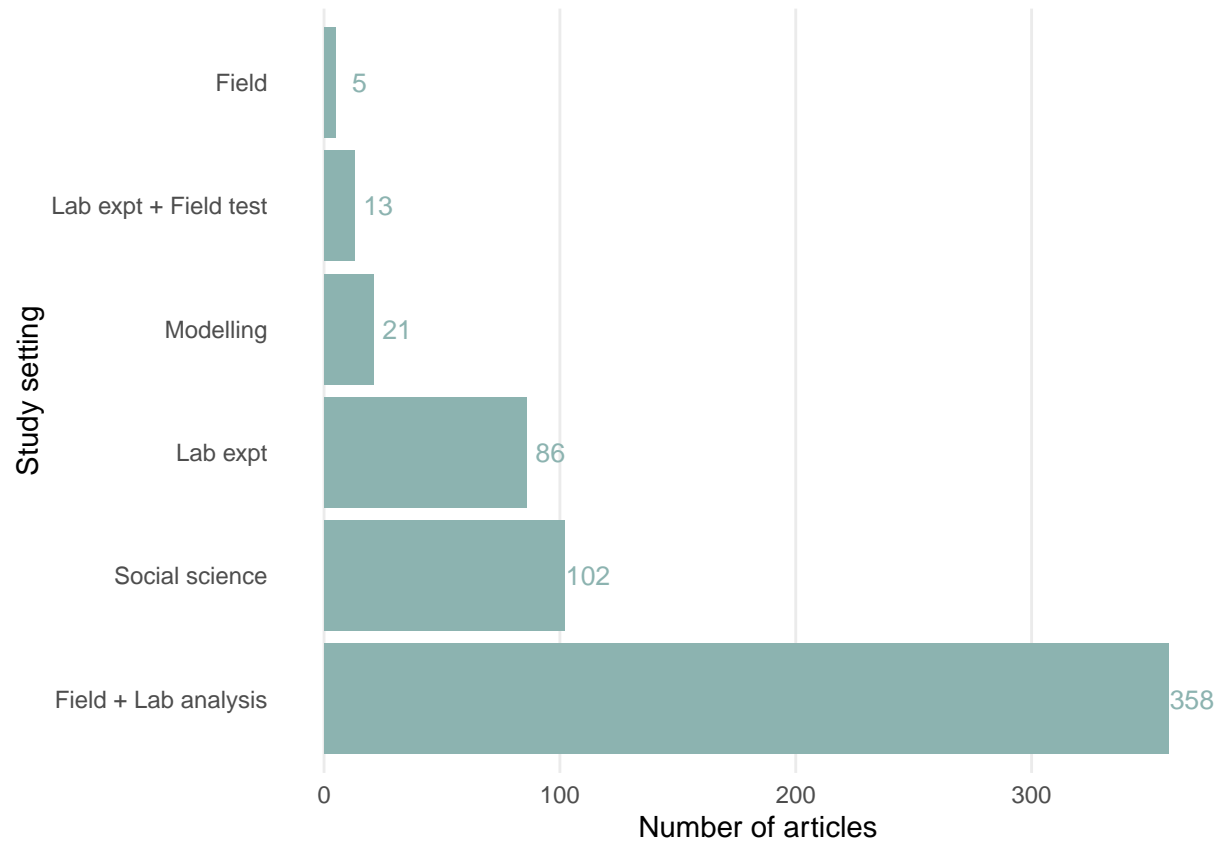
###Study designs



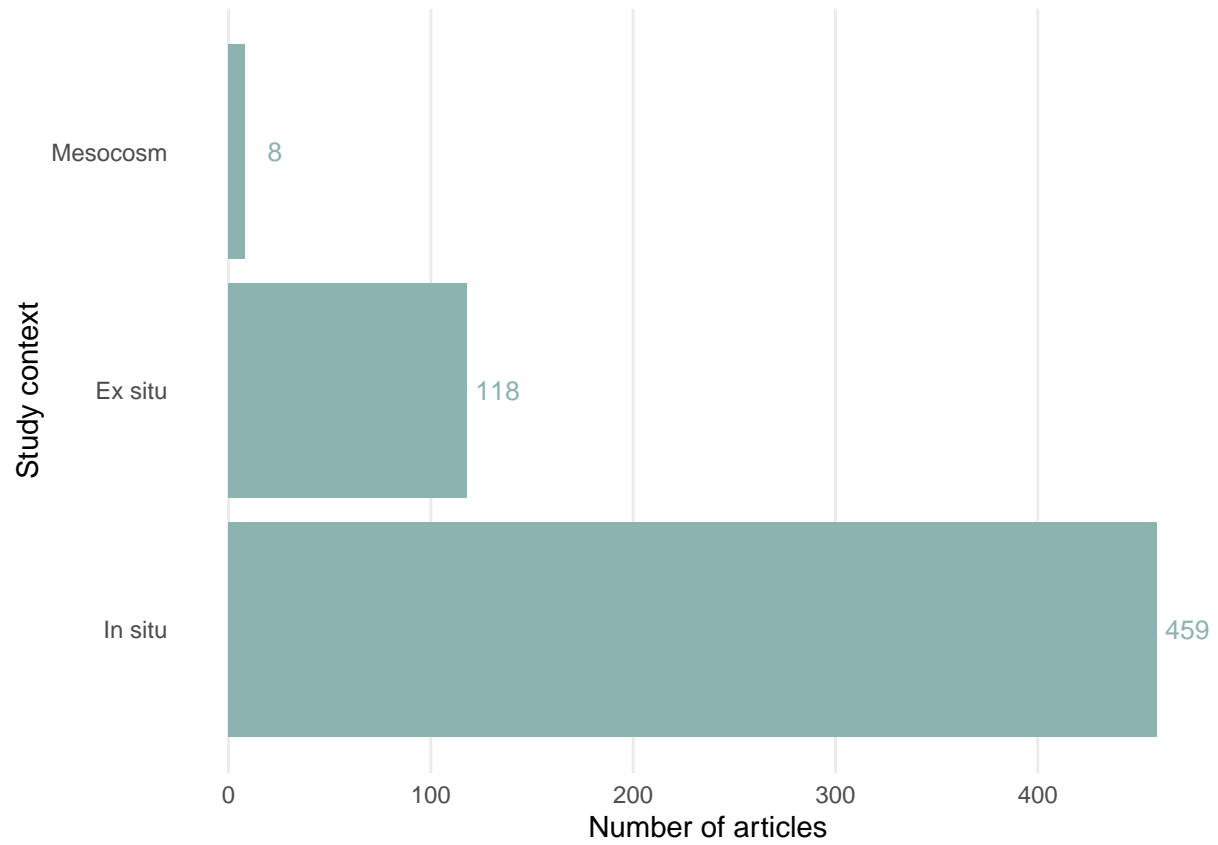
###Comparators



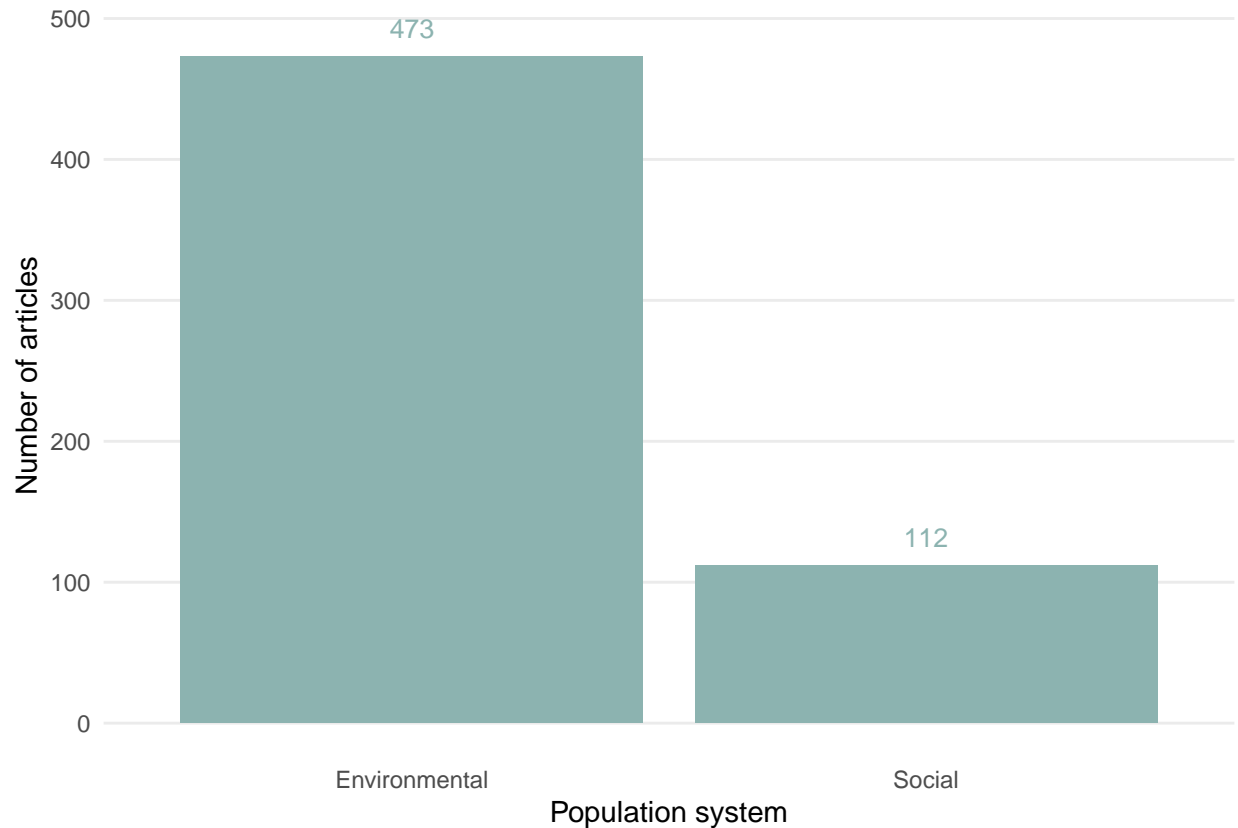
###Settings



###Contexts



###Population systems



Affected systems, components, factors

Warning: Removed 1 rows containing missing values (geom_interactive_point).

