Sciatica PubMed

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This script takes ten articles from the abstracts on sciatic articles from NCBI's PubMed

This creates a directory to stem the abstracts and preprocess from the csv file into a corpus of 20 files in a folder called Sciatica.

This code preprocesses and stems the corpus

```
library(tm)
library(SnowballC)
library(wordcloud)
library(ggplot2)
Sciatica <- Corpus(DirSource("Sciatica"))</pre>
Sciatica
## <<SimpleCorpus>>
## Metadata: corpus specific: 1, document level (indexed): 0
## Content: documents: 20
#Sciatica <- tm_map(Sciatica, removePunctuation)
#Sciatica <- tm_map(Sciatica, removeNumbers)</pre>
Sciatica <- tm_map(Sciatica, tolower)</pre>
Sciatica <- tm_map(Sciatica, removeWords, stopwords("english"))</pre>
Sciatica <- tm_map(Sciatica, stripWhitespace)</pre>
Sciatica <- tm map(Sciatica, stemDocument)
dtmSciatica <- DocumentTermMatrix(Sciatica)</pre>
```

freq <- colSums(as.matrix(dtmSciatica))</pre>

This code orders words stemmed by frequency and finds input correlations

```
FREQ <- data.frame(freq)
ord <- order(freq, decreasing=TRUE)
freq[head(ord, 25)]</pre>
```

##	nerv	sciatic	patient	pain	studi	muscl	control
##	68	59	56	41	29	22	20
##	postop	piriformi	clinic	signific	associ	block	measur
##	18	17	16	16	16	16	14
##	sciatica	group	show	produc	effect	assess	am1710
##	13	13	13	12	12	12	12
##	imag	outcom	receiv	result			
##	11	11	11	10			

findAssocs(dtmSciatica, "effect", corlimit=0.7)

##	\$effect		
##	manag	hour	(cg).
##	0.85	0.78	0.76
##	(f?=?16.26;	(partial	(pisp)
##	0.76	0.76	0.76
##	(stens)	(tens)	(tg)
##	0.76	0.76	0.76
##	10?week	10th	19%
##	0.76	0.76	0.76
##	2018.	29th	2nd,
##	0.76	0.76	0.76
##	4th,	6th,	8th
##	0.76	0.76	0.76
##	account	administr	africa
##	0.76	0.76	0.76
##	allot	analysi	anova
##	0.76	0.76	0.76
##	appli	azikiw	blind
##	0.76	0.76	0.76
##	carri	come	comparison
##	0.76	0.76	0.76
##	consent	conveni	counterpart
##	0.76	0.76	0.76
##	current	determin	electr
##	0.76	0.76	0.76
##	enrol	eta	hospital,
##	0.76	0.76	0.76
##	howev	inform	intervent
##	0.76	0.76	0.76
##	interventions.	intramuscular	landmark
##	0.76	0.76	0.76

##	lasted.	mani	may,
##	0.76	0.76	0.76
##	mean?ñ?sd,	medicals/surg	nnamdi
##	0.76	0.76	0.76
##	nnewi	nnewi.	non-random
##	0.76	0.76	0.76
##	none,	obtain	p? 0.05.</th
##	0.76	0.76	0.76
##	p?=?0.01);	pactr201805003408271	pair
##	0.76	0.76	0.76
##	pan	physiotherapi	physiotherapy,
##	0.76	0.76	0.76
##	pisp	pisp.	post-inject
##	0.76	0.76	0.76
##	prove	recommend	recruit
##	0.76	0.76	0.76
##	registri	reliev	repeat
##	0.76	0.76	0.76
##	respectively,	result:	services,
##	0.76	0.76	0.76
##	session	set	sham
##	0.76	0.76	0.76
##	squared)?=?0.19.	sten	stimul
##	0.76	0.76	0.76
##	t-test.	tens;	test/experiment
##	0.76	0.76	0.76
##	transcutan	undertaken	varianc
##	0.76	0.76	0.76
##	variance.	week,	weeks)
##	0.76	0.76	0.76
##	well	wise	written
##	0.76	0.76	0.76
##	wrong	ten	particip
##	0.76	0.74	0.73
##	test		
##	0.72		

findAssocs(dtmSciatica, "assess", corlimit=0.5)

