

alm tutorial

What are article level metrics?

Glad you asked. The canonical URL for this is perhaps altmetrics.org. Basically it is a metric that measures something about an article. This is in stark contrast to journal level metrics, like the Journal Impact Factor.

Are there other altmetrics data providers?

Yes indeedy.

- [ImpactStory](#)
- [Altmetric.com](#)
- [PlumAnalytics](#)

Authentication

You aren't currently not required to use an API key to access the PLoS ALM API, but soon will need to.

Get your PLoS API key [here](#)

Put your API key in your .Rprofile file using exactly this: `options(PlosApiKey = "YOUalmAPIKEY")`, and the functions within this package will be able to use your API key without you having to enter it every time you run a search.

Install and load

You can get this package by installing via `install_github()` within Hadley Wickham's devtools package.

```
install.packages("devtools")
require(devtools)
install_github("alm", "rOpenSci")

library(alm)
```

The default call with either doi, pmid, pmcid, or mdid without specifying an argument for info

(We'll not print a few columns so the table prints nicely)

```
alm(doi = "10.1371/journal.pone.0029797")[, -c(6:8)]
```

	.id	pdf	html	shares	groups	total
1	bloglines	NA	NA	NA	NA	0
2	citeulike	NA	NA	1	NA	1
3	connotea	NA	NA	NA	NA	0
4	crossref	NA	NA	NA	NA	7
5	nature	NA	NA	NA	NA	4
6	postgenomic	NA	NA	NA	NA	0
7	pubmed	NA	NA	NA	NA	1

8	scopus	NA	NA	NA	NA	4
9	counter	2274	26410	NA	NA	28790
10	researchblogging	NA	NA	NA	NA	1
11	biod	0	0	NA	NA	21
12	pmc	57	334	NA	NA	391
13	facebook	NA	NA	147	NA	228
14	mendeley	NA	NA	50	0	50
15	twitter	NA	NA	NA	NA	9
16	wikipedia	NA	NA	NA	NA	49
17	scienceseeker	NA	NA	NA	NA	0
18	relativemetric	NA	NA	NA	NA	46986
19	f1000	NA	NA	NA	NA	0
20	figshare	NA	NA	NA	NA	0

Details for a single DOI

```
out <- alm(doi = "10.1371/journal.pone.0029797", info = "detail")
## totals
out[["totals"]][, -c(6:8)]
```

	.id	pdf	html	shares	groups	total
1	bloglines	NA	NA	NA	NA	0
2	citeulike	NA	NA	1	NA	1
3	connotea	NA	NA	NA	NA	0
4	crossref	NA	NA	NA	NA	7
5	nature	NA	NA	NA	NA	4
6	postgenomic	NA	NA	NA	NA	0
7	pubmed	NA	NA	NA	NA	1
8	scopus	NA	NA	NA	NA	4
9	counter	2274	26410	NA	NA	28790
10	researchblogging	NA	NA	NA	NA	1
11	biod	0	0	NA	NA	21
12	pmc	57	334	NA	NA	391
13	facebook	NA	NA	147	NA	228
14	mendeley	NA	NA	50	0	50
15	twitter	NA	NA	NA	NA	9
16	wikipedia	NA	NA	NA	NA	49
17	scienceseeker	NA	NA	NA	NA	0
18	relativemetric	NA	NA	NA	NA	46986
19	f1000	NA	NA	NA	NA	0
20	figshare	NA	NA	NA	NA	0

```
## history
head(out[["history"]])
```

	.id	dates	totals
1	citeulike	2013-09-16	1
2	citeulike	2013-08-15	1
3	citeulike	2013-07-13	1
4	citeulike	2013-06-10	1
5	citeulike	2013-05-08	1
6	citeulike	2013-04-05	1

Search using various identifiers, including pubmed id, pmc id, and mendeley id

```
# A single PubMed ID (pmid)
alm(pmid = 22590526)[, -c(6:8)]
```

	.id	pdf	html	shares	groups	total
1	bloglines	NA	NA	NA	NA	0
2	citeulike	NA	NA	5	NA	5
3	connotea	NA	NA	NA	NA	0
4	crossref	NA	NA	NA	NA	3
5	nature	NA	NA	NA	NA	1
6	postgenomic	NA	NA	NA	NA	0
7	pubmed	NA	NA	NA	NA	1
8	scopus	NA	NA	NA	NA	2
9	counter	909	13050	NA	NA	13995
10	researchblogging	NA	NA	NA	NA	1
11	biod	NA	NA	NA	NA	0
12	wos	NA	NA	NA	NA	2
13	pmc	23	113	NA	NA	136
14	facebook	NA	NA	59	NA	161
15	mendeley	NA	NA	39	0	39
16	twitter	NA	NA	NA	NA	118
17	wikipedia	NA	NA	NA	NA	0
18	scienceseeker	NA	NA	NA	NA	1
19	relativemetric	NA	NA	NA	NA	27039
20	f1000	NA	NA	NA	NA	0
21	figshare	NA	NA	NA	NA	0

```
# A single PubMed Central ID (pmcid)
alm(pmcid = 212692)[, -c(6:8)]
```

	.id	pdf	html	shares	groups	total
1	bloglines	NA	NA	NA	NA	0
2	citeulike	NA	NA	8	NA	8
3	connotea	NA	NA	NA	NA	0
4	crossref	NA	NA	NA	NA	143
5	nature	NA	NA	NA	NA	0
6	postgenomic	NA	NA	NA	NA	0
7	pubmed	NA	NA	NA	NA	143
8	scopus	NA	NA	NA	NA	318
9	counter	2447	19698	NA	NA	22292
10	researchblogging	NA	NA	NA	NA	0
11	biod	NA	NA	NA	NA	0
12	pmc	2305	4752	NA	NA	7057
13	facebook	NA	NA	0	NA	0
14	mendeley	NA	NA	71	0	71
15	twitter	NA	NA	NA	NA	0
16	wikipedia	NA	NA	NA	NA	0
17	scienceseeker	NA	NA	NA	NA	0
18	relativemetric	NA	NA	NA	NA	2192654
19	f1000	NA	NA	NA	NA	0
20	figshare	1	2	NA	NA	3

```
# A single Mendeley UUID (mdid)
alm(mdid = "35791700-6d00-11df-a2b2-0026b95e3eb7"), -c(6:8)]
```

	.id	pdf	html	shares	groups	total
1	bloglines	NA	NA	NA	NA	0
2	citeulike	NA	NA	8	NA	8
3	connotea	NA	NA	NA	NA	0
4	crossref	NA	NA	NA	NA	143
5	nature	NA	NA	NA	NA	0
6	postgenomic	NA	NA	NA	NA	0
7	pubmed	NA	NA	NA	NA	143
8	scopus	NA	NA	NA	NA	318
9	counter	2447	19698	NA	NA	22292
10	researchblogging	NA	NA	NA	NA	0
11	biod	NA	NA	NA	NA	0
12	pmc	2305	4752	NA	NA	7057
13	facebook	NA	NA	0	NA	0
14	mendeley	NA	NA	71	0	71
15	twitter	NA	NA	NA	NA	0
16	wikipedia	NA	NA	NA	NA	0
17	scienceseeker	NA	NA	NA	NA	0
18	relativemetric	NA	NA	NA	NA	2192654
19	f1000	NA	NA	NA	NA	0
20	figshare	1	2	NA	NA	3