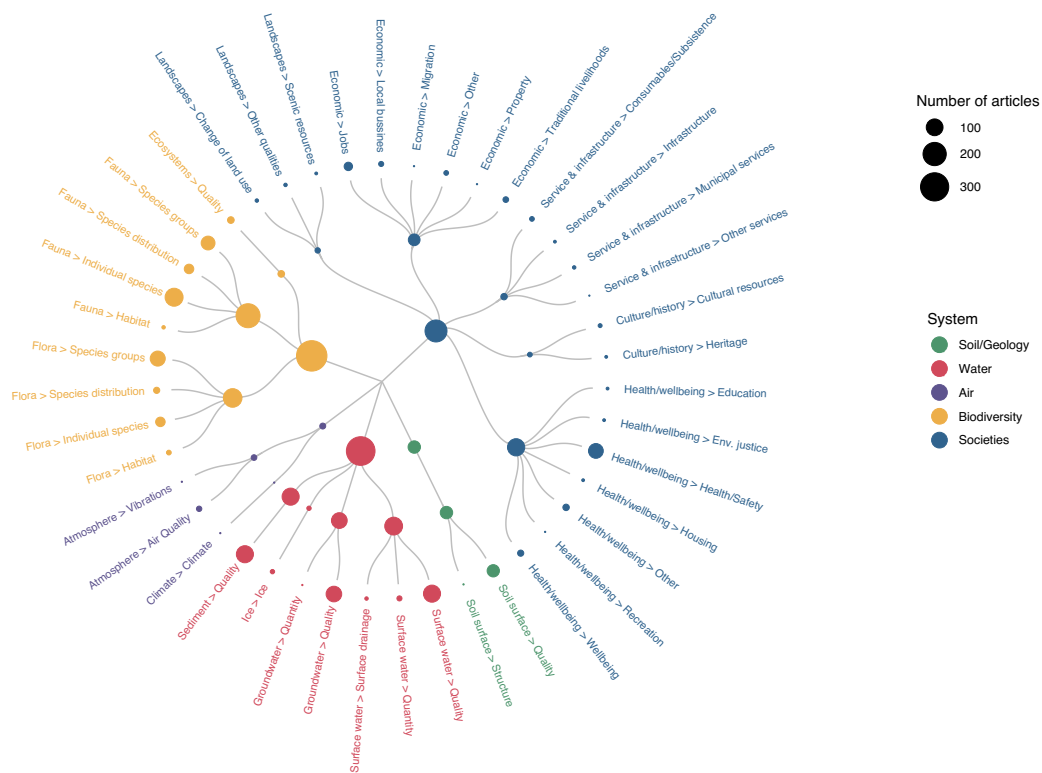
Warning: It is deprecated to specify 'guide = FALSE' to remove a guide. Please use 'guide = "none"' instead.

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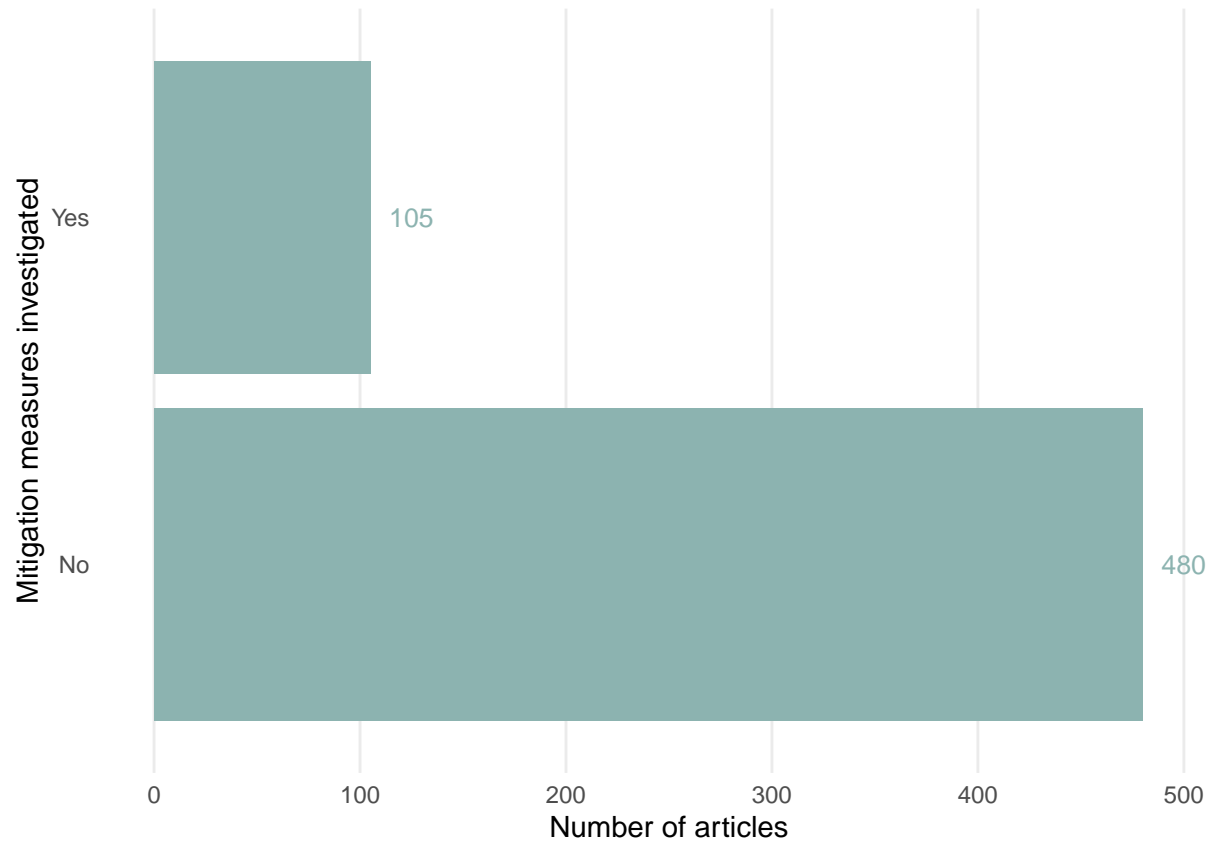
Warning: Removed 1 rows containing missing values (geom_interactive_point).

