Assignment2

Hannah Cronin

2023-03-26

1. Separating the data like in the book.

```
library(keras)
original_dataset_dir <- "/Users/hannahcronin/Desktop/kaggle_original_data"
base_dir <- "/Users/hannahcronin/Desktop/kaggle_original_data"</pre>
dir.create(base_dir)
## Warning in dir.create(base_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data' already exists
train_dir <- file.path(base_dir, "train")</pre>
dir.create(train dir)
## Warning in dir.create(train_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/train' already exists
validation_dir <- file.path(base_dir, "validation")</pre>
dir.create(validation dir)
## Warning in dir.create(validation_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/validation' already exists
test_dir <- file.path(base_dir, "test")</pre>
dir.create(test_dir)
## Warning in dir.create(test_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/test' already exists
train cats dir <- file.path(train dir, "cats")
dir.create(train_cats_dir)
## Warning in dir.create(train cats dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/train/cats' already exists
train_dogs_dir <- file.path(train_dir, "dogs")</pre>
dir.create(train_dogs_dir)
## Warning in dir.create(train_dogs_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/train/dogs' already exists
validation cats dir <- file.path(validation dir, "cats")</pre>
dir.create(validation_cats_dir)
## Warning in dir.create(validation_cats_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/validation/cats' already exists
validation_dogs_dir <- file.path(validation_dir, "dogs")</pre>
dir.create(validation_dogs_dir)
## Warning in dir.create(validation_dogs_dir): '/Users/hannahcronin/Desktop/
## kaggle original data/validation/dogs' already exists
test_cats_dir <- file.path(test_dir, "cats")</pre>
dir.create(test_cats_dir)
## Warning in dir.create(test_cats_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/test/cats' already exists
```

file.path(train_cats_dir))

```
test_dogs_dir <- file.path(test_dir, "dogs")
dir.create(test_dogs_dir)

## Warning in dir.create(test_dogs_dir): '/Users/hannahcronin/Desktop/
## kaggle_original_data/test/dogs' already exists

fnames <- paste0("cat.", 1:1000, ".jpg")
file.copy(file.path(train_dir, fnames),</pre>
```

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cat("total training cat images:", length(list.files(train_cats_dir)), "\n")
## total training cat images: 1000
## total training cat images: 1000
cat("total training dog images:", length(list.files(train dogs dir)), "\n")
## total training dog images: 1000
## total training dog images: 1000
cat("total validation cat images:",length(list.files(validation_cats_dir)), "\n")
## total validation cat images: 500
## total validation cat images: 500
cat("total validation dog images:",length(list.files(validation_dogs_dir)), "\n")
## total validation dog images: 500
## total validation dog images: 500
cat("total test cat images:", length(list.files(test cats dir)), "\n")
## total test cat images: 500
## total test cat images: 500
cat("total test dog images:", length(list.files(test_dogs_dir)), "\n")
```

```
## total test dog images: 500
```

total test dog images: 500

```
## Using data augmentation as model optimizer
datagen <- image_data_generator(
  rescale = 1/255,
  rotation_range = 40,
  width_shift_range = 0.2,
  height_shift_range = 0.2,
  shear_range = 0.2,
  zoom_range = 0.2,
  horizontal_flip = TRUE,
  fill_mode = "nearest"
)</pre>
```

```
model <- keras_model_sequential() %>%
 layer_conv_2d(filters = 16, kernel_size = c(3, 3), activation = "relu",
               input_shape = c(150, 150, 3)) %>%
 layer max pooling 2d(pool size = c(2, 2)) %>%
 layer\_conv\_2d(filters = 32, kernel\_size = c(3, 3), activation = "relu") %>%
 layer_max_pooling_2d(pool_size = c(2, 2)) %>%
 layer_conv_2d(filters = 64, kernel_size = c(3, 3), activation = "relu") %>%
 layer_max_pooling_2d(pool_size = c(2, 2)) %>%
 layer_conv_2d(filters = 64, kernel_size = c(3, 3), activation = "relu") %>%
 layer_max_pooling_2d(pool_size = c(2, 2)) %>%
 layer_flatten() %>%
 layer_dropout(rate = 0.5) %>%
 layer_dense(units = 512, activation = "relu") %>%
 layer_dense(units = 1, activation = "sigmoid")
model %>% compile(
 loss = "binary_crossentropy",
 optimizer = optimizer rmsprop(learning rate = 0.001),
 metrics = c("acc")
```