Search on many identifiers

```
dois <- c("10.1371/journal.pone.0001543", "10.1371/journal.pone.0040117", "10.1371/journal.pone.0029797",
"10.1371/journal.pone.0039395")
out <- alm(doi = dois)
lapply(out, head)
```

```
[[1]]
```

	.id	pdf	html	shares	groups	comments	likes	citations	total
1	bloglines	NA	NA	NA	NA	NA	NA	0	0
2	citeulike	NA	NA	0	NA	NA	NA	NA	0
3	connotea	NA	NA	NA	NA	NA	NA	0	0
4	crossref	NA	NA	NA	NA	NA	NA	1	1
5	nature	NA	NA	NA	NA	NA	NA	0	0
6	postgenomic	NA	NA	NA	NA	NA	NA	0	0

```
[[2]]
```

	.id	pdf	html	shares	groups	comments	likes	citations	total
1	bloglines	NA	NA	NA	NA	NA	NA	0	0
2	citeulike	NA	NA	0	NA	NA	NA	NA	0
3	connotea	NA	NA	NA	NA	NA	NA	0	0
4	crossref	NA	NA	NA	NA	NA	NA	0	0
5	nature	NA	NA	NA	NA	NA	NA	0	0
6	postgenomic	NA	NA	NA	NA	NA	NA	0	0

```
[[3]]
```

	.id	pdf	html	shares	groups	comments	likes	citations	total
--	-----	-----	------	--------	--------	----------	-------	-----------	-------

1	bloglines	NA	NA	NA	NA	NA	NA	0	0
2	citeulike	NA	NA	1	NA	NA	NA	NA	1
3	connotea	NA	NA	NA	NA	NA	NA	0	0
4	crossref	NA	NA	NA	NA	NA	NA	7	7
5	nature	NA	NA	NA	NA	NA	NA	4	4
6	postgenomic	NA	NA	NA	NA	NA	NA	0	0

```
[[4]]
```

	.id	pdf	html	shares	groups	comments	likes	citations	total
1	bloglines	NA	NA	NA	NA	NA	NA	0	0
2	citeulike	NA	NA	0	NA	NA	NA	NA	0
3	connotea	NA	NA	NA	NA	NA	NA	0	0
4	crossref	NA	NA	NA	NA	NA	NA	5	5
5	nature	NA	NA	NA	NA	NA	NA	0	0
6	postgenomic	NA	NA	NA	NA	NA	NA	1	1

Get altmetrics by year

You can also get metrics by day (`sum_metrics='day'`) or month (`sum_metrics='month'`)

```
alm(doi = "10.1371/journal.pone.0036240", sum_metrics = "year")[, -c(6:8)]
```

	.id	x	year	pdf	html	likes	citations	total
1	bloglines	NA	NA	NA	NA	NA	NA	NA
2	citeulike	NA	2012	NA	NA	NA	NA	5
3	connotea	NA	NA	NA	NA	NA	NA	NA
4	crossref	NA	2013	NA	NA	NA	3	3
5	nature	NA	NA	NA	NA	NA	NA	NA
6	postgenomic	NA	NA	NA	NA	NA	NA	NA
7	pubmed	NA	NA	NA	NA	NA	NA	NA
8	scopus	NA	NA	NA	NA	NA	NA	NA
9	counter	NA	2012	699	10502	NA	NA	11234
10	counter	NA	2013	210	2548	NA	NA	2761
11	researchblogging	NA	2013	NA	NA	NA	1	1
12	biod	NA	NA	NA	NA	NA	NA	NA
13	pmc	NA	2012	16	53	NA	NA	69
14	pmc	NA	2013	7	60	NA	NA	67
15	facebook	NA	NA	NA	NA	NA	NA	NA
16	mendeley	NA	NA	NA	NA	NA	NA	NA
17	twitter	NA	2012	NA	NA	NA	NA	103
18	twitter	NA	2013	NA	NA	NA	NA	15
19	wikipedia	NA	NA	NA	NA	NA	NA	NA
20	scienceseeker	NA	2013	NA	NA	NA	1	1
21	relativemetric	NA	NA	NA	NA	NA	NA	NA
22	f1000	NA	NA	NA	NA	NA	NA	NA
23	figshare	NA	NA	NA	NA	NA	NA	NA

Output an-easy-to-combine-with-other-results dataframe

```
alm(doi = "10.1371/journal.pone.0035869", total_details = TRUE)[, 3:10]
```

```
publication_date bloglines_pdf bloglines_html bloglines_shares
```

```

1 2012-05-11T07:00:00Z      NA      NA      NA
  bloglines_groups bloglines_comments bloglines_likes bloglines_citations
1              NA              NA              NA              0

```

Get altmetrics data for a single paper, and visualize the total data across dates

```

out <- alm(doi = "10.1371/journal.pone.0001543", info = "detail")
almplot(out, type = "totalmetrics")