Warning: It is deprecated to specify 'guide = FALSE' to remove a guide. Please use 'guide = "none"' instead.
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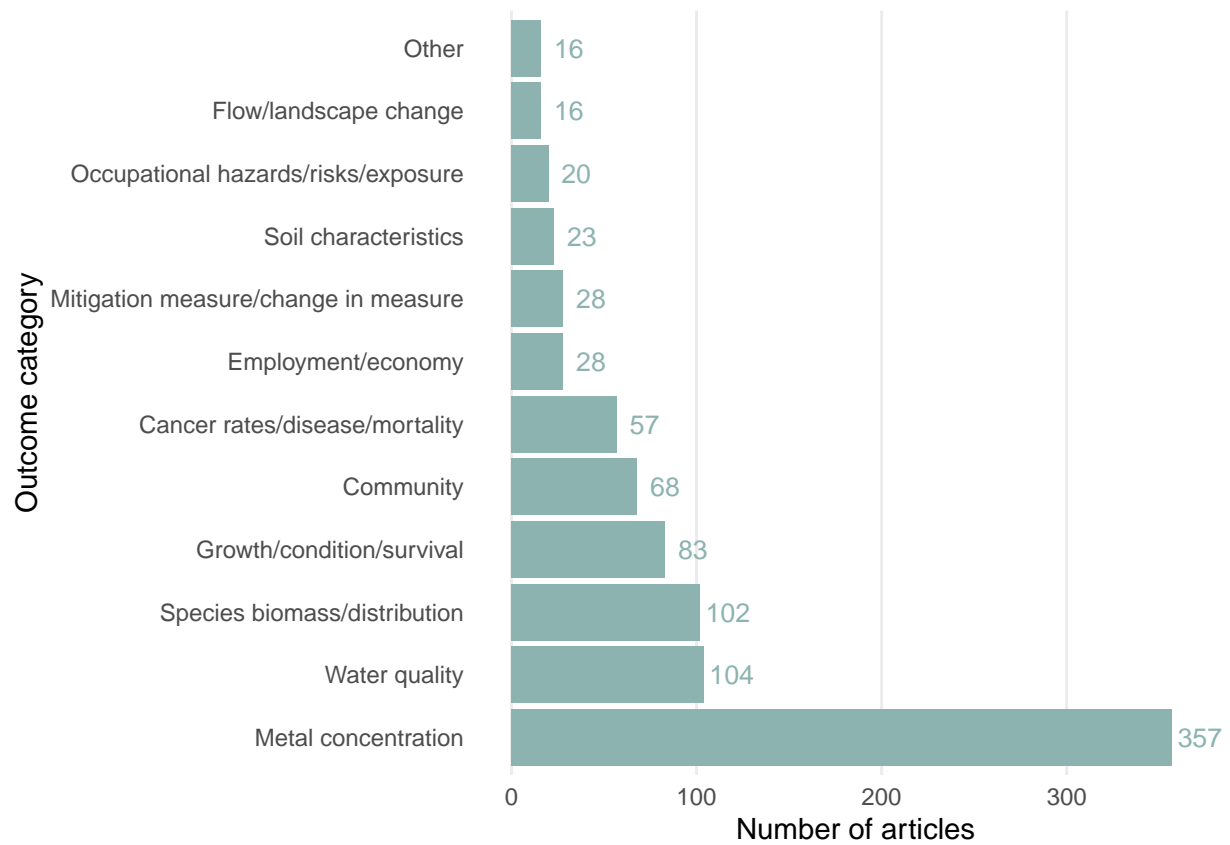


[note - exported interactive_dendrogram.html and output.html (itself using the script in database.R) that each become an iframe within visualisation.html (itself knitted from visualisation.Rhtml). All three need to be updated and hosted for the main site to work]

