Assignment 2

Jacob Fabian

2023-03-26

```
library(reticulate)
use_condaenv('r-reticulate')
library(tensorflow)
library(keras)
original_dataset_dir <- "~/Downloads/dogs-vs-cats"</pre>
base_dir <- "~/Downloads/cats_and_dogs_small"</pre>
dir.create(base_dir)
## Warning in dir.create(base_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small' already exists
train_dir <- file.path(base_dir, "train")</pre>
dir.create(train_dir)
## Warning in dir.create(train_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/train' already exists
validation_dir <- file.path(base_dir, "validation")</pre>
dir.create(validation_dir)
## Warning in dir.create(validation_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/validation' already exists
test_dir <- file.path(base_dir, "test")</pre>
dir.create(test_dir)
## Warning in dir.create(test_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/test' already exists
train_cats_dir <- file.path(train_dir, "cats")</pre>
dir.create(train_cats_dir)
## Warning in dir.create(train_cats_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/train/cats' already exists
```

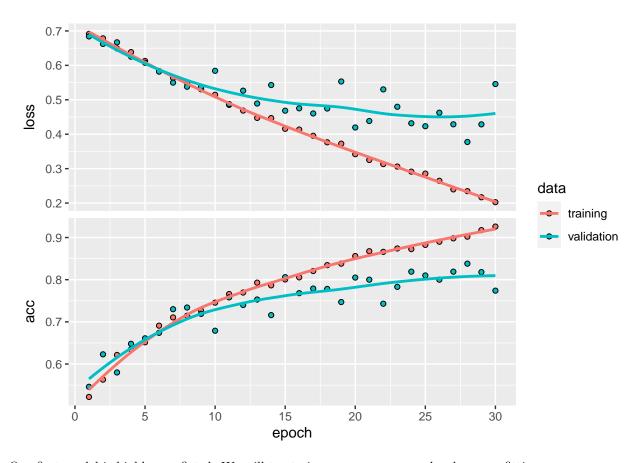
```
train_dogs_dir <- file.path(train_dir, "dogs")</pre>
dir.create(train_dogs_dir)
## Warning in dir.create(train_dogs_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/train/dogs' already exists
validation_cats_dir <- file.path(validation_dir, "cats")</pre>
dir.create(validation_cats_dir)
## Warning in dir.create(validation_cats_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/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/jacobfabian/Downloads/cats_and_dogs_small/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/jacobfabian/Downloads/cats_and_dogs_small/test/cats' already exists
test_dogs_dir <- file.path(test_dir, "dogs")</pre>
dir.create(test_dogs_dir)
## Warning in dir.create(test_dogs_dir):
## '/Users/jacobfabian/Downloads/cats_and_dogs_small/test/dogs' already exists
fnames <- paste0("cat.", 1:1000, ".jpg")</pre>
file.copy(file.path(original_dataset_dir, fnames),
          file.path(train_cats_dir))
fnames <- paste0("cat.", 1001:1500, ".jpg")</pre>
file.copy(file.path(original_dataset_dir, fnames),
          file.path(validation_cats_dir))
fnames <- paste0("cat.", 1501:2000, ".jpg")</pre>
file.copy(file.path(original_dataset_dir, fnames),
          file.path(test_cats_dir))
fnames <- paste0("dog.", 1:1000, ".jpg")</pre>
file.copy(file.path(original_dataset_dir, fnames),
          file.path(train_dogs_dir))
fnames <- paste0("dog.", 1001:1500, ".jpg")</pre>
```

```
file.copy(file.path(original_dataset_dir, fnames),
          file.path(validation_dogs_dir))
fnames <- paste0("dog.", 1501:2000, ".jpg")</pre>
file.copy(file.path(original_dataset_dir, fnames),
          file.path(test_dogs_dir))
library(keras)
library(tensorflow)
model <- keras_model_sequential() %>%
  layer_conv_2d(filters = 32, 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 = 64, kernel_size = c(3, 3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer_conv_2d(filters = 128, kernel_size = c(3, 3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer_conv_2d(filters = 128, kernel_size = c(3, 3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer flatten() %>%
  layer_dense(units = 512, activation = "relu") %>%
  layer_dense(units = 1, activation = "sigmoid")
```