```

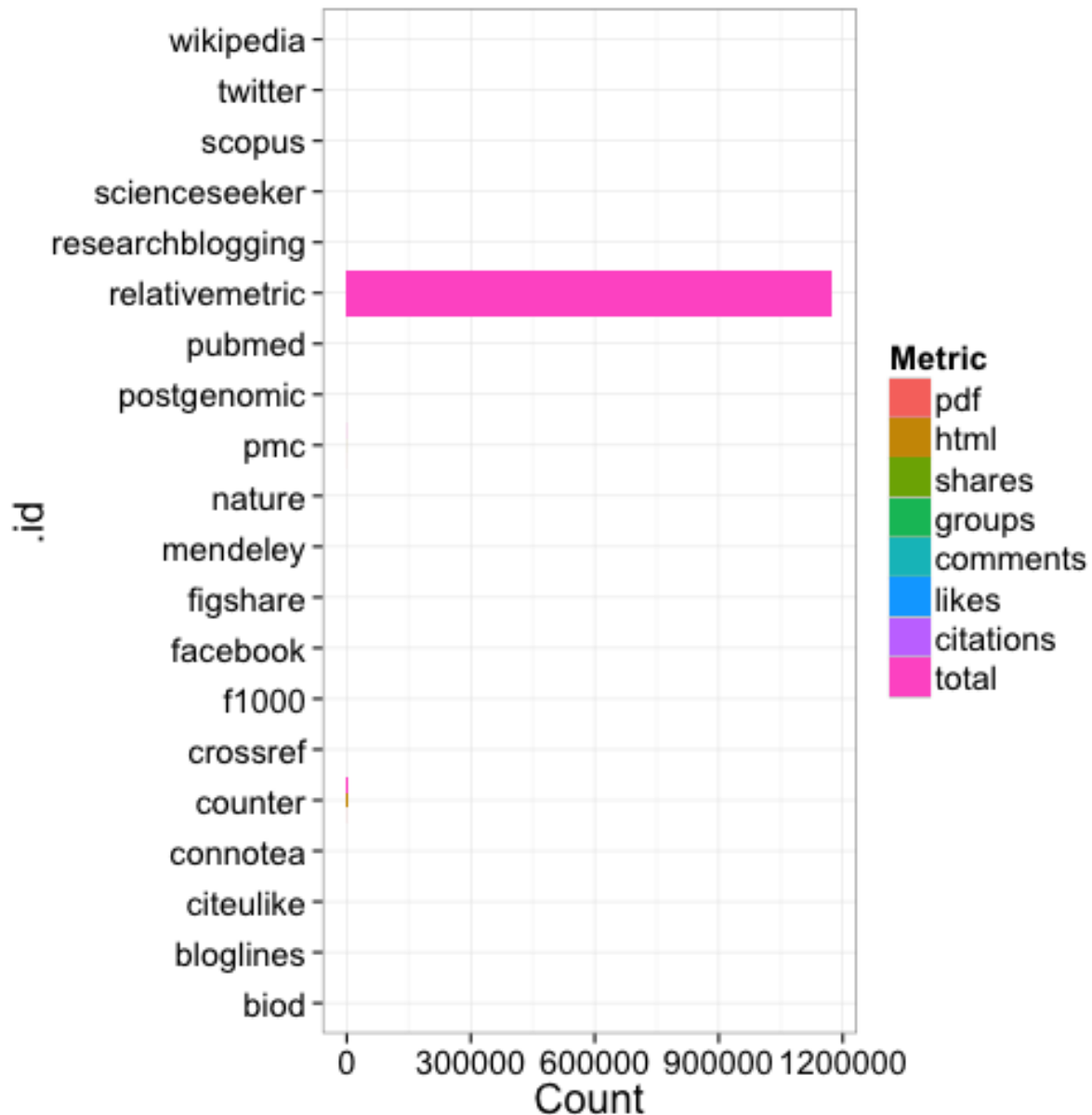


Figure 1: plot of chunk totalmets

Get detailed data for altmetrics using almevents

```
out <- almevents(doi = "10.1371/journal.pone.0029797")
names(out) # names of sources
```

```
[1] "bloglines"      "citeulike"      "connotea"
[4] "crossref"       "nature"         "postgenomic"
[7] "pubmed"         "scopus"         "counter"
[10] "researchblogging" "biod"           "pmc"
[13] "facebook"       "mendeley"       "twitter"
[16] "wikipedia"      "scienceseeker"  "relativemetric"
[19] "f1000"          "figshare"
```

```
out <- out[!out %in% c("sorry, no events content yet", "parser not written yet")] # remove those with
out[["pmc"]] # get the results for PubMed Central
```

	abstract	cited-by	figure	full-text	month	pdf	scanned-page-browse
1	1	0	9	51	1	8	0
2	0	0	11	15	2	4	0
3	0	0	0	11	3	4	0
4	1	0	0	6	4	2	0
5	0	0	0	5	5	1	0
6	0	0	2	7	6	2	0
7	1	0	3	6	7	3	0
8	1	0	0	5	8	0	0
9	0	0	3	14	9	5	0
10	1	0	1	20	10	4	0
11	0	0	1	13	12	1	0
12	1	0	0	10	11	1	0

	scanned-summary	supp-data	unique-ip	year
1		0	0	42 2012
2		0	2	11 2012
3		0	0	12 2012
4		0	0	6 2012
5		0	0	5 2012
6		0	0	9 2012
7		0	0	8 2012
8		0	0	4 2012
9		0	0	13 2012
10		0	0	16 2012
11		0	0	12 2012
12		0	0	9 2012

```
out[["twitter"]] # get the results for twitter (boo, there aren't any)
```

```
id
1 237088032224849920
2 237088322290331648
3 263798980054487041
4 263799348842872832
5 263960642589949953
6 282528931503038464
7 282528931612082177
```

```

8 284131287810338818
9 313850610174799873

```

```

1 #PLOS: Ecological Guild Evolution and the Discovery of the World's S
2 #PLOS: Ecological Guild Evolution and the Discovery of the World's S
3 Happy #Halloween from Maria @PLOSONE, dressed as a tiny frog, complete with dime for scale! ht
4 RT @multidiscipline: Happy #Halloween from Maria @PLOSONE, dressed as a tiny frog, complete with dime
5 RT @multidiscipline: Happy #Halloween from Maria @PLOSONE, dressed as a tiny frog, complete with dime
6 2012: Discovery of the World's Smallest Vertebrate: a frog http://t.co/BwxxRTdm in @PLOS
7 2012: Descubrimiento del vertebrado más pequeño: una rana http://t.co/BwxxRTdm in @PLOS
8
9 Ecological Guild Evolution and the Discovery of the World's Sma

```

	created_at	user	user_profile_image
1	Sun Aug 19 07:26:06 +0000 2012	opdebult	http://a0.twimg.com/profile_images/1741153180/Tidan_normal.jpg
2	Sun Aug 19 07:27:15 +0000 2012	forestalis	http://a0.twimg.com/profile_images/654250700/ForestalisIco_normal.jpg
3	Thu Nov 01 00:25:53 +0000 2012	multidiscipline	http://a0.twimg.com/profile_images/1910116023/261235_920680811178_6708085_43508969_7138379_n_normal.jpg
4	Thu Nov 01 00:27:20 +0000 2012	BernScience	http://a0.twimg.com/profile_images/1788875907/new_normal.jpg
5	Thu Nov 01 11:08:16 +0000 2012	mattjhodgkinson	http://a0.twimg.com/profile_images/2595571976/bc2za9tnyui0wxobreb0_normal.jpg
6	Sat Dec 22 16:52:01 +0000 2012	sferrebenedicto	http://a0.twimg.com/profile_images/2935384666/94c858315bbf621ae3916019026a6c24_normal.jpg
7	Sat Dec 22 16:52:01 +0000 2012	sferrebenedicto	http://a0.twimg.com/profile_images/2935384666/94c858315bbf621ae3916019026a6c24_normal.jpg
8	Thu Dec 27 02:59:12 +0000 2012	LeeAnaconda	http://a0.twimg.com/profile_images/2765018809/5b540749006aaf85c5661c67d93b68e7_normal.jpg
9	Tue Mar 19 03:13:11 +0000 2013	didicikit	http://a0.twimg.com/profile_images/1809269429/hop_normal.jpg

```

out[c("twitter", "crossref")] # get the results for two sources

```

```

$twitter
      id
1 237088032224849920
2 237088322290331648
3 263798980054487041
4 263799348842872832
5 263960642589949953
6 282528931503038464
7 282528931612082177
8 284131287810338818
9 313850610174799873