```
##training the data
set.seed(100)
datagen <- image_data_generator(</pre>
 rescale = 1/255,
 rotation_range = 40,
 width_shift_range = 0.2,
 height shift range = 0.2,
 shear_range = 0.2,
 zoom range = 0.2,
 horizontal flip = TRUE
test datagen <- image data generator(rescale = 1/255)
train_generator <- flow_images_from_directory(</pre>
 train dir,
 datagen,
 target size = c(150, 150),
 batch_size = 20,
 class_mode = "binary"
validation_generator <- flow_images_from_directory(</pre>
 validation_dir,
 test datagen,
  target_size = c(150, 150),
 batch size = 20,
 class_mode = "binary"
history <- model %>% fit(
 train_generator,
 steps_per_epoch = 50,
 epochs = 10,
 validation_data = validation_generator,
 validation_steps = 5
```

```
history
```

```
## Final epoch (plot to see history):
  ##
                       loss: 0.6411
  ##
                                 acc: 0.638
  ## val loss: 0.5913
  ## val_acc: 0.68
  ## This model had a validation accuracy of 0.7 and an accuracy of 0.687.
Adding more samples to training data.
 r train_dir2a <- file.path(base_dir, "train2a") dir.create(train_dir2a)
  ## Warning in dir.create(train dir2a): '/Users/hannahcronin/Desktop/ ## kaggle original data/train2a' already exists
 r train cats dir2a <- file.path(train dir2a, "cats2a") dir.create(train cats dir2a)
  ## Warning in dir.create(train cats dir2a): '/Users/hannahcronin/Desktop/ ## kaggle original data/train2a/cats2a' already exists
 r train_dogs_dir2a <- file.path(train_dir2a, "dogs2a") dir.create(train_dogs_dir2a)
  ## Warning in dir.create(train_dogs_dir2a): '/Users/hannahcronin/Desktop/ ## kaggle_original_data/train2a/dogs2a' already exists
 r fnames <- paste0("cat.", 1:1000, ".jpg") file.copy(file.path(train_dir2a, fnames), file.path(train_cats_dir2a))
                       [1] FALSE FA
 r fnames <- paste0("cat.", 2001:2500, ".jpg") file.copy(file.path(train_dir2a, fnames), file.path(train_cats_dir2a))
  ## [1] FALSE ## [13] FALSE FAL
 r fnames <- paste0("dog.", 1:1000, ".jpg") file.copy(file.path(train_dir2a, fnames), file.path(train_dogs_dir2a))
                       [1] FALSE FA
 r fnames <- paste0("dog.", 2001:2500, ".jpg") file.copy(file.path(train_dir2a, fnames), file.path(train_dogs_dir2a))
  ## [1] FALSE ## [13] FALSE FAL
 r cat("total training cat images:", length(list.files(train_cats_dir2a)), "\n")
  ## total training cat images: 1500
r ## total training cat images: 1500 cat("total training dog images:", length(list.files(train dogs dir2a)), "\n")
 ## total training dog images: 1500
 r ## total training dog images: 1500
r ## Using data augmentation as model optimizer datagen <- image_data_generator( rescale = 1/255, rotation_range = 40, width_shift_range = 40, width_s
r library(keras) model2a <- keras_model_sequential() %>% layer_conv_2d(filters = 16, kernel_size = c(3, 3), activation = "relu", inpu
 r ##training the data ## scaling training images set.seed(100) datagen <- image_data_generator( rescale = 1/255, rotation_range = 40,
 r history2a
  ## ## Final epoch (plot to see history): ##
                                                                                                                                                                                          loss: 0.645 ##
                                                                                                                                                                                                                                                                         acc: 0.631 ## val_loss: 0.7637 ## val_acc: 0.5
  r ## Accuracy decreased overall when I increased the number of training images.
  train_dir3a <- file.path(base_dir, "train3a")</pre>
  dir.create(train_dir3a)
  ## Warning in dir.create(train_dir3a): '/Users/hannahcronin/Desktop/
  ## kaggle original data/train3a' already exists
  train_cats_dir3a <- file.path(train_dir3a, "cats3a")</pre>
  dir.create(train cats dir3a)
  ## Warning in dir.create(train_cats_dir3a): '/Users/hannahcronin/Desktop/
  ## kaggle original data/train3a/cats3a' already exists
  train_dogs_dir3a <- file.path(train_dir3a, "dogs3a")</pre>
  dir.create(train_dogs_dir3a)
  ## Warning in dir.create(train_dogs_dir3a): '/Users/hannahcronin/Desktop/
   ## kaggle_original_data/train3a/dogs3a' already exists
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```
fnames <- paste0("dog.", 1:1000, ".jpg")
file.copy(file.path(train_dir3a, fnames),
    file.path(train_dogs_dir3a))</pre>
```

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[1] FALSE FA
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               [445] FALSE 
               [457] FALSE FALSE
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              [589] FALSE FALSE
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               [613] FALSE FALSE
             [625] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
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               [637] FALSE FALSE
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               [661] FALSE FALSE
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             [685] FALSE FALSE
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               [697] FALSE 
               [709] FALSE FALSE
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              [721] FALSE FALSE
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               [817] FALSE FALSE
              [829] FALSE FALSE
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               [841] FALSE FALSE
                [853] FALSE FALSE
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               [865] FALSE FALSE
             [877] FALSE FALSE
               [889] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
               [901] FALSE FALSE
               [913] FALSE FALSE
            [925] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
```