##	\$assess			
##	intern	less	(32.9	(g.
##	0.84	0.82	0.76	0.76
##	(ifis),	(odi)	(sf-36),	/s.
##	0.76	0.76	0.76	0.76
##	0.01)	0.05).	159	16th
##	0.76	0.76	0.76	0.76
##	34th	36-item	4.7	agility;
##	0.76	0.76	0.76	0.76
##	bodili	bodily,	cardiorespiratori	compon
##	0.76	0.76	0.76	0.76
##	course.	disabl	employ	fit
##	0.76	0.76	0.76	0.76
##	fitness.	fitness;	flexibility;	flexibl
##	0.76	0.76	0.76	0.76

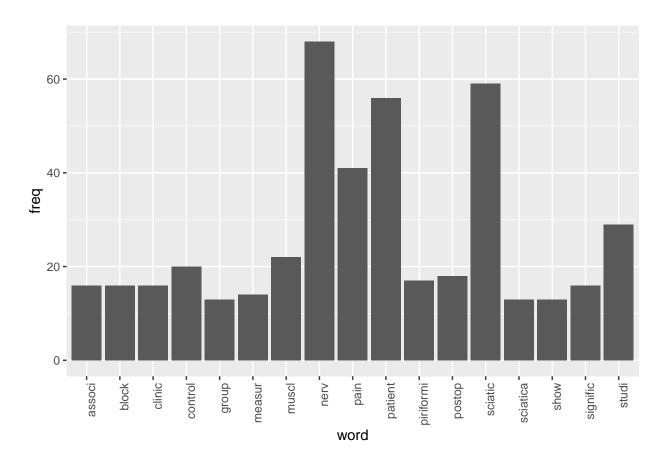
##	gestat	gestation;	greater	index
##	0.76	0.76	0.76	0.76
##	john	low-moder	medium	muscular
##	0.76	0.76	0.76	0.76
##	old).	oswestri	overal	pregnanc
##	0.76	0.76	0.76	0.76
##	pregnancy.	pregnant	questionnair	relev
##	0.76	0.76	0.76	0.76
##	scale;	self-report	son	speed-agil
##	0.76	0.76	0.76	0.76
##	stage.	strength	strength;	w.).
##	0.76	0.76	0.76	0.76
##	₩.,	women	year	analogu
##	0.76	0.76	0.76	0.73
##	explor	physic	associ	pain,
##	0.73	0.72	0.71	0.69
##	pain	wiley	scale	compris
##	0.67	0.64	0.63	0.62
##	week	health	survey	score
##	0.62	0.62	0.62	0.59
##	ltd.	need	higher	lumbar
##	0.58	0.58	0.55	0.53

findAssocs(dtmSciatica, "piriformi", corlimit=0.5)

##	<pre>\$piriformi</pre>				
##	muscle.	twenti	"nerv	(5%),	(70%),
##	0.95	0.88	0.87	0.87	0.87
##	(85%),	32:282-286,	80%	abandoned.	anat.
##	0.87	0.87	0.87	0.87	0.87
##	anatomi	anatomy;	anterior	approach	arthroplasty,
##	0.87	0.87	0.87	0.87	0.87
##	cadav	cadaver;	clarifi	clin.	disorders,
##	0.87	0.87	0.87	0.87	0.87
##	dissect	exist	fresh-frozen	identified,	identified.
##	0.87	0.87	0.87	0.87	0.87
##	implic	inc.	inferior	innerv	<pre>joint;</pre>
##	0.87	0.87	0.87	0.87	0.87
##	nerves.	origin.	periodicals,	piriformis"	plexus,
##	0.87	0.87	0.87	0.87	0.87
##	plexus;	posterior	rami	ramus	s2.
##	0.87	0.87	0.87	0.87	0.87
##	sacroiliac	said	sides,	sides.	singl
##	0.87	0.87	0.87	0.87	0.87
##	sourc	specimen	study,	suppli	surround
##	0.87	0.87	0.87	0.87	0.87
##	term	vague.	ventral	side	2018
##	0.87	0.87	0.87	0.85	0.80
##	syndrome;	sacral	superior	one	branch
##	0.80	0.75	0.75	0.73	0.67
##	gluteal	common	iatrogen	studied.	origin
##	0.65	0.64	0.64	0.64	0.62
##	thus	nerve,	descript	specifically,	trace
##	0.57	0.57	0.57	0.57	0.57

```
## proxim respectively. therefore, literatur ## 0.57 0.57 0.57 0.56
```