```

```

1 #PLOS: Ecological Guild Evolution and the Discovery of the World's S
2 #PLOS: Ecological Guild Evolution and the Discovery of the World's S
3 Happy #Halloween from Maria @PLOSONE, dressed as a tiny frog, complete with dime for scale! ht
4 RT @multidiscipline: Happy #Halloween from Maria @PLOSONE, dressed as a tiny frog, complete with dime
5 RT @multidiscipline: Happy #Halloween from Maria @PLOSONE, dressed as a tiny frog, complete with dime
6 2012: Discovery of the World's Smallest Vertebrate: a frog http://t.co/BwxxRTdm in @PLOS
7 2012: Descubrimiento del vertebrado más pequeño: una rana http://t.co/BwxxRTdm in @PLOS

```


8
9 Ecological Guild Evolution and the Discovery of the World's Smal

	created_at	user
1	Sun Aug 19 07:26:06 +0000 2012	opdebult
2	Sun Aug 19 07:27:15 +0000 2012	forestalis
3	Thu Nov 01 00:25:53 +0000 2012	multidiscipline
4	Thu Nov 01 00:27:20 +0000 2012	BernScience
5	Thu Nov 01 11:08:16 +0000 2012	mattjhodgkinson
6	Sat Dec 22 16:52:01 +0000 2012	sferrebenedicto
7	Sat Dec 22 16:52:01 +0000 2012	sferrebenedicto
8	Thu Dec 27 02:59:12 +0000 2012	LeeAnaconda
9	Tue Mar 19 03:13:11 +0000 2013	didicikit

user_profile_image

1	http://a0.twimg.com/profile_images/1741153180/Tidan_normal.jpg
2	http://a0.twimg.com/profile_images/654250700/ForestalisIco_normal.jpg
3	http://a0.twimg.com/profile_images/1910116023/261235_920680811178_6708085_43508969_7138379_n_normal.jpg
4	http://a0.twimg.com/profile_images/1788875907/new_normal.jpg
5	http://a0.twimg.com/profile_images/2595571976/bc2za9tnyui0wxobreb0_normal.jpg
6	http://a0.twimg.com/profile_images/2935384666/94c858315bbf621ae3916019026a6c24_normal.jpg
7	http://a0.twimg.com/profile_images/2935384666/94c858315bbf621ae3916019026a6c24_normal.jpg
8	http://a0.twimg.com/profile_images/2765018809/5b540749006aaf85c5661c67d93b68e7_normal.jpg
9	http://a0.twimg.com/profile_images/1809269429/hop_normal.jpg

\$crossref

	issn	journal_title
1	1439-6092; 1618-1077	Organisms Diversity & Evolution
2	1313-2970; 1313-2989	ZooKeys
3	00218790	Journal of Animal Ecology
4	1936-6426; 1936-6434	Evolution: Education and Outreach
5	0018-0831; 1938-5099	Herpetologica
6	10557903	Molecular Phylogenetics and Evolution
7	00244066	Biological Journal of the Linnean Society

	journal_abbreviation
1	Org Divers Evol
2	ZOOKEYS
3	J Anim Ecol
4	Evo Edu Outreach
5	Herpetologica
6	Molecular Phylogenetics and Evolution
7	Biol J Linn Soc Lond

1 New insights into the systematics and molecular phylogeny of the Malagasy snake genus Liopholidophis s
2 Accele
3
4
5 A New Species of Miniaturized Toadlet, GenusBrachycephalus
6 Genetic diversity, phylogeny and evolution of alkaloid s
7 Are dim
contribu

1 Frank Glaw; Christoph Kucharzewski; Zoltán T. Nagy; Oliver Hawlitschek; Miguel Vences
2 Terry Erwin; Lyubomir Penev; Pavel Stoev; Teodor Georgiev
3 Andrew D. C. MacColl; Aliya El Nagar; Job de Roij; Tom W. S. van Balbeek
4 Kenneth J. McNamara
5 Rute B. G. Clemente-Carvalho; Ariovaldo A. Giaretta; Thais H. Condez; Célio F. B. Haddad; Sergio F. dos Reis

```

6                                Ariel Rodríguez; Dennis Poth; Stefan Schulz; Marcelo Gehara; Miguel Ven
7                                Kenneth D. Angielczyk; Chris R. Feldr
  year publication_type          doi fl_count volume
1 2013      full_text    10.1007/s13127-013-0152-4      0  <NA>
2 2012      full_text    10.3897/zookeys.251.4516      1   251
3 2013      full_text    10.1111/1365-2656.12028      0    82
4 2012      full_text    10.1007/s12052-012-0420-3      1     5
5 2012      full_text 10.1655/HERPETOLOGICA-D-11-00085.1      0    68
6 2013      full_text    10.1016/j.ympev.2013.04.031      0    68
7 2013      full_text    10.1111/bij.12010      0   108
  issue first_page
1  <NA>      <NA>
2     0         1
3     3       642
4     2       203
5     3       365
6     3       541
7     4       727

```

Retrieve and plot PLOS article-level metrics signposts.

```

dat <- signposts(doi = "10.1371/journal.pone.0029797")
plot_signposts(input = dat)

```

Or plot many identifiers gives a line chart

```

dois <- c("10.1371/journal.pone.0001543", "10.1371/journal.pone.0040117", "10.1371/journal.pone.0029797",
          "10.1371/journal.pone.0039395")
dat <- signposts(doi = dois)
plot_signposts(input = dat)

```

Or make an interactive chart by doing `plot_signposts(input=dat, type="multiBarChart")`. Try it out! It should open in your browser and you can interact with it.

Density and histogram plots from PLOS Article Level Metrics data

Note: Do you the key below in the `searchplos` call in this example, but if you plan to use `rplos` more, get your own API key [here](#).

```

library(rplos)
library(plyr)
dois <- searchplos(terms = ":*:", fields = "id", toquery = list("cross_published_journal_key:PLoS ONE",
  "doc_type:full", "publication_date:[2010-01-01T00:00:00Z TO 2010-12-31T23:59:59Z]"),
  limit = 100, key = "WQcDSXm12VSWx3P")
alm <- alm(doi = do.call(c, dois$id), total_details = TRUE)
alm <- ldply(alm)

