# Impact\_Journal\_Count

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# **Import Author Information**

Import list of scholars. Some of the scolars do not have a Google Scholar page so their unique ID was mared as "NA." They are not considered in the analysis. If it is necessary to have their citation information we can count their citations manually.

```
df <- readxl::read_xlsx("./PI_google_scholar_ids.xlsx")</pre>
head(df, n = 3)
## # A tibble: 3 x 5
     title last_name first_name institution
                                                           google_scholar_id
                                                           <chr>
     <chr> <chr>
                     <chr>
                                <chr>>
                     Huimin
                                U of Ill Urbana-Champaign R8waAyOAAAAJ
## 1 PI
           Zhao
## 2 coPI Burke
                     Martin
                                U of Ill Urbana-Champaign u4hmNXAAAAAJ
## 3 coPI Denmark
                     Scott
                                U of Ill Urbana-Champaign NA
print(dim(df)) #Rows, Columns
```

## [1] 16 5

### Extract High Impact Journal Information

Any journal with "Nature" in it's title is captured with str\_extract and articles published in "Science" were captured with an exact character match.

```
pubs <- lapply(df$google_scholar_id, get_publications)

impact_journal_count <- c()
nature <- c()
nature_sister <- c()
science <- c()
log_check <- c()
for (i in 1:length(pubs)){
        author <- pubs[[i]]
        #nature count
        nature_tot_count <- length(author$journal[as.logical(str_count(author$journal,"Nature"))])
        nature_sister_count <- nature_tot_count - nature_count
        nature_log <- as.logical(str_count(author$journal,"Nature"))</pre>
```

```
#science count
      science_count <- length(author$journal[author$journal == "Science"])</pre>
      science_log <- author$journal == "Science"</pre>
      total_count <- nature_tot_count + science_count</pre>
      total_log <- list(nature_log + science_log)</pre>
      impact_journal_count <- c(impact_journal_count,total_count)</pre>
      log_check <- c(log_check,total_log)</pre>
      nature <- c(nature,nature_count)</pre>
      nature_sister <- c(nature_sister, nature_sister_count)</pre>
      science <- c(science, science_count)</pre>
}
num_cites <- c()</pre>
for (i in 1:nrow(df)){
      if (df$google_scholar_id[i] != "NA"){
             num_cites <- c(num_cites, get_profile(df$ google_scholar_id[i])$total_cites)</pre>
      else {
             num_cites <- c(num_cites, 0)</pre>
      }
}
df_impact <- mutate(df, nature, nature_sister, science, impact_journal_count, num_cites)</pre>
```

# Missing Google Scholar Profiles

For Authors without a Google Scholar profile the data was manually downloaded from the author profile page on Web of Science. Profile pages for Scott E. Denmark and Kenton McHenry exist but a profile page for Rachel Switzky could not be found. Scott E. Denmark has two search result hits, these were combined then downloaded. Kenton McHenry only has one one search result hit. This portion of the analysis is not completely reproducible as Web of Science requires log in credentials.

```
# Denmark, Scott E.
denmark <- read.table("Denmark_Scott_2020_05_22.txt", header = TRUE, sep = ",", skip = 2)</pre>
denmark_nature_tot <- sum(as.logical(str_count(denmark$Source.Title, "NATURE")))</pre>
df_impact$nature[3] <- sum(denmark$Source.Title == "NATURE")</pre>
df_impact$nature_sister[3] <- denmark_nature_tot - df_impact$nature[3]</pre>
df_impact$science[3] <- sum(denmark$Source.Title == "SCIENCE")</pre>
df_impact$impact_journal_count[3] <- df_impact$nature[3] + df_impact$nature_sister[3] + df_impact$scien</pre>
df_impact$num_cites[3] <- sum(denmark$Total.Citations)</pre>
denmark_log <- as.logical(str_count(denmark$Source.Title, "NATURE") + as.integer(denmark$Source.Title =</pre>
# McHenry, Kenton
mchenry <- read.table("McHenry_Kenton_2020_05_22.txt", header = TRUE, sep = ",", skip = 2)</pre>
mchenry_nature_tot <- sum(as.logical(str_count(mchenry$Source.Title, "NATURE")))</pre>
df_impact$nature[10] <- sum(mchenry$Source.Title == "NATURE")</pre>
df_impact$nature_sister[10] <- mchenry_nature_tot - df_impact$nature[10]</pre>
df_impact$science[10] <- sum(mchenry$Source.Title == "SCIENCE")</pre>
df_impact$impact_journal_count[10] <- df_impact$nature[10] + df_impact$nature_sister[10] + df_impact$sc
df_impact$num_cites[10] <- sum(mchenry$Total.Citations)</pre>
```