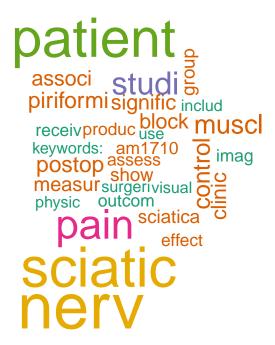
```
wf <- data.frame(word=names(freq), freq=freq)
p <- ggplot(subset(wf, freq>12), aes(word, freq))
p <- p + geom_bar(stat= 'identity')
p <- p + theme(axis.text.x=element_text(angle=90, hjust=1))
p</pre>
```



wordcloud(names(freq), freq, min.freq=10,colors=brewer.pal(3,'Dark2'))

```
sciatic
muscl nerv
outcom imag
group physic includ
clinic of effect includ
receiv signific
result
am1710 measur
includ
clinic of effect includ
clinic of effect includ
clinic of effect includ
receiv signific
result
am1710 measur
includ
clinic of effect includ
clinic of e
```

wordcloud(names(freq), freq, max.words=30,colors=brewer.pal(6,'Dark2'))



The above stemmed the corpus, this will lemmatize the original csv file

}

setwd('../')

and add the field to the table and write out to csv, followed by plot the word count frequencies that were lemmatized and the word clouds

```
library(textstem)

lemma <- lemmatize_strings(auto$abstract, dictionary=lexicon::hash_lemmas)

Lemma <- as.data.frame(lemma)
Lemma <- cbind(Lemma, auto)

colnames(Lemma) <- c('lemmatizedAbstract', 'abstract', 'source')

write.csv(Lemma, 'LemmatizedSciatica.csv', row.names=FALSE)

dir.create('./Sciatica-Lemma')

ea <- as.character(Lemma$lemmatizedAbstract)
setwd('./Sciatica-Lemma')

for (j in 1:length(ea)){
    write(ea[j], paste(paste('EAL',j, sep='.'), '.txt', sep=''))</pre>
```

```
library(tm)
library(SnowballC)
library(wordcloud)
library(ggplot2)
Sciatica <- Corpus(DirSource("Sciatica-Lemma"))</pre>
Sciatica
## <<SimpleCorpus>>
## Metadata: corpus specific: 1, document level (indexed): 0
## Content: documents: 20
#Sciatica <- tm_map(Sciatica, removePunctuation)</pre>
#Sciatica <- tm_map(Sciatica, removeNumbers)</pre>
Sciatica <- tm_map(Sciatica, tolower)</pre>
Sciatica <- tm_map(Sciatica, removeWords, stopwords("english"))</pre>
Sciatica <- tm_map(Sciatica, stripWhitespace)</pre>
dtmSciatica <- DocumentTermMatrix(Sciatica)</pre>
dtmSciatica
## <<DocumentTermMatrix (documents: 20, terms: 1425)>>
## Non-/sparse entries: 2238/26262
## Sparsity
## Maximal term length: 21
## Weighting
                   : term frequency (tf)
freq <- colSums(as.matrix(dtmSciatica))</pre>
FREQ <- data.frame(freq)</pre>
ord <- order(freq, decreasing=TRUE)</pre>
freq[head(ord, 25)]
##
           nerve
                        sciatic
                                       patient
                                                        pain
                                                                      study
##
                             60
                                                                          29
              69
                                            55
                                                           41
##
          muscle
                        control
                                   piriformis postoperative
                                                                         ca8
##
              22
                             21
                                           18
                                                           17
                                                                         17
##
                            low
                                         block
                                                    clinical
                                                                   sciatica
           group
##
              16
                             16
                                            16
                                                           14
                                                     produce ultrasound
##
            show
                            can
                                          much
                                                           12
##
              13
                             12
                                           12
                                                                          12
##
             tka
                      associate
                                        am1710 conclusion:
                                                                    include
##
              12
                             12
                                            12
                                                           11
                                                                          11
piriformis <- as.data.frame(findAssocs(dtmSciatica, "piriformis", corlimit=0.70))</pre>
ultrasound <- as.data.frame(findAssocs(dtmSciatica, "ultrasound", corlimit=0.5))</pre>
```

```
am1710 <- as.data.frame(findAssocs(dtmSciatica, "am1710", corlimit=0.55))
ca8 <- as.data.frame(findAssocs(dtmSciatica, "ca8", corlimit=0.85))
piriformis</pre>
```

```
piriformis
                        0.89
## 2019.?
## 282
                        0.89
## 286,
                        0.89
## 32:
                        0.89
## abandon.
                        0.89
## anat.
                        0.89
## anatomy
                        0.89
## anatomy;
                        0.89
## anterior
                        0.89
## approach
                        0.89
## arthroplasty,
                        0.89
## cadaver
                        0.89
## cadaver;
                        0.89
## clarify
                        0.89
## clin.
                        0.89
## description
                        0.89
## disorder,
                        0.89
## dissection
                        0.89
## exist
                        0.89
## freeze
                        0.89
## fresh
                        0.89
## identify,
                        0.89
## identify.
                        0.89
## implicate
                        0.89
## inc.
                        0.89
## inferior
                        0.89
## innervate
                        0.89
## innervation
                        0.89
## joint;
                        0.89
## majority
                        0.89
## origin.
                        0.89
## periodical,
                        0.89
## plexus,
                        0.89
## plexus;
                        0.89
## posterior
                        0.89
## proximally
                        0.89
## rami
                        0.89
## ramus
                        0.89
## s2.
                        0.89
## sacroiliac
                        0.89
                        0.89
## say
## side,
                        0.89
## side.
                        0.89
## source
                        0.89
## specimen
                        0.89
                        0.89
## study,
```

