Neural Net Documentation

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Setup

Following packages were used in this script:

```
# loading packages
library(keras)
library(gutenbergr)
library(dplyr)
library(tensorflow)
library(tidyr)
library(stringr)
library(tidytext)
library(udpipe)
library(sampling)
```

Get data

This is the essential step for setting up the neural net. These functions include the sampling procedure from the *gutenbergr* library

```
sampling_books <- function(seed=1234, n=20){
  # sample n books from the whole library
  set.seed(seed)
  gutenberg_works() %>%
    # select works with title
    dplyr::filter(!is.na(title)) %>%
    # set the sample sitze
    sample_n(n) %>%
    # set a special download link
    gutenberg_download(
      mirror = "http://mirrors.xmission.com/gutenberg/")
}

set_up_books <- function(n_books=4, seed=1992){
  # initial book sample
  books <- sampling_books(n=n_books, seed=seed)
  by_chapter <- books %>%
    group_by(gutenberg_id) %>%
    # split in chapters
    mutate(chapter = cumsum(str_detect(text, regex("^chapter ", ignore_case = TRUE)))) %>%
    ungroup() %>%
    # exclude books without chapters
    dplyr::filter(chapter > 0)
  return(by_chapter)
}
```

```
shorten_titles <- function(titles){
  # shorten very long book titles by setting
  # a subset of characters of the first line
  # of the title
  sub_inds <- titles %>%
    regexpr(pattern="\\n|\\r")-1
  sub_inds[sub_inds<0] <- nchar(titles)[sub_inds<0]
  titles %>%
    substr(1,sub_inds)
}

get_titles <- function(x, n_books){
  # get the sampled gutenberg_ids
  unique_ids <- x %>%
    select(gutenberg_id) %>%
    unique() %>% unlist()
  # get the titles
  titles <- gutenberg_works() %>%
    dplyr::filter(gutenberg_id %in% unique_ids) %>%
    select(gutenberg_id, title, author) %>%
    mutate(title=shorten_titles(title))
  # get the number of gutenberg ids
  len <- nrow(titles)
  if(n_books!=len) warning(paste("--- ",n_books-len,
                                 " books have 0 chapters --- "))
  # the output as a list
  ret <- list(
    titles=titles,
    len=len
  )
  return(ret)
}

append_by_chapter <- function(x=by_chapter, n_books, seed_index=1){
  # append the books matrix until
  # we get the desired number of books n_books
  titles <- get_titles(x, n_books)
  n <- titles$len
  while (n<n_books) {
    book2add <- sampling_books(n=1, seed=seed_index)
    by_chapter_add <- book2add %>%
      group_by(gutenberg_id) %>%
      # split in chapters
      mutate(chapter = cumsum(str_detect(text, regex("^chapter ", ignore_case = TRUE)))) %>%
      ungroup() %>%
      # exclude books without chapters
      dplyr::filter(chapter > 2)
    titles2add <- get_titles(by_chapter_add, 1)
    # adding the book to by_chapter if there are chapters in the
    # book plus it is not in the data already
    if (titles2add$len==1) if(!titles2add$titles$gutenberg_id%in%titles$titles$gutenberg_id) {
      x <- bind_rows(x, by_chapter_add)
    }
```

```
    n<-get_titles(x, n)$len
    seed_index <- seed_index+1
  }
  return(x)
}

exclude_stop_words <- function(x){
  # unite chapter and document title
  by_chapter_word <- x %>%
    unite(document, gutenberg_id, chapter) %>%
    # split into words
    unnest_tokens(word, text)
  # import tibble stop words
  data(stop_words)
  # find document-word counts
  word_counts <- by_chapter_word %>%
    # exclude stop words
    anti_join(stop_words) %>%
    # count each word by chapter
    count(document, word, sort = TRUE) %>%
    ungroup()
  return(word_counts)
}

convert_to_dtm <- function(x, minfq = 2){
  # get into a format lda can handle
  chapters_dtm <- x %>%
    select(doc_id=document, term=word, freq=n) %>%
    document_term_matrix() %>%
    # reduce by low frequencies
    dtm_remove_lowfreq(minfreq = minfq)
  return(chapters_dtm)
}


convert_to_dtm_2 <- function(x, n=n, minfq = 2, top=10000){
  # get into a format lda can handle
  chapters_dtm <- x %>%
    select(doc_id=document, term=word, freq=n) %>%
    document_term_matrix() %>%
    # reduce by low frequencies
    dtm_remove_tfidf(top=top)
  return(chapters_dtm)
}

# convert x matrix into a form such that it can be used for tensorflow
adjust_tensor_format <- function(x){
  x_chapters <- apply(x, 1, function(x) as.matrix(x)) %>% t()
  topics <- x %>% rownames() %>% as_tibble() %>%
    separate(value, c("gutenberg_id", "chapter"), sep = "_", convert = TRUE) %>%
    select(gutenberg_id) %>%
    # split joint name of book and chapter
    as.matrix %>% as.factor() %>% as.integer()
```

```
  # one hot encoding for the chapters (y)
  topics_categorical <- topics %>% -1 %>%
    to_categorical()
  ret <- list(
    x=x_chapters,
    y=topics_categorical,
    topics=topics
  )
  return(ret)
}
```