mchenry\_log <- as.logical(str\_count(mchenry\$Source.Title, "NATURE") + as.integer(mchenry\$Source.Title =</pre>

# **Team Totals**

Total number of publications from the team classified as: Nature publication, Nature Sister publication, Science publication

```
publications_totals <- data.frame(sum(nature), sum(nature_sister), sum(science))
names(publications_totals) <- c("Nature Publications", "Nature Sister Journal Publications", "Science Pub
knitr::kable(publications_totals, caption = "Team Publication Totals")</pre>
```

Table 1: Team Publication Totals

Nature Publications	Nature Sister Journal Publications	Science Publications
13	95	16

# Average Citation per Author

The average number of citations per team member including the three team members without google scholar profiles.

knitr::kable(mean(df\_impact\$num\_cites), col.name = "Average", caption = "Average citation per team memb

Table 2: Average citation per team member

 $\frac{\text{Average}}{20368.88}$ 

# All Results

Table of Results

knitr::kable(df\_impact)

title	$last\_name$	$first\_name$	institution	$google\_scholar\_id$	nature	$nature\_sister$	science
PI	Zhao	Huimin	U of Ill Urbana-Champaign	R8waAy0AAAAJ	2	19	0
coPI	Burke	Martin	U of Ill Urbana-Champaign	u4hmNXAAAAAJ	1	9	3
coPI	Denmark	Scott	U of Ill Urbana-Champaign	NA	1	4	2
coPI	Peng	Jian	U of Ill Urbana-Champaign	H2JX-RQAAAAJ	0	9	1
coPI	Sinha	Saurabh	U of Ill Urbana-Champaign	7BM1uyYAAAAJ	1	4	3
srPe	Diao	Ying	U of Ill Urbana-Champaign	ZbEs7rsAAAAJ	0	7	0
srPe	Han	Jiawei	U of Ill Urbana-Champaign	Kv9AbjMAAAAJ	0	0	0
$\operatorname{srPe}$	Ji	Heng	U of Ill Urbana-Champaign	z7GCqT4AAAAJ	0	1	0
srPe	Maranas	Costas	PA St U University Park	l3M1PW0AAAAJ	0	10	0
srPe	McHenry	Kenton	U of Ill Urbana-Champaign	NA	0	0	0
srPe	Schroeder	Charles	U of Ill Urbana-Champaign	wIY9shsAAAAJ	0	4	1
srPe	Shukla	Diwakar	U of Ill Urbana-Champaign	${ m Cm2dWNYAAAAJ}$	0	6	0
srPe	Switzky	Rachel	U of Ill Urbana-Champaign	NA	0	0	0
$\operatorname{srPe}$	White	Maria	U of Ill Urbana-Champaign	ICPpNtkAAAAJ&hl	2	5	3

title	$last\_name$	$first\_name$	institution	$google\_scholar\_id$	nature	$nature\_sister$	science
	Zanibbi	Richard	Rochester Inst of Tech	k6xVb1QAAAAJ	0	0	0
coPI	Grzybowski	Bartosz A.	Northwestern University	hjS0xZ0AAAAJ	7	21	5

knitr::kable(select(df\_impact, last\_name, first\_name, impact\_journal\_count, num\_cites))

last_name	first_name	$impact\_journal\_count$	num_cites
Zhao	Huimin	21	18376
Burke	Martin	13	7091
Denmark	Scott	7	26388
Peng	Jian	10	6268
Sinha	Saurabh	8	10301
Diao	Ying	7	5430
Han	Jiawei	0	187215
Ji	Heng	1	6559
Maranas	Costas	10	15798
McHenry	Kenton	0	105
Schroeder	Charles	5	3144
Shukla	Diwakar	6	2740
Switzky	Rachel	0	0
White	Maria	10	9075
Zanibbi	Richard	0	2119
Grzybowski	Bartosz A.	33	25293