```

The default plot

```
plot_density(alm)
```

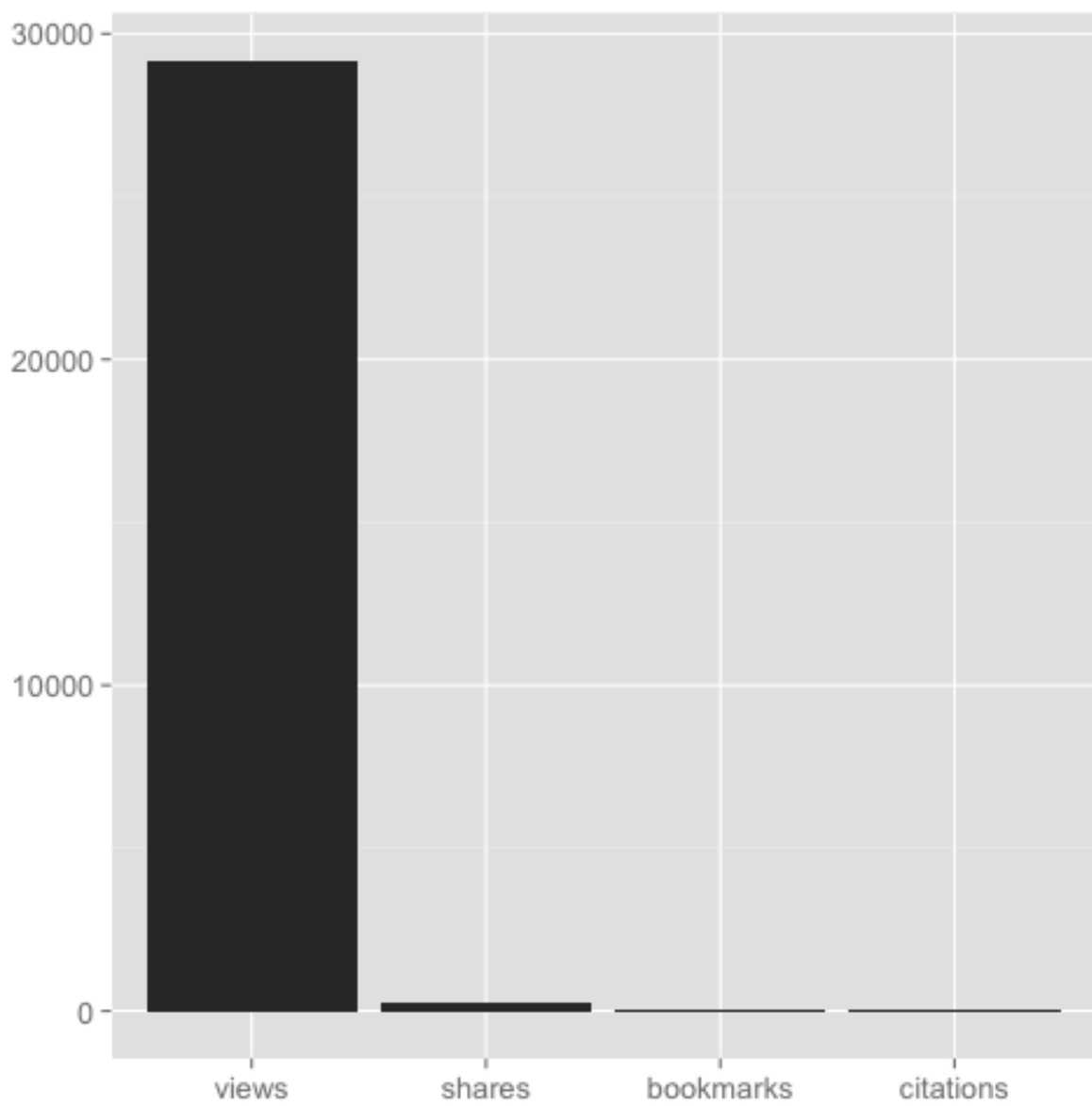


Figure 2: plot of chunk signposts1

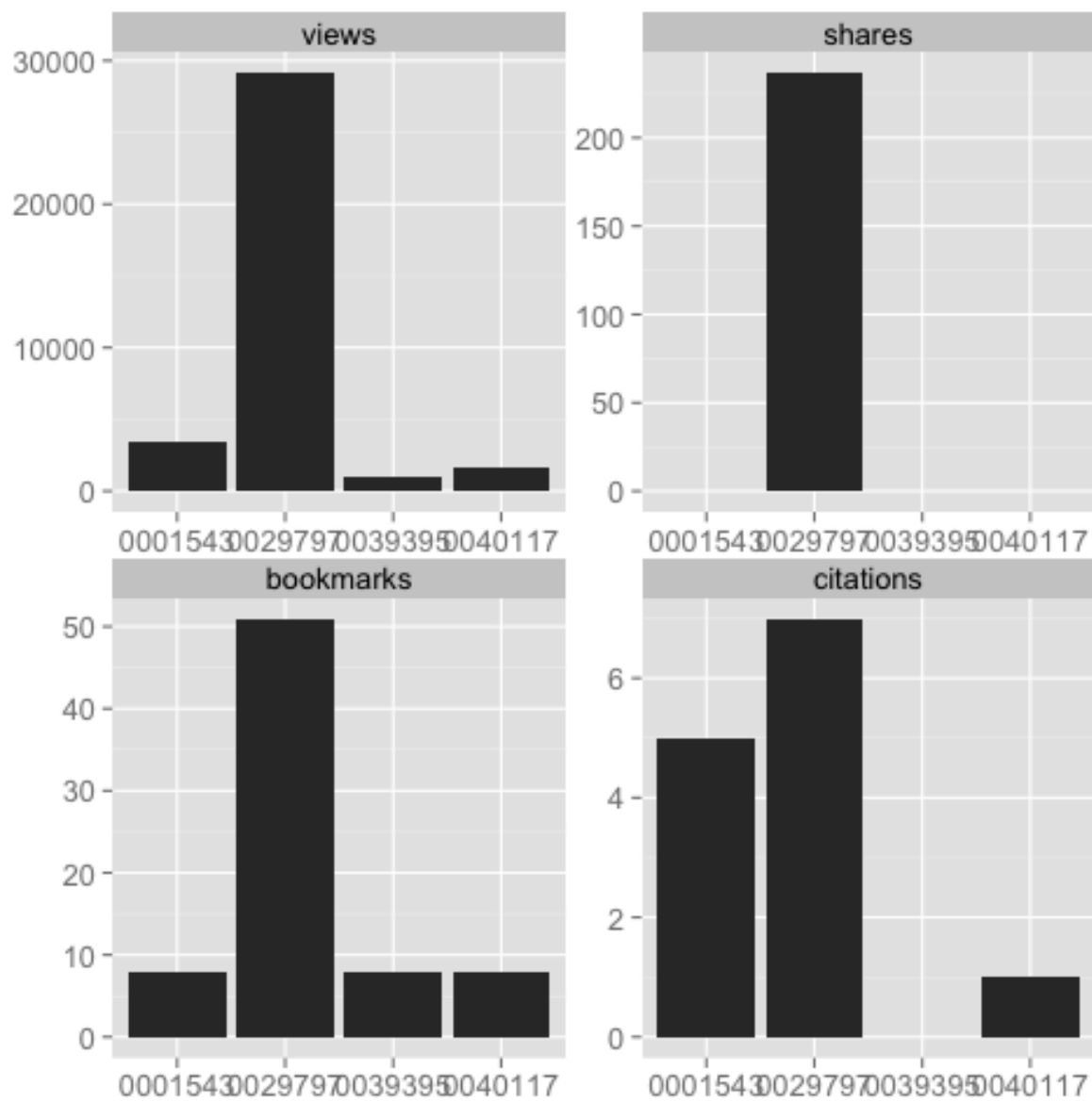


Figure 3: plot of chunk signposts2

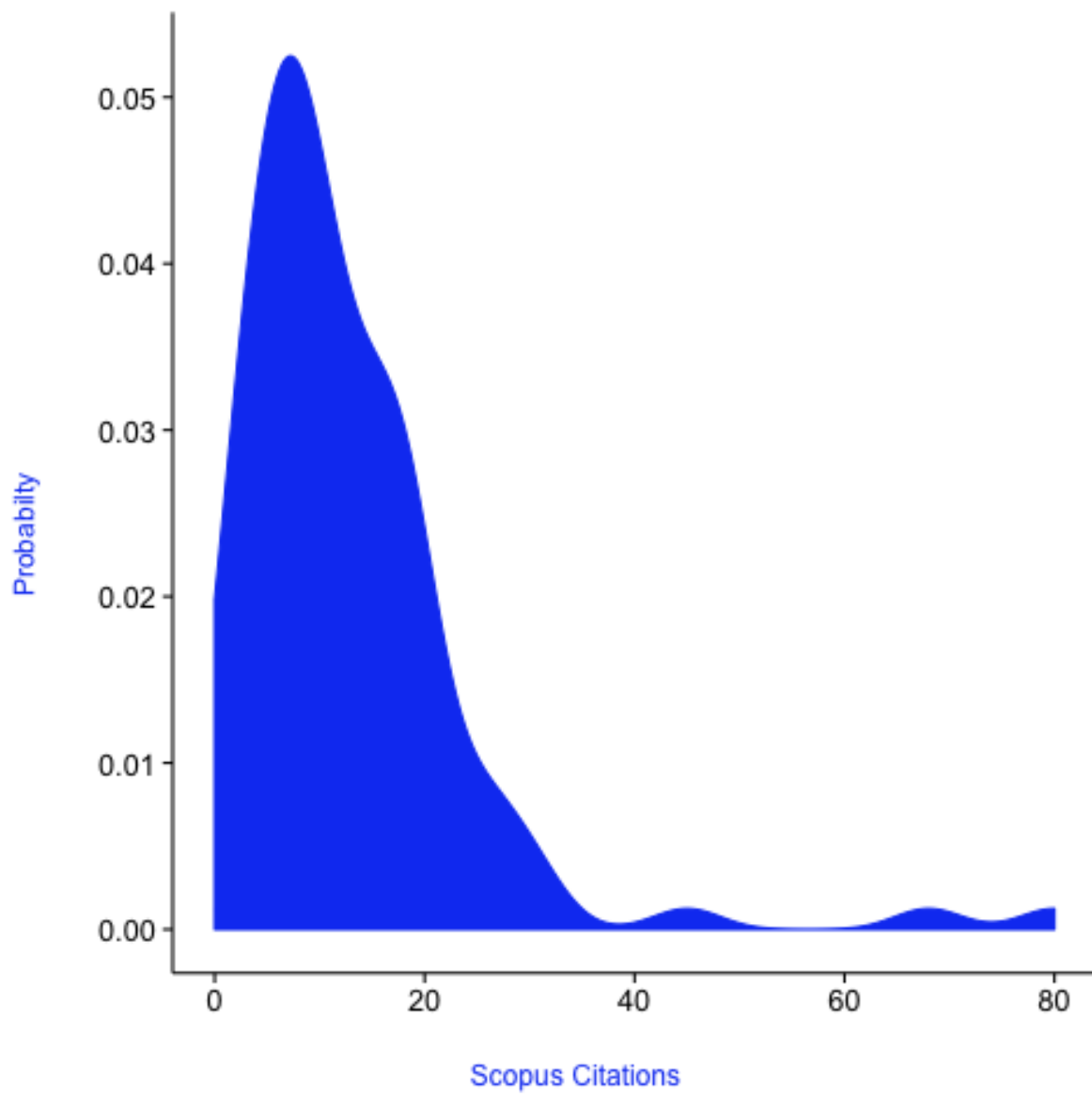


Figure 4: plot of chunk plot_densityplot1