Sample corpus

Now we can use all these functions to get to the initial corpus sample. In this example 6 books are choosen.

```
n_books <- 6
by_chapter <- set_up_books(n_books=n_books, seed=222)
get_titles(by_chapter, n_books)
```

```
## Warning in get_titles(by_chapter, n_books): --- 1 books have 0 chapters ---
```

```
## $titles
## # A tibble: 5 x 3
##   gutenberg_id title                                    author
##          <int> <chr>                                    <chr>
## 1           11 Alice's Adventures in Wonderland         Carroll, Lewis
## 2         3096 Beatrice                                 Haggard, H. Rider~
## 3        25603 Detailed Minutiae of Soldier life in the~ McCarthy, Carlton
## 4        47402 Along Alaska's Great River              Schwatka, Frederi~
## 5        49675 Hawkins Electrical Guide v. 5 (of 10)   Hawkins, N. (Nehe~
##
## $len
## [1] 5
```

The function set_up_books() returns a warning that one book seems to consist of only one chapter. In order to get a copus consisting out of 6 books, the function append_by_chaper() is used, which fills up the corpus to the desired number of books.

```
appended_by_chapter <- append_by_chapter(x=by_chapter, n_books = n_books)
word_counts <- exclude_stop_words(appended_by_chapter)
```

```
## Joining, by = "word"
```

In table 1 the sampled titles for the book sample with the seed 222 are displayed. It appears through the function append_by:chapter() one book was added, called "My Novel" — Volume 04".

```
titles <- get_titles(appended_by_chapter, n_books)
titles$titles %>% stargazer(summary=FALSE, font.size = "footnotesize",
                            header=FALSE, title="Book-titles", rownames=FALSE,
                            label="titles:6books")
```

Reduction of the dimensionality

In the set up we have another parameter to adjust. The function *convert_to_dtm* takes the parameter *minfq*, which is used to reduce the "bag of words" (i.e. dimensionality). *minfq* is the minimum frequency for the

Table 1: Book-titles

gutenberg_id	title	author
11	Alice's Adventures in Wonderland	Carroll, Lewis
3096	Beatrice	Haggard, H. Rider (Henry Rider)
7705	"My Novel" — Volume 04	Lytton, Edward Bulwer Lytton, Baron
25603	Detailed Minutiae of Soldier life in the Army of Northern Virginia, 1861-1865	McCarthy, Carlton
47402	Along Alaska's Great River	Schwatka, Frederick
49675	Hawkins Electrical Guide v. 5 (of 10)	Hawkins, N. (Nehemiah)

bag of words dictionary. I will refer to this as "embedding". Let us set it to 2 in this case, meaning that we include a word only if the frequency is 2 or more.

```
chapters_dtm <- convert_to_dtm(word_counts, minfq=2)
adjusted_format <- adjust_tensor_format((chapters_dtm))
ncol(chapters_dtm)
```

```
## [1] 10685
```

Let us compare it to the case if we include all words.

```
chapters_dtm_all <- convert_to_dtm(word_counts, minfq=0)
adjusted_format_all <- adjust_tensor_format((chapters_dtm))
ncol(chapters_dtm_all)
```

```
## [1] 17961
```

We also want to compare this to a reduction of the word dictionary by the tfidf. For the sake of comparison the reduction is made to the same value as used above via minfreq=2 (i.e. 10685 words).

```
chapters_dtm_tfidf <- convert_to_dtm_2(word_counts, top=10685)
adjusted_format_tfidf <- adjust_tensor_format((chapters_dtm))
ncol(chapters_dtm_tfidf)
```

```
## [1] 10685
```

Splitting and Fitting

The following function splits the sample in an manner, such that each cluster is eually to its size represented in the test data and the validation data.

```
sample_cluster_wise <- function(data, test_ratio=0.1, val_ratio=0.2, seed=1234){
  X <- data$x; y <- data$y
  cluster=data$topics
  set.seed(seed)
  {
  # setting the absolute number of observations for the sample of each cluster
  n_test <- (table(cluster)*test_ratio) %>% floor() %>%
    # use at least one observation of each cluster
    sapply(., function(x) max(x,1))
  n_val <- (table(cluster)*val_ratio) %>% floor() %>%
    # use at least one observation of each cluster
    sapply(., function(x) max(x,1))
  # function to get the correct sample indices for validation and test sample
  samp_ind <- function(i, n_list) which(cluster==i) %>% sample(n_list[i])
  test_indices <- unique(cluster) %>% sort() %>%
    sapply(function(i) samp_ind(i, n_test)) %>%
```

```
    unlist()
  val_indices <- unique(cluster) %>% sort() %>%
    sapply(function(i) samp_ind(i, n_val)) %>%
    unlist()
  }
  ret <- list(partial_x_train=X[-c(val_indices,test_indices),],
              partial_y_train=y[-c(val_indices,test_indices),],
              x_val=X[val_indices,], y_val = y[val_indices,],

              x_test=X[test_indices,], y_test = y[test_indices,])
  return(ret)
}
```