# Titles of Author's High Impact Publications

The following are titles of the high impact papers (Nature, Nature Sister, and Science) published by each Author. Authors that did not have Google Scholar Profiles were omitted from this part of the analysis.

```
for (i in 1:nrow(df_impact)){
      last_name <- df_impact$last_name[i]</pre>
      author_col_name <- (sprintf("%s's High Impact Publications",last_name))</pre>
      if (i != 3 & i != 10){
             authors_pubs <- pubs[[i]]$title[as.logical(log_check[[i]])]</pre>
             print(i)
      }
      else if (i == 3){
             #Scott E Denmark
             authors_pubs <- denmark$Title[denmark_log]</pre>
      else if (i == 10){
             #Kenton McHenry
             authors_pubs <- mchenry$Title[mchenry_log]</pre>
      }
      print(knitr::kable(authors_pubs, col.names = "High Impact Publication Titles",
                    caption = author_col_name))
      cat('\n')
```

## Table 5: Zhao's High Impact Publications

### High Impact Publication Titles

Molecular evolution by staggered extension process (StEP) in vitro recombination

Chromatin architecture reorganization during stem cell differentiation

Activation and characterization of a cryptic polycyclic tetramate macrolactam biosynthetic gene cluster

CRISPR-Cas9 strategy for activation of silent Streptomyces biosynthetic gene clusters

Automated multiplex genome-scale engineering in yeast

In vitro 'sexual' evolution through the PCR-based staggered extension process (StEP)

Combinatorial metabolic engineering using an orthogonal tri-functional CRISPR system

Direct observation of TALE protein dynamics reveals a two-state search mechanism

Cooperative asymmetric reactions combining photocatalysis and enzymatic catalysis

Genome-scale engineering of Saccharomyces cerevisiae with single-nucleotide precision

Direct cloning of large genomic sequences

TALE proteins search DNA using a rotationally decoupled mechanism

Building a global alliance of biofoundries

Genome-wide identification of natural RNA aptamers in prokaryotes and eukaryotes

Activation of silent biosynthetic gene clusters using transcription factor decoys

Indoleacetate decarboxylase is a glycyl radical enzyme catalysing the formation of malodorant skatole

Radical-mediated CS bond cleavage in C2 sulfonate degradation by anaerobic bacteria

Towards a fully automated algorithm driven platform for biosystems design

An efficient gene knock-in strategy using 5'-modified double-stranded DNA donors with short homology

Multi-functional genome-wide CRISPR system for high throughput genotype-phenotype mapping

DNA punch cards for storing data on native DNA sequences via enzymatic nicking

[1] 2

Table 6: Burke's High Impact Publications

#### High Impact Publication Titles

Generating diverse skeletons of small molecules combinatorially

Synthesis of many different types of organic small molecules using one automated process

Amphotericin forms an extramembranous and fungicidal sterol sponge

Synthesis of most polyene natural product motifs using just 12 building blocks and one coupling reaction Nontoxic antimicrobials that evade drug resistance

MIDA boronates are hydrolysed fast and slow by two different mechanisms

Electronic tuning of site-selectivity

Restored iron transport by a small molecule promotes absorption and hemoglobinization in animals

Towards the generalized iterative synthesis of small molecules

Small-molecule ion channels increase host defences in cystic fibrosis airway epithelia

Axial shielding of Pd(II) complexes enables perfect stereoretention in Suzuki-Miyaura cross-coupling of Csp 3 boronic acids