You can change the color of the density plot

```
plot_density(alm, color = "#EFA5A5")
```

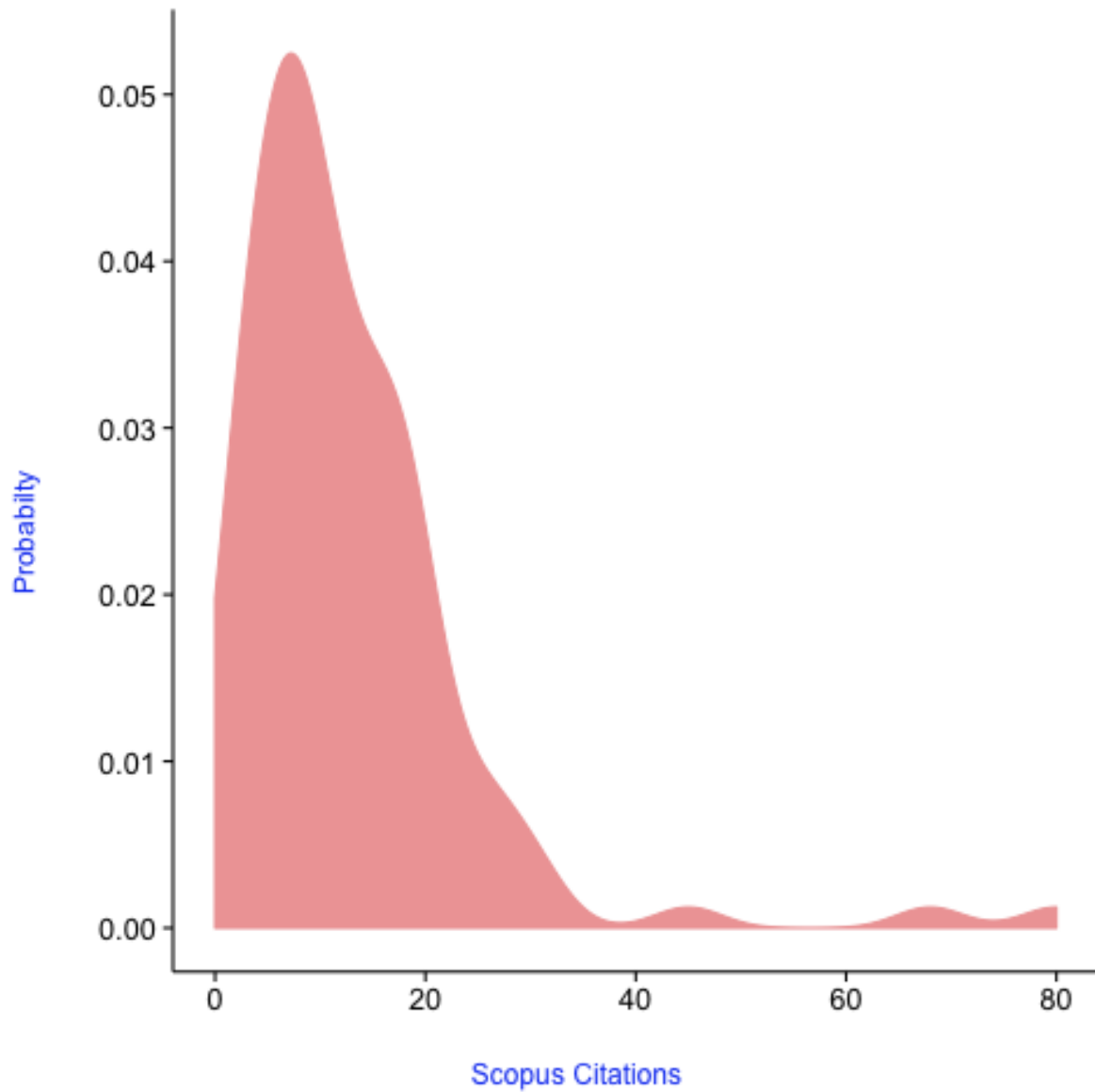


Figure 5: plot of chunk plot_densityplot2

Pass in a title or description subtending the title

```
plot_density(alm, title = "Scopus citations from 2010")
```