##	supply	0.89
##	surround	0.89
##	term	0.89
##	vague.	0.89
##	ventral	0.89
##	twenty	0.87
##	side	0.85
##	2018	0.80
##	origin	0.80
##	sacral	0.77
##	superior	0.77
##	muscle.	0.71

ultrasound

##		${\tt ultrasound}$
##	femoris,	0.72
##	medial	0.72
##	soleus	0.72
##	${\tt ultrasonography}$	0.72
##	july	0.72
##	contraction	0.70
##	area	0.68
##	bicep	0.68
##	change	0.68
##	low	0.68
##	ratio	0.65
##	2017.	0.65
##	cross	0.64
##	sectional	0.64
##	evaluate	0.63
##	intensity	0.58
##	muscle	0.57
##	femoris	0.57
##	injection,	0.57
##	background:	0.55
##	copyright?	0.53
##	elsevier	0.53
##	<pre>pain;</pre>	0.53
##	structure	0.53
##	guide	0.53
##	image	0.52
##	measure	0.50
##	maximum	0.50

am1710

##		am1710
##	activation	1.00
##	adjuvant	1.00
##	agent	1.00
##	agonist,	1.00
##	allodynia	1.00

##	antagonist	1.00
##	antiallodynic	1.00
##	antinociceptive	1.00
##	appropriate	1.00
##	attenuate	1.00
##	behave	1.00
##	benzo	1.00
##	broad	1.00
##	camp	1.00
##	cannabilactone	1.00
##	cannabinoid	1.00
##	cb1	1.00
##	cb2	1.00
##	cfa	1.00
##	characterize.	1.00
##	chromen	1.00
##	contrast,	1.00
##	day,	1.00
##	day?	1.00
##	delay	1.00
##	dependence.	1.00
##	development	1.00
##	dimethyl	1.00
##	dose	1.00
##	efficacy	1.00
##	embryonic	1.00
##	endure	1.00
##	extracellular	1.00
##	forskolin	1.00
##	freund's	1.00
##	gabapentin	1.00
##	hcb2.	1.00
##	hek	1.00
##	heptyl	1.00
##	hydroxy	1.00
##	incompletely	1.00
##		1.00
##	inhibition	1.00
##	kg,	1.00
##	kidney	1.00
##	kinase	1.00
##	ligation	1.00
##	mcb2	1.00
##	mcb2.	1.00
##	mechanical	1.00
##	mediate	1.00
##	methoxy	1.00
##	model.	1.00
##	modest	1.00
##	mouse	1.00
##	mouse,	1.00
##	mouse.	1.00
##	paclitaxel	1.00
##	pathway	1.00

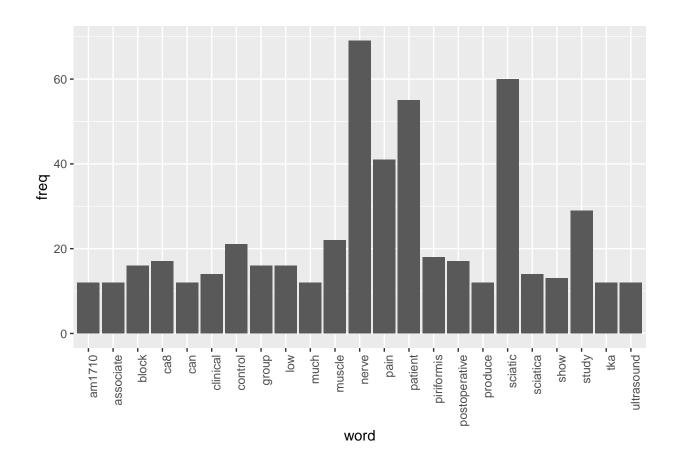
##	nhognhorylation	1.00
##	phosphorylation precipitate	1.00
	production	1.00
	profile	1.00
##	-	1.00
	= = =	1.00
##	psnl	1.00
##	,	1.00
	rodent	1.00
	similarly,	1.00
	species	1.00
	spectrum	1.00
	stably	1.00
	stimulate	1.00
	suppress	1.00
	sustain	1.00
	tetrahydrocannabinol,	1.00
	tolerance	1.00
	tolerant	1.00
	translation.	1.00
	underlie	1.00
	vitro,	1.00
	vivo,	1.00
	vivo.	1.00
##	withdrawal	1.00
##	signal	0.94
##		0.89
##	-	0.89
##		0.81
##		0.79
##	suggest	0.70
##	unwanted	0.69
##	know	0.69
##	chemotherapy	0.69
##	neuropathic	0.69
##	partial	0.69
##	establish	0.69
##	express	0.69
##	human	0.69
##	inhibit	0.69
##	observe	0.69
##	therapeutic	0.69
##	whether	0.69