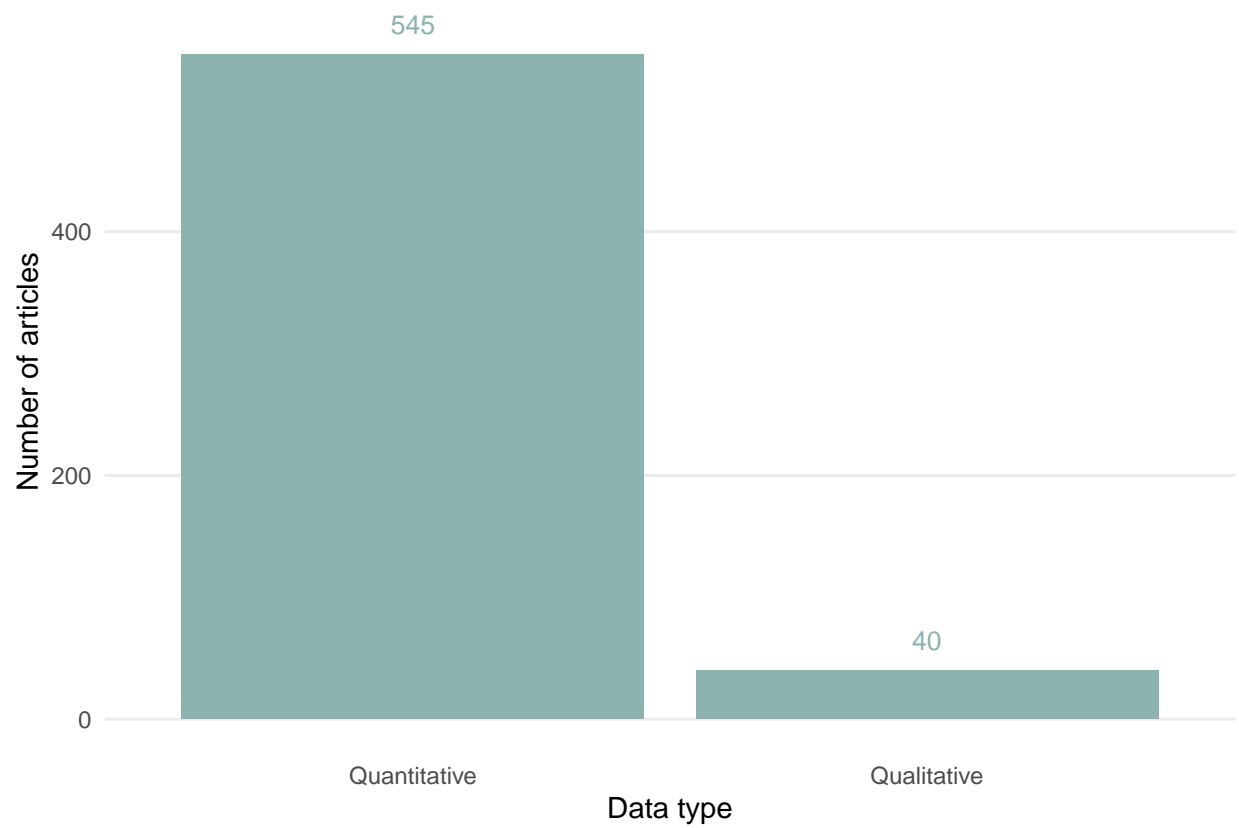
###Mitigation tested



###Outcome category

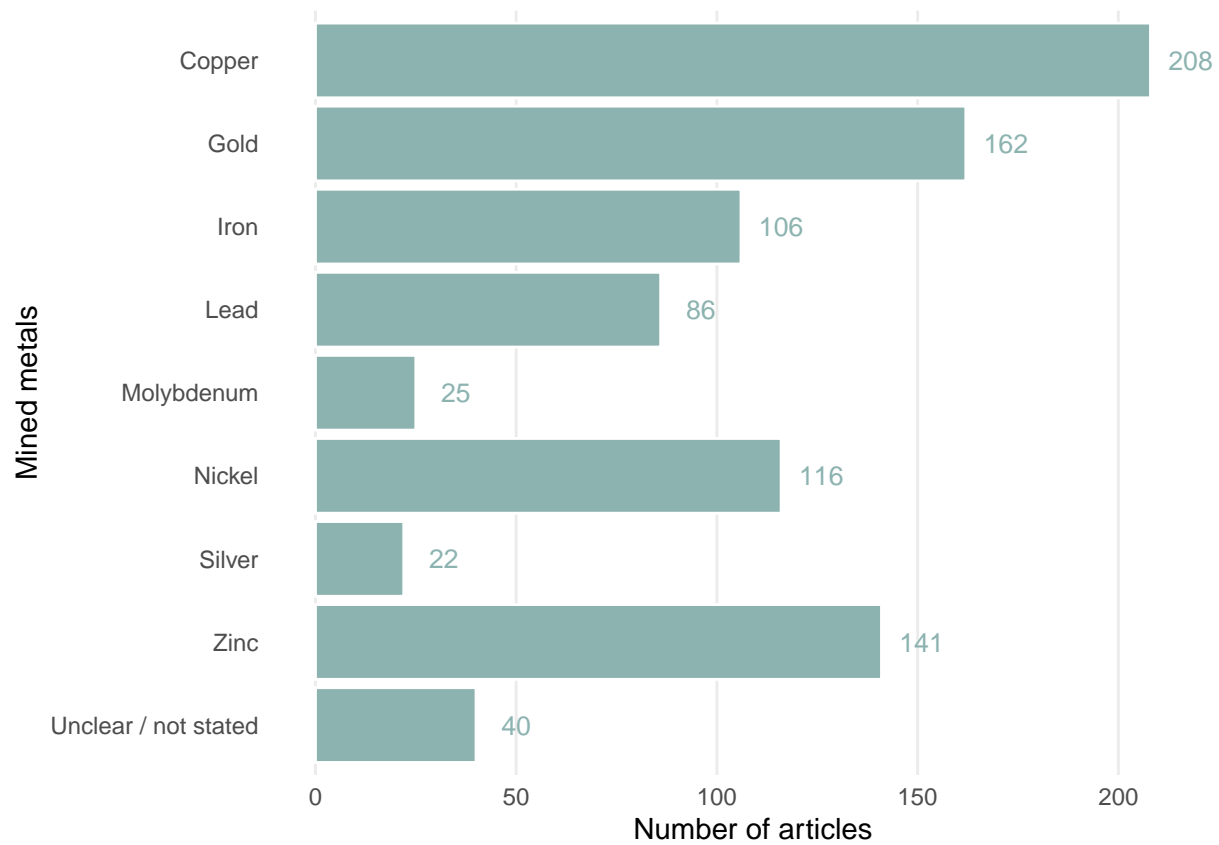


###Data type