Plot a particular source

```
names(alm)[1:35]
```

Scopus citations from 2010

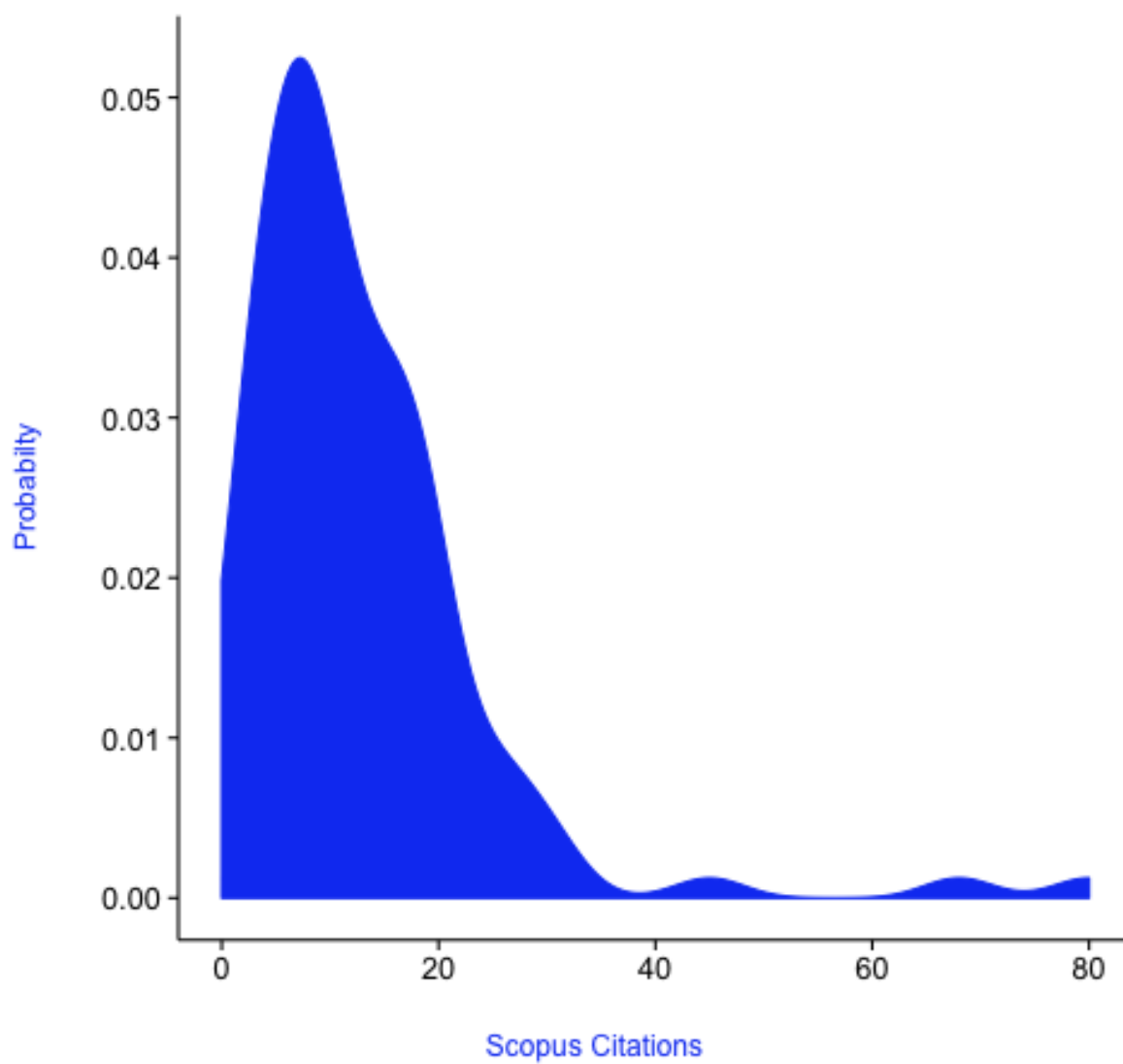


Figure 6: plot of chunk plot_densityplot3

```

[1] ".id"           "doi"           "title"
[4] "publication_date" "bloglines_pdf" "bloglines_html"
[7] "bloglines_shares" "bloglines_groups" "bloglines_comments"
[10] "bloglines_likes" "bloglines_citations" "bloglines_total"
[13] "citeulike_pdf" "citeulike_html" "citeulike_shares"
[16] "citeulike_groups" "citeulike_comments" "citeulike_likes"
[19] "citeulike_citations" "citeulike_total" "connotea_pdf"
[22] "connotea_html" "connotea_shares" "connotea_groups"
[25] "connotea_comments" "connotea_likes" "connotea_citations"
[28] "connotea_total" "crossref_pdf" "crossref_html"
[31] "crossref_shares" "crossref_groups" "crossref_comments"
[34] "crossref_likes" "crossref_citations"

```

```
plot_density(input = alm, source = "crossref_citations")
```

Plot many sources in different panels in the same plot, and pass in colors just for fun

```

plot_density(input = alm, source = c("counter_total", "crossref_citations",
  "twitter_total", "wos_citations"), color = c("#83DFB4", "#EFA5A5", "#CFD470",
  "#B2C9E4"))

```

Alt-metrics total citations from all sources.

```
almtotals(doi = "10.1371/journal.pbio.0000012")
```

```

  views shares bookmarks citations
1 29349      0         79       143

```

Get title of article by inputting the doi for the article.

```
almtitle(doi = "10.1371/journal.pbio.0000012")
```

```
[1] "Genome-Wide RNAi of C. elegans Using the Hypersensitive rrf-3 Strain Reveals Novel Gene Functions"
```