Toward generalization of iterative small molecule synthesis

Flexible tetracycline synthesis yields promising antibiotics

Table 7: Denmark's High Impact Publications

Pre-transmetalation intermediates in the Suzuki-Miyaura reaction revealed: The missing link Catalytic, stereospecific syn-dichlorination of alkenes

Mechanistic, crystallographic, and computational studies on the catalytic, enantioselective sulfenofunctionalization of alkenes

N-silyl oxyketene imines are underused yet highly versatile reagents for catalytic asymmetric synthesis Prediction of higher-selectivity catalysts by computer-driven worlflow and machine learning Organic chemistry - Catalysts break symmetry

Demystifying the asymmetry-amplifying, autocatalytic behaviour of the Soai reaction through structural, mechanistic and computational studies

[1] 4

Table 8: Peng's High Impact Publications

#### High Impact Publication Titles

Template-based protein structure modeling using the RaptorX web server

A community effort to assess and improve drug sensitivity prediction algorithms

Computational solutions for omics data

Selection bias at the heterosexual HIV-1 transmission bottleneck

A community computational challenge to predict the activity of pairs of compounds

A network integration approach for drug-target interaction prediction and computational drug repositioning from heterogeneous information

Quality score compression improves genotyping accuracy

Community assessment to advance computational prediction of cancer drug combinations in a pharmacogenomic screen

Compressive mapping for next-generation sequencing

Typing tumors using pathways selected by somatic evolution

[1] 5

Table 9: Sinha's High Impact Publications

### High Impact Publication Titles

Assessing computational tools for the discovery of transcription factor binding sites. The genome of a songbird

Functional and evolutionary insights from the genomes of three parasitoid Nasonia species

Genomic signatures of evolutionary transitions from solitary to group living

Core and region-enriched networks of behaviorally regulated genes and the singing genome Intravital imaging by simultaneous label-free autofluorescence-multiharmonic microscopy

Towards a fully automated algorithm driven platform for biosystems design

ChIPs and regulatory bits

[1] 6

Solution coating of large-area organic semiconductor thin films with aligned single-crystalline domains Flow-enhanced solution printing of all-polymer solar cells

The role of nanopore shape in surface-induced crystallization

One-dimensional self-confinement promotes polymorph selection in large-area organic semiconductor thin films

Dynamic-template-directed multiscale assembly for large-area coating of highly-aligned conjugated polymer thin films

Rotator side chains trigger cooperative transition for shape and function memory effect in organic semiconductors

Repurposing DNA-binding agents as H-bonded organic semiconductors

[1] 7

Table 11: Han's High Impact Publications

High Impact Publication Titles

[1] 8

Table 12: Ji's High Impact Publications

High Impact Publication Titles

Moralization in Social Networks and the Emergence of Violent Protests

[1] 9

Table 13: Maranas's High Impact Publications

### High Impact Publication Titles

Creation and analysis of biochemical constraint-based models using the COBRA Toolbox v. 3.0 A genome-scale Escherichia coli kinetic metabolic model k-ecoli457 satisfying flux data for multiple mutant strains

Pathway design using de novo steps through uncharted biochemical spaces

PoreDesigner for tuning solute selectivity in a robust and highly permeable outer membrane pore EcoFABs: advancing microbiome science through standardized fabricated ecosystems

A microbial factory for diverse chemicals

Artificial water channels enable fast and selective water permeation through water-wire networks Bacterial colonization reprograms the neonatal gut metabolome

Author Correction: Artificial water channels enable fast and selective water permeation through water-wire networks

Enzymes: Orchestrating hi-fi annotations

Table 14: McHenry's High Impact Publications

[1] 11

Table 15: Schroeder's High Impact Publications

# High Impact Publication Titles

Observation of polymer conformation hysteresis in extensional flow

Multiplexed single-molecule assay for enzymatic activity on flow-stretched DNA

Direct observation of TALE protein dynamics reveals a two-state search mechanism

TALE proteins search DNA using a rotationally decoupled mechanism

Effect of molecular architecture on ring polymer dynamics in semidilute linear polymer solutions