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## [937] FALSE FALSE
     [949] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [961] FALSE FALSE
## [973] FALSE FALSE
## [985] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [997] FALSE FALSE FALSE
fnames <- paste0("dog.", 2001:2200, ".jpg")</pre>
file.copy(file.path(train_dir3a, fnames),
               file.path(train_dogs_dir3a))
    [1] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [13] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [25] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [37] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
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## [157] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [169] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [181] FALSE FALSE
## [193] FALSE FALSE FALSE FALSE FALSE FALSE
cat("total training cat images:", length(list.files(train_cats_dir3a)), "\n")
## total training cat images: 1300
## total training cat images: 1300
cat("total training dog images:", length(list.files(train_dogs_dir3a)), "\n")
## total training dog images: 1300
## total training dog images: 1300
## Using data augmentation as model optimizer
datagen <- image_data_generator(</pre>
  rescale = 1/255,
  rotation range = 40.
  width shift range = 0.2,
  height shift range = 0.2,
  shear_range = 0.2,
  zoom_range = 0.2,
  horizontal flip = TRUE,
  fill_mode = "nearest"
library(keras)
model3a <- keras model sequential() %>%
  layer_conv_2d(filters = 16, kernel_size = c(3, 3), activation = "relu",
                        input_shape = c(150, 150, 3)) %>%
  layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer conv 2d(filters = 32, kernel size = c(3, 3), activation = "relu") %>%
   layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer conv 2d(filters = 64, kernel size = c(3, 3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer_conv_2d(filters = 64, kernel_size = c(3, 3), activation = "relu") %>%
   layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer flatten() %>%
  layer_dropout(rate = 0.5) %>%
  layer_dense(units = 512, activation = "relu") %>%
  layer_dense(units = 1, activation = "sigmoid")
model3a %>% compile(
   loss = "binary_crossentropy",
  optimizer = optimizer_rmsprop(learning_rate = 0.001),
   metrics = c("acc")
```

```
##training the data
## scaling training images
set.seed(100)
datagen <- image_data_generator(</pre>
 rescale = 1/255,
 rotation_range = 40,
 width_shift_range = 0.2,
 height_shift_range = 0.2,
 shear range = 0.2,
 zoom_range = 0.2,
 horizontal_flip = TRUE
## scaling testing images
test_datagen <- image_data_generator(rescale = 1/255)</pre>
train_generator3a <- flow_images_from_directory(</pre>
 train_dir3a,
 datagen,
 target size = c(150, 150),
 batch size = 20,
 class_mode = "binary"
validation_generator <- flow_images_from_directory(</pre>
 validation_dir,
 test datagen,
 target_size = c(150, 150),
 batch_size = 20,
 class_mode = "binary"
history3a <- model3a %>% fit(
 train_generator3a,
 steps_per_epoch = 50,
 epochs = 10.
 validation_data = validation_generator,
 validation steps = 5
```

history3a

```
##
## Final epoch (plot to see history):
## loss: 0.6678
## acc: 0.604
## val_loss: 0.7189
## val_acc: 0.54
```

Validation is 0.75 and training accuracy is 0.69. I think with more epochs and more steps per epoch, this numb er could rise. I've had to lower the number of steps and epochs for performance reasons. Initially, each run was taking ~15 minutes for my R browser to complete. Given the size of the file, I did not want the document to take over an hour to knit/export (as it may overheat my computer).

Using a pre-trained network 1)