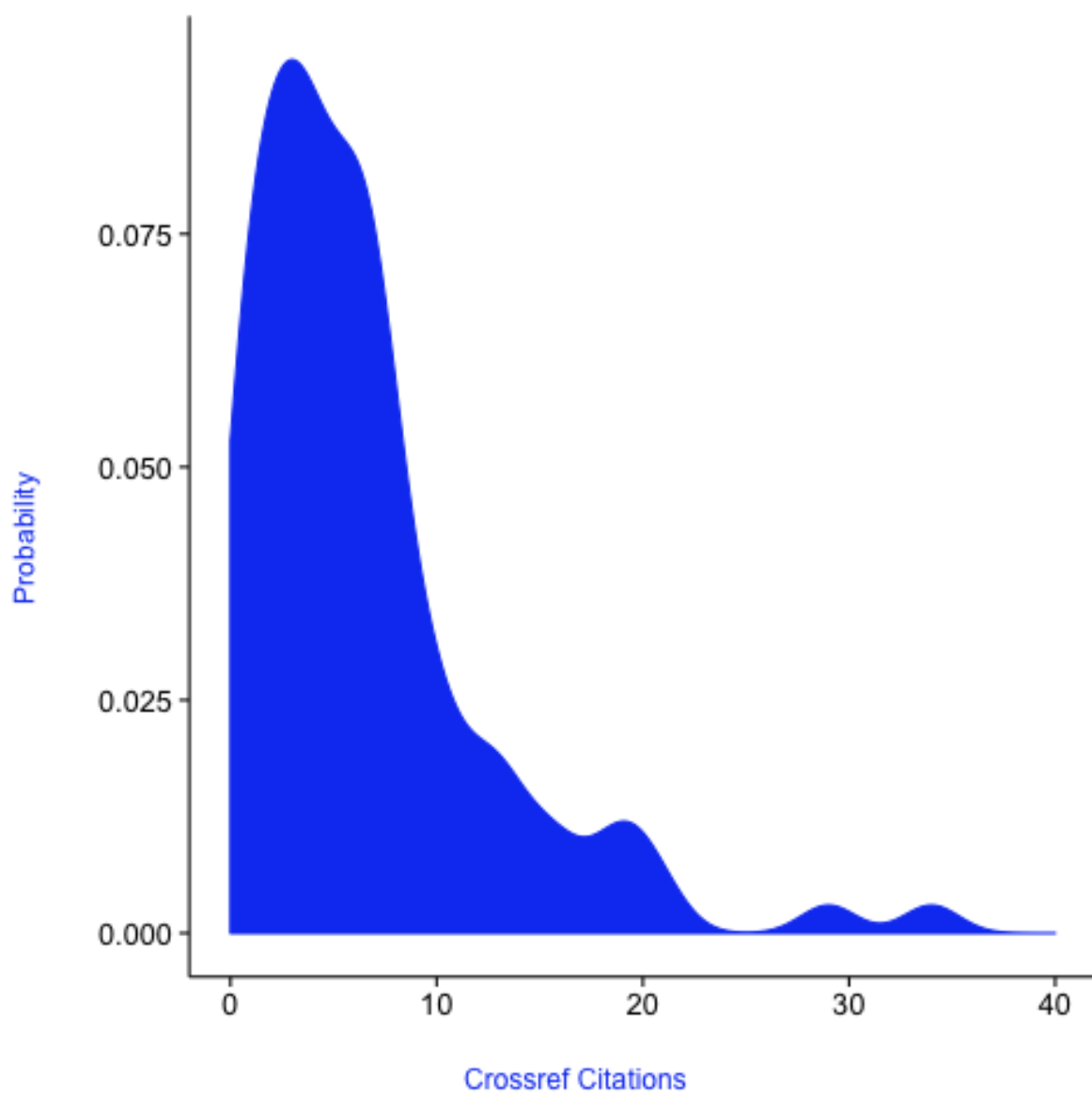



Figure 7: plot of chunk plot_densityplot4

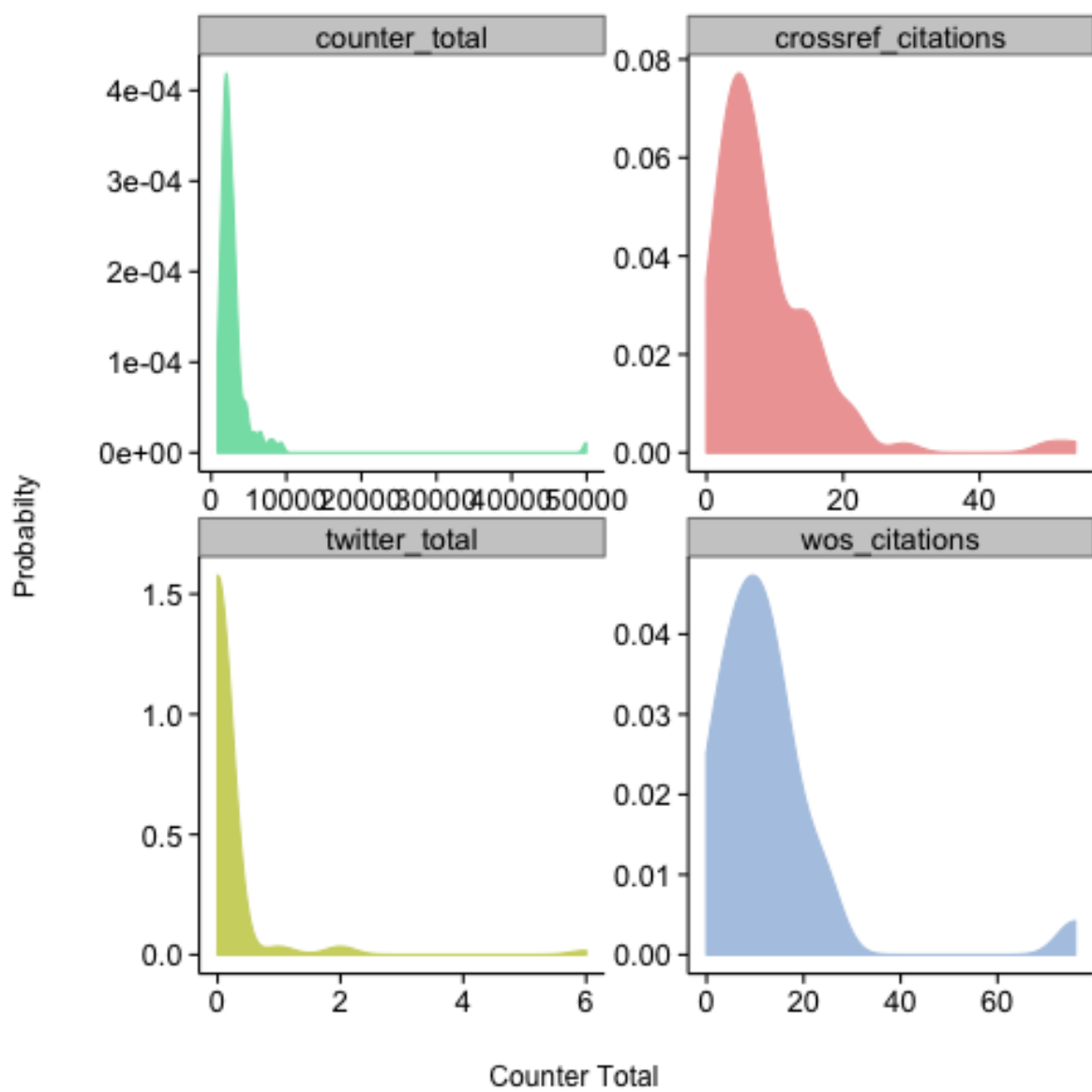


Figure 8: plot of chunk plot_densityplot5