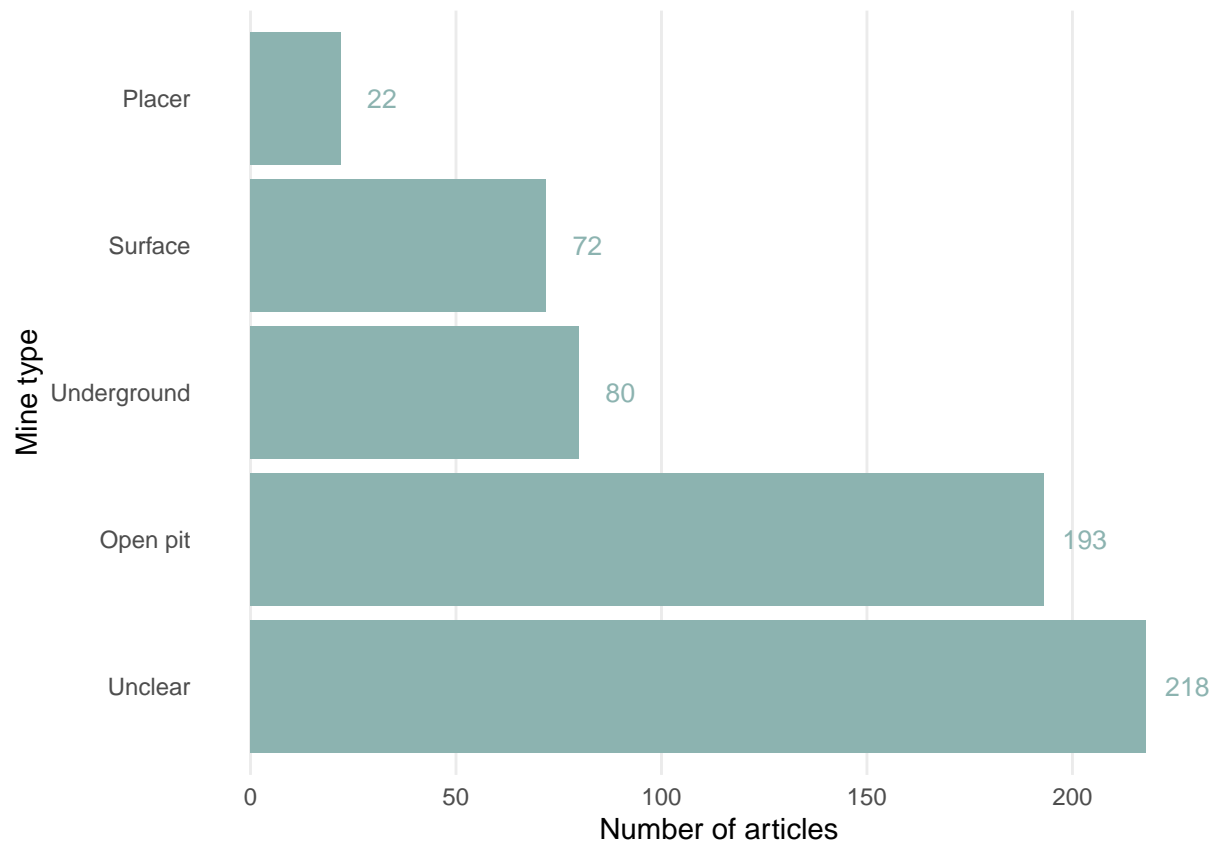


##Mines ###Unique mines

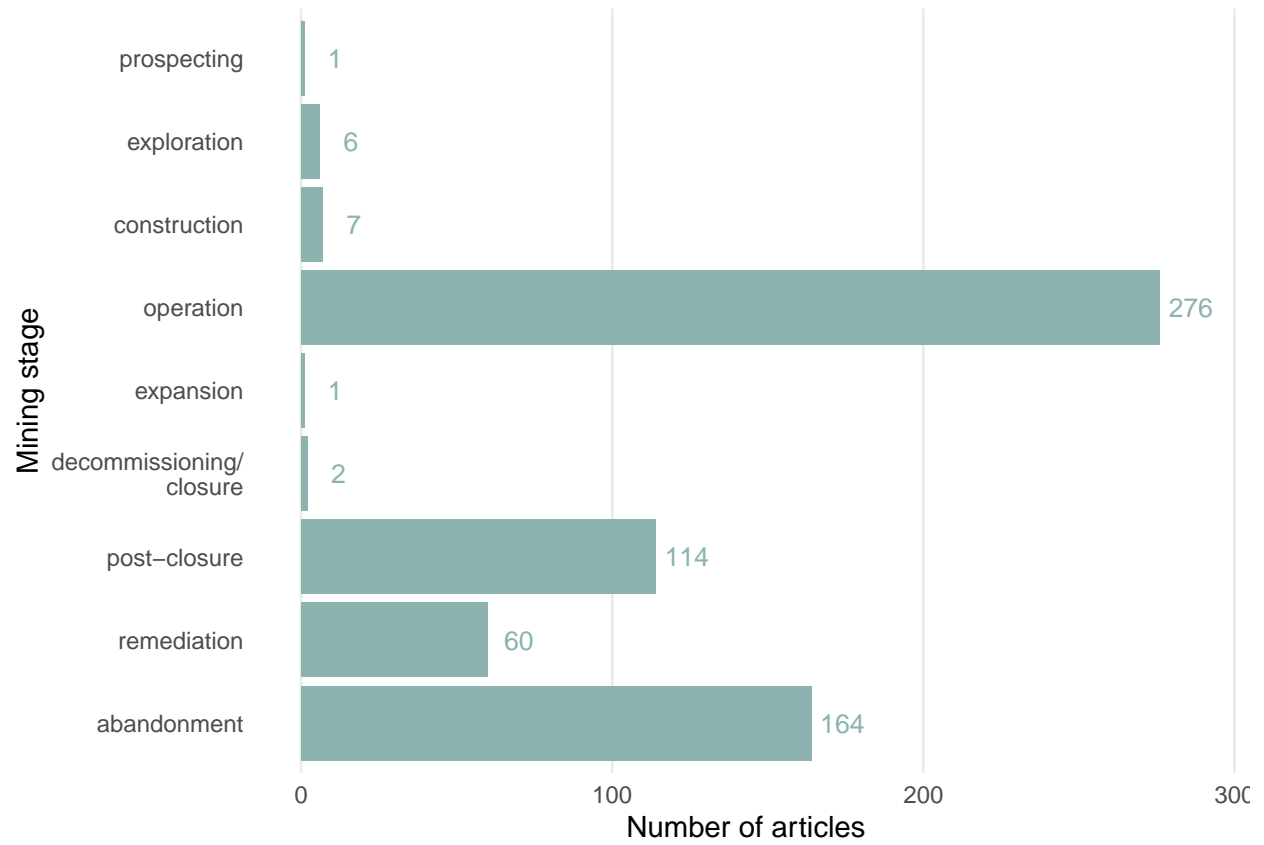
###Metals mined