[1] 12

Table 16: Shukla's High Impact Publications

#### High Impact Publication Titles

Cloud-based simulations on Google Exacycle reveal ligand modulation of GPCR activation pathways Activation pathway of Src kinase reveals intermediate states as targets for drug design

Dynamic-template-directed multiscale assembly for large-area coating of highly-aligned conjugated polymer thin films

Conformational heterogeneity of the calmodulin binding interface

A network of molecular switches controls the activation of the two-component response regulator NtrC Corrigendum: Cloud-based simulations on Google Exacycle reveal ligand modulation of GPCR activation pathways

[1] 13

Table 17: Switzky's High Impact Publications

 ${\bf High\ Impact\ Publication\ Titles}$ 

[1] 14

Table 18: White's High Impact Publications

## High Impact Publication Titles

A predictably selective aliphatic C-H oxidation reaction for complex molecule synthesis

Combined effects on selectivity in Fe-catalyzed methylene oxidation

Adding aliphatic C-H bond oxidations to synthesis

Total synthesis and study of 6-deoxyerythronolide B by late-stage C-H oxidation

A manganese catalyst for highly reactive yet chemoselective intramolecular C(sp3)-H amination

Diverting non-haem iron catalysed aliphatic C-H hydroxylations towards desaturations

Oxidative diversification of amino acids and peptides by small-molecule iron catalysis

Manganese-catalysed benzylic C (sp 3)-H amination for late-stage functionalization

Chemoselective methylene oxidation in aromatic molecules

Late-stage oxidative C (sp 3)-H methylation

[1] 15

Table 19: Zanibbi's High Impact Publications

High Impact Publication Titles

[1] 16

Table 20: Grzybowski's High Impact Publications

## High Impact Publication Titles

Self-assembly at all scales

The mosaic of surface charge in contact electrification

Dynamic self-assembly of magnetized, millimetre-sized objects rotating at a liquid-air interface

Directing cell motions on micropatterned ratchets

Plastic and moldable metals by self-assembly of sticky nanoparticle aggregates

Photoconductance and inverse photoconductance in films of functionalized metal nanoparticles

Electrostatic self-assembly of macroscopic crystals using contact electrification

The nanotechnology of life-inspired systems

Colloidal assembly directed by virtual magnetic moulds

Ultrasensitive detection of toxic cations through changes in the tunnelling current across films of striped nanoparticles

Dynamic hook-and-eye nanoparticle sponges

Geometric curvature controls the chemical patchiness and self-assembly of nanoparticles

The 'wired' universe of organic chemistry

Multicolour micropatterning of thin films of dry gels

Control of surface charges by radicals as a principle of antistatic polymers protecting electronic circuitry Self-assembly of polymeric microspheres of complex internal structures

Molecular dynamics imaging in micropatterned living cells

Dynamic aggregation of chiral spinners

Chemoelectronic circuits based on metal nanoparticles

Systems of mechanized and reactive droplets powered by multi-responsive surfactants

Charged nanoparticles as supramolecular surfactants for controlling the growth and stability of microcrystals

Dynamic internal gradients control and direct electric currents within nanostructured materials Vortex flows impart chirality-specific lift forces

A long-lasting concentration cell based on a magnetic electrolyte

Automatic mapping of atoms across both simple and complex chemical reactions

Lévy-like movement patterns of metastatic cancer cells revealed in microfabricated systems and implicated in vivo

Oscillating droplet trains in microfluidic networks and their suppression in blood flow

Interference-like patterns of static magnetic fields imprinted into polymer/nanoparticle composites

Targeted crystallization of mixed-charge nanoparticles in lysosomes induces selective death of cancer cells

Enhancing crystal growth using polyelectrolyte solutions and shear flow

Author Correction: Systems of mechanized and reactive droplets powered by multi-responsive surfactants.

Systems of mechanized and reactive droplets powered by multi-responsive surfactants (vol 553, pg 313, 2018)

Nanosystem: Programmed communication