The following two functions are setting up the model and evaluate the goodness of fit.

```
# The whole model is set up and trained within this function
set_up_n_fit <- function(split, books_n=n_books){
  # starting with 64 neurons and scaling it down to 46 in the
  # mid layer turned out to be a well predicting model
  model <- keras_model_sequential() %>%
    layer_dense(units=64, activation="relu", input_shape=ncol(split$partial_x_train)) %>%
    layer_dense(units=46, activation="relu") %>%
    # we want to classify for as many categories as books
    layer_dense(units=books_n, activation="softmax")

  model %>% compile(
    optimizer="rmsprop",
    loss="categorical_crossentropy",
    metrics=c("accuracy"))

  history <- model %>% fit(
    split$partial_x_train,
    split$partial_y_train,
    # from experience the model tends to
    # overfit for more than 5 epochs
    epochs=5,
    batch_size=512,
    validation_data=list(split$x_val,split$y_val)
  )
  return(
    list(history=history,
         model=model))
}


# making a prediction on the test data and calculating the
# misspecification rate; we also want to save the true categories and the predicted ones
evaluate_model <- function(model_fit, y=split$y_test, x=split$x_test) {
  prediction <- model_fit %>% predict(x)
  pred <- apply(prediction, 1, which.max)
  true_value <- apply(y, 1, which.max)
  misspecified <- sum(!pred==true_value)/length(pred)
  ret <- list(misspecified=misspecified,
              # the function also dicloses the true and the predicted
```

```
              # values for exact evaluation, if needed
              pred=pred, true_value=true_value)
  return(ret)
}
```

Now we can start applying the model:

```
tim1 <- Sys.time()
n <- 59
results <- rep(NA,n)
for(i in 1:n){
  split <- sample_cluster_wise(adjusted_format, seed=i*2)
  results[i] <- set_up_n_fit(split) %>% .$model %>%
    evaluate_model() %>% .$misspecified
}
tim2 <- Sys.time()
```

The results of the misclassification rate over 59 splits and fits of the model are:

```
results %>% summary
```

```
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.00000 0.00000 0.00000 0.02203 0.00000 0.20000
```

```
results %>% var
```

```
## [1] 0.002092344
```

and the time:

```
# mfreq=2
(u1 <- tim2-tim1)
```

```
## Time difference of 3.646245 mins
```

We now try this procedure using the full bag of words:

```
tim1_all <- Sys.time()
n <- 59
results_all <- rep(NA,n)
for(i in 1:n){
  split <- sample_cluster_wise(adjusted_format_all, seed=i*2)
  results_all[i] <- set_up_n_fit(split) %>% .$model %>%
    evaluate_model() %>% .$misspecified
}
tim2_all <- Sys.time()
```

The results of the error rate

```
results_all %>% summary
```

```
##     Min.  1st Qu.   Median     Mean  3rd Qu.     Max.
## 0.000000 0.000000 0.000000 0.008475 0.000000 0.100000
```

```
results_all %>% var
```

```
## [1] 0.0007890123
```

and the time:

```
# mfreq=2
(u2 <- tim2_all-tim1_all)
```

```
## Time difference of 10.28905 mins
```

Now using the tfidf reduced bag of words:

```
tim1_tfidf <- Sys.time()
n <- 59
results_tfidf <- rep(NA,n)
for(i in 1:n){
  split <- sample_cluster_wise(adjusted_format_tfidf, seed=i*2)
  results_tfidf[i] <- set_up_n_fit(split) %>% .$model %>%
    evaluate_model() %>% .$misspecified
}
tim2_tfidf <- Sys.time()
```

The results of the error rate

```
results_tfidf %>% summary
```

```
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.00000 0.00000 0.00000 0.01864 0.00000 0.20000
```

```
results_tfidf %>% var
```

```
## [1] 0.001887785
```

and the time:

```
# mfreq=2
(u3 <- tim2_tfidf-tim1_tfidf)
```

```
## Time difference of 17.29859 mins
```

The following matrix summarises the results.

```
performance_matrix <- data.frame(freq2.embedding=c(results%>%mean, u1),
           all.embedding=c(results_all%>%mean, u2),
           tfidf=c(results_tfidf%>%mean, u3))
rownames(performance_matrix) <- c("missc. rate", "time")
performance_matrix %>%  stargazer(summary=FALSE, header=F)
```

Table 2:

	freq2.embedding	all.embedding	tfidf
missc. rate	0.022	0.008	0.019
time	3.646	10.289	17.299

The time for the embedding using more than frequency 2 is very short. Whereas the embedding via tfidf with the same dimensionality takes very long.