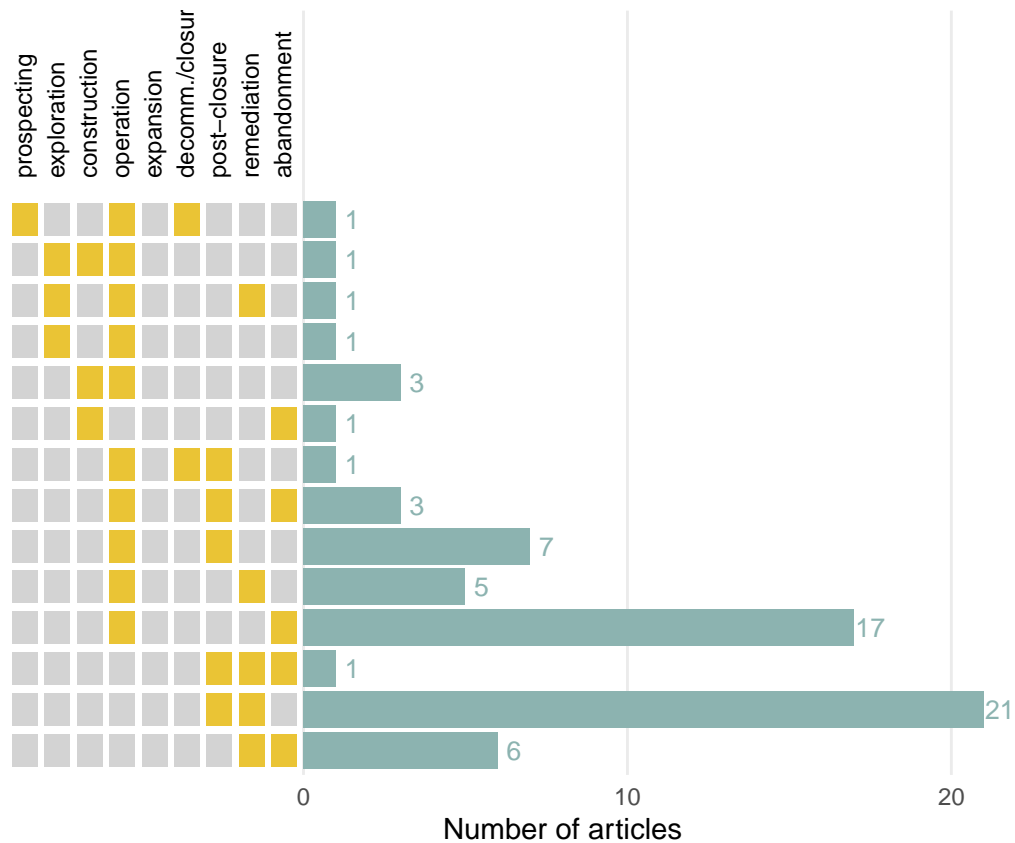


###Mine type



###Extraction stage

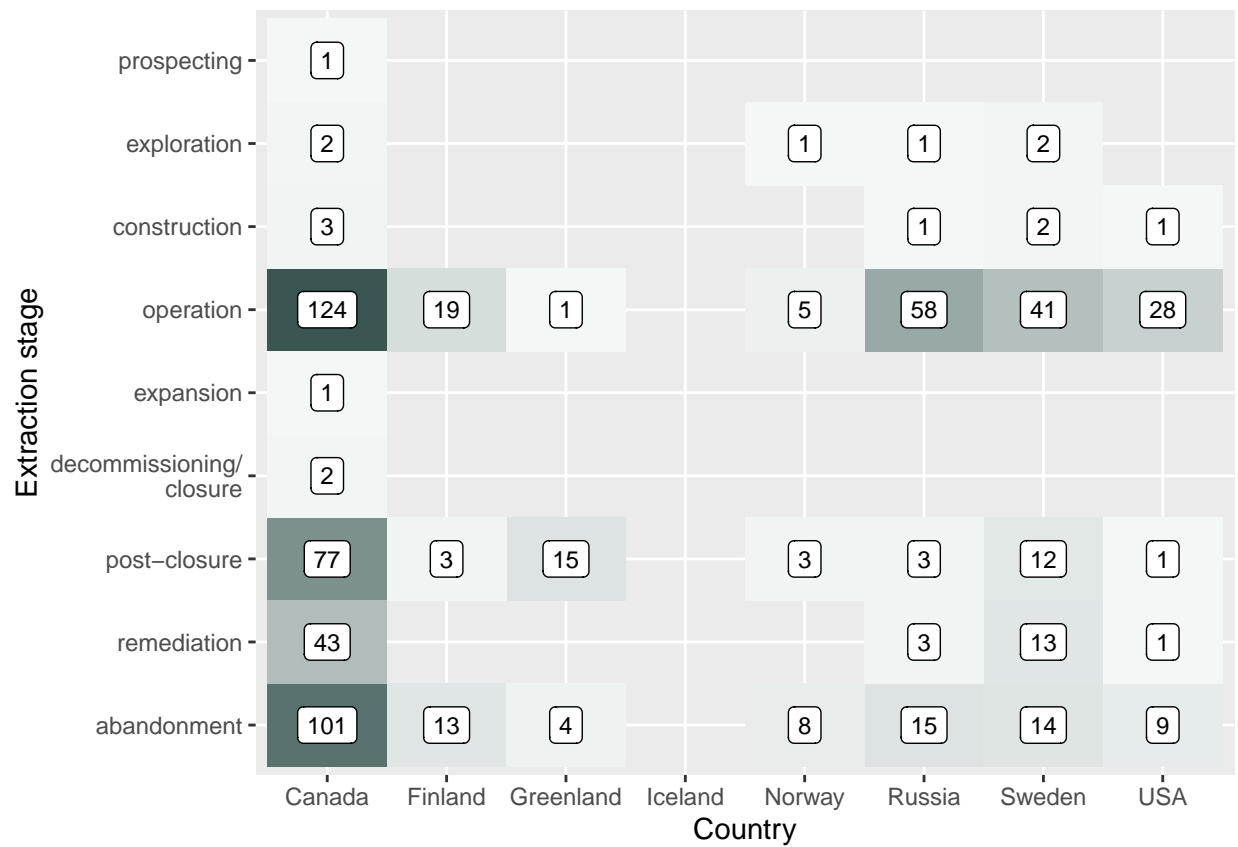




1 Cross Tabs

1.0.0.1 Country -vs- stage

Warning: Removed 9 rows containing missing values (geom_label).



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Gap analysis