```
Larger training dataset
```

```
r train_dir2b <- file.path(base_dir2, "train2b") datagen1b <- image_data_generator(rescale = 1/255) batch_size <- 20 extract_features
r model2b <- keras_model_sequential() %>% conv_base %>% layer_flatten() %>% layer_dense(units = 256, activation = "relu") %>% layer_c
r set.seed(100) train_datagen2b = image_data_generator( rescale = 1/255, rotation_range = 40, width_shift_range = 0.2, height_shift_r
r history2b
## ## Final epoch (plot to see history): ## loss: 0.5735 ## acc: 0.678 ## val_loss: 0.5682 ## val_acc: 0.74
```

r ## Performance is slightly poorer with training data samples of 1500 each for cats & dogs. The model does seem to have some more va

Trying to improve model

```
train_dir3b <- file.path(base_dir2, "train3b")
dir.create(train_dir3b)

## Warning in dir.create(train_dir3b): '/Users/hannahcronin/Desktop/
## cats_and_dogs_small/train3b' already exists</pre>
```

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[1] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
  [13] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
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## [721] FALSE FALSE
## [733] FALSE FALSE
## [745] FALSE FALSE
## [757] FALSE FALSE
## [769] FALSE FALSE
## [781] FALSE FALSE
## [793] FALSE FALSE FALSE FALSE FALSE FALSE
```

```
train_dir3b <- file.path(base_dir2, "train3b")</pre>
datagen1b <- image_data_generator(rescale = 1/255)</pre>
batch size <- 20
extract_features <- function(directory, sample_count) {</pre>
 features <- array(0, dim = c(sample_count, 4, 4, 512))</pre>
  labels <- array(0, dim = c(sample_count))</pre>
  generator <- flow_images_from_directory(</pre>
   directory = directory,
   generator = datagen,
   target_size = c(150, 150),
   batch_size = batch_size,
   class mode = "binary"
 while(TRUE) {
   batch <- generator_next(generator)</pre>
   inputs_batch <- batch[[1]]</pre>
   labels batch <- batch[[2]]
   features_batch <- conv_base %>% predict(inputs_batch)
   index_range <- ((i * batch_size)+1):((i + 1) * batch_size)</pre>
   features[index_range,,,] <- features_batch</pre>
   labels[index_range] <- labels_batch</pre>
   i <- i + 1
   if (i * batch_size >= sample_count)
      break
 list(
    features = features,
    labels = labels
  )
train3b <- extract_features(train_dir3b, 1600)</pre>
train3b$features <- reshape_features(train$features)</pre>
```

```
model3b <- keras_model_sequential() %>%
  conv_base %>%
  layer_flatten() %>%
  layer_dense(units = 256, activation = "relu") %>%
  layer_dense(units = 1, activation = "sigmoid")
```

```
set.seed(100)
train_datagen3b = image_data_generator(
 rescale = 1/255,
 rotation_range = 40,
 width shift range = 0.2,
 height_shift_range = 0.2,
 shear range = 0.2,
 zoom_range = 0.2,
 horizontal_flip = TRUE,
 fill_mode = "nearest"
test datagen3b <- image data generator(rescale = 1/255)
train_generator3b <- flow_images_from_directory(</pre>
 train_dir3b,
 train datagen3b,
 target_size = c(150, 150),
 batch_size = 20,
 class mode = "binary"
validation_generator1b <- flow_images_from_directory(</pre>
 validation_dir1b,
 test_datagen1b,
  target_size = c(150, 150),
 batch size = 20,
 class_mode = "binary"
model3b %>% compile(
 loss = "binary_crossentropy",
 optimizer = optimizer_rmsprop(learning_rate = 0.001),
 metrics = c("accuracy")
history3b <- model %>% fit(
 train_generator3b,
 steps_per_epoch = 50,
 epochs = 10,
 validation_data = validation_generator1b,
  validation_steps = 5
```

history3b

```
##
## Final epoch (plot to see history):
## loss: 0.56
## acc: 0.724
## val_loss: 0.4988
## val_acc: 0.77
```

Performance increased with less data in the training samples. Accuracy was 0.79 and validation accuracy was 0.76. This suggests their may be some overfitting due to the smaller size of the sample data, but the accuracies we re not consistently higher than validation accuracy.

Conclusion, with the pretrained covnet, the models had overall better performance. For the model's built from scratch, I tried to use less training data and got worse results, used more training data and also got worse results from the initial run. However, slightly more data (not as much as run 2) procured the best results of the runs.

When it came to the pretrained covnet, there was better results immmediately. Again, more training data resulted in worse results. However, less training data gave the model better accuracy.

I am wondering if hyperparameters or if more epochs/steps per epoch were done in my "built" from scratch model that these results would be replicated.

Overall, less training data with pretrained worked better and a data samples from 1000-1300 worked best for my built from scratch model.