ca8

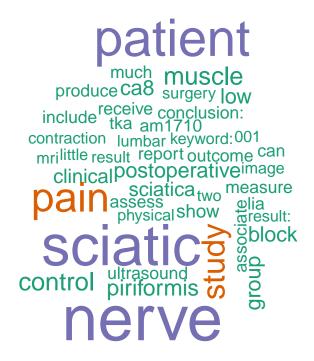
##		ca8
##	1,697bp	1.00
##	201	1.00
##	204	1.00
##	204.	1.00
##	204c	1.00
##	204g	1.00
##	3'utr	1.00

##	aav8	1.00
##	able	1.00
##	advance	1.00
##	advillin	1.00
##	allele	1.00
##	allosteric	1.00
##	also	1.00
##	anhydrase	1.00
##	ataxia,	1.00
##	barely	1.00
##	ca++	1.00
##	calcium	1.00
	carbonic	1.00
##	cell,	1.00
	cell.	1.00
	cerebellar	1.00
##	cis	1.00
	construct	1.00
	critical	1.00
	cryptic	1.00
	deliver	1.00
	derive	1.00
	develop	1.00
	dorsal	1.00
	drg	1.00
	eqtl	1.00
	eqtl.	1.00
	evade	1.00
	excitability	1.00
	excitability.	1.00
##	exclusively	1.00
##	exon	1.00
	explain	1.00
## ##	1	1.00
##	extent flag	1.00
##	-	1.00
##	g., ganglion	1.00
	gene	1.00
##	gene	1.00
	glial	1.00
	greatly	1.00
##	•	1.00
##	homozygote	1.00
##	homozygous	1.00
##	hyperalgesic	1.00
##	ihc	1.00
##	immunohistochemistry	
##	impact	1.00
##	imply	1.00
##	- *	1.00
	inositol	1.00
##		1.00
##	itpr1	1.00
	-	

```
## lesser
                       1.00
## murine
                       1.00
## mutation
                       1.00
## naturally
                       1.00
## nbl
                       1.00
## neural
                       1.00
## neuronal
                       1.00
## nociception
                      1.00
## null
                       1.00
## occur
                       1.00
## pathway,
                       1.00
## peptide
                       1.00
## phenotype.
                       1.00
## pitpr1
                       1.00
## population,
                       1.00
## predominantly
                       1.00
## previously
                       1.00
## question,
                       1.00
## release,
                       1.00
## report,
                       1.00
## restrict
                       1.00
## reversion
                       1.00
## rs6471859
                       1.00
## rs6471859,
                       1.00
## splice
                       1.00
## splice,
                       1.00
## stable
                       1.00
## synaptic
                       1.00
## tissue
                       1.00
## transcript
                       1.00
## transduce
                       1.00
## transfer
                       1.00
## trisphosphate
                       1.00
## truncate
                       1.00
## viral
                       1.00
## vitro.
                       1.00
## vivo
                       1.00
## function
                       0.96
## regulate
                       0.95
## response
                       0.89
## cell
                       0.87
                       0.86
## produce
wf <- data.frame(word=names(freq), freq=freq)</pre>
p <- ggplot(subset(wf, freq>11), aes(word, freq))
p <- p + geom_bar(stat= 'identity')</pre>
p <- p + theme(axis.text.x=element_text(angle=90, hjust=1))</pre>
```



wordcloud(names(freq), freq, min.freq=10,colors=brewer.pal(3,'Dark2'))



wordcloud(names(freq), freq, max.words=40,colors=brewer.pal(6,'Dark2'))