summary(model)

```
## Model: "sequential"
## Layer (type)
                                Output Shape
                                                          Param #
## -----
## conv2d 3 (Conv2D)
                                (None, 148, 148, 32)
                                                           896
## max_pooling2d_3 (MaxPooling2D)
                                (None, 74, 74, 32)
## conv2d_2 (Conv2D)
                                (None, 72, 72, 64)
                                                           18496
                                (None, 36, 36, 64)
## max_pooling2d_2 (MaxPooling2D)
## conv2d_1 (Conv2D)
                                (None, 34, 34, 128)
                                                           73856
## max_pooling2d_1 (MaxPooling2D)
                                (None, 17, 17, 128)
## conv2d (Conv2D)
                                 (None, 15, 15, 128)
                                                           147584
## max_pooling2d (MaxPooling2D)
                                (None, 7, 7, 128)
                                (None, 6272)
## flatten (Flatten)
                                                           0
## dense_1 (Dense)
                                (None, 512)
                                                           3211776
## dense (Dense)
                                (None, 1)
                                                           513
## Total params: 3,453,121
## Trainable params: 3,453,121
## Non-trainable params: 0
model %>% compile(
 loss = "binary_crossentropy",
 optimizer = optimizer rmsprop(learning rate = 1e-4),
 metrics = c("acc")
```

```
train_datagen <- image_data_generator(rescale = 1/255)</pre>
validation_datagen <- image_data_generator(rescale = 1/255)</pre>
train_generator <- flow_images_from_directory(</pre>
  train_dir,
 train_datagen,
 target_size = c(150, 150),
batch_size = 20,
 class_mode = "binary"
validation_generator <- flow_images_from_directory(</pre>
  validation_dir,
 validation_datagen,
 target_size = c(150, 150),
 batch_size = 20,
  class_mode = "binary"
batch<- generator_next(train_generator)</pre>
str(batch)
## List of 2
## $ : num [1:20, 1:150, 1:150, 1:3] 0.4353 0.0902 0.4314 0.4824 0.1882 ...
## $ : num [1:20(1d)] 1 0 0 0 1 0 0 1 0 1 ...
history <- model %>% fit(
 train_generator,
 steps_per_epoch = 100,
 epochs = 30,
 validation_data = validation_generator,
  validation_steps = 50
plot(history)
```



Our first model is highly overfitted. We will try to increase accuracy and reduce overfitting.

Question 1:

The following code improves upon the base model by adding a dropout layer. This should help the model to stop overfitting to try to improve preformance.

```
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
fnames <- list.files(file.path(train_dir,"cats"), full.names = T)</pre>
img_path <- fnames[[round(runif(1,1,length(fnames)))]]</pre>
img <- image_load(img_path, target_size = c(150,150))</pre>
img_array <- image_to_array(img)</pre>
img_array <- array_reshape(img_array, c(1,150,150,3))</pre>
augmentation_generator <- flow_images_from_data(</pre>
    img array,
    generator = datagen,
    batch size = 1
```

```
op <- par(mfrow=c(2,2), pty="s", mar=c(1,0,.1,0))
for (i in 1:4) {
  batch <- generator_next(augmentation_generator)
  plot(as.raster(batch[1,,,]))
}</pre>
```



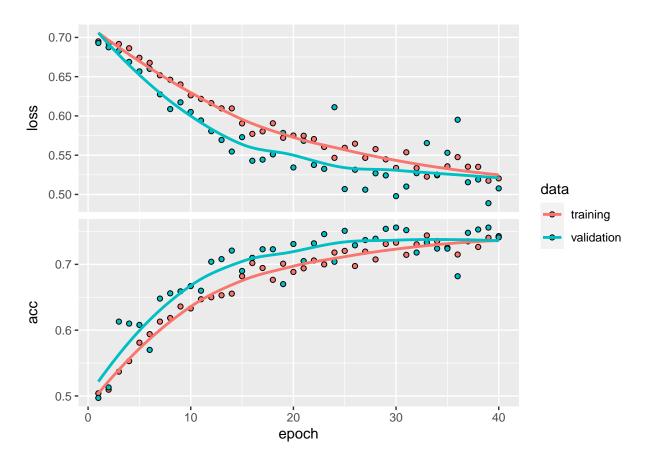






```
par(op)
model2 <- keras_model_sequential() %>%
  layer_conv_2d(filters = 32, kernel_size = c(3,3), activation = "relu", input_shape = c(150,150,3)) %>
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 64, kernel_size = c(3,3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 128, kernel_size = c(3,3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 128, 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")
model2 %>% compile(
  loss = "binary crossentropy",
 optimizer = optimizer_rmsprop(learning_rate = 1e-4),
```

```
metrics = c("acc")
)
test_datagen <- image_data_generator(rescale = 1/255)</pre>
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"
history2 <- model2 %>% fit(
  train_generator,
  steps_per_epoch = 100,
 epochs = 40,
 validation_data = validation_generator,
  validation_steps = 50
test_datagen <- image_data_generator(rescale = 1/255)</pre>
test_generator <- flow_images_from_directory(</pre>
 test_dir,
 test_datagen,
 target_size = c(150, 150),
 batch_size = 20,
  class_mode = "binary"
plot(history2)
```



model2 %>% evaluate_generator(test_generator, steps = 50)

```
## loss acc
## 0.5156532 0.7480000
```

40 epochs later, the loss is now .508, and the accuracy is now .77

Question 2

Adding more pictures to the training data, while keeping the same number of validation and test images

```
dir.create(base_dir)
train_dir <- file.path(base_dir, "train")
dir.create(train_dir)
validation_dir <- file.path(base_dir, "validation")
dir.create(validation_dir)
test_dir <- file.path(base_dir, "test")
dir.create(test_dir)
train_cats_dir <- file.path(train_dir, "cats")
dir.create(train_cats_dir)
train_dogs_dir <- file.path(train_dir, "dogs")
dir.create(train_dogs_dir)
validation_cats_dir <- file.path(validation_dir, "cats")</pre>
```

```
[1] FALSE FA
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fnames <- paste0("cat.", 1501:2000, ".jpg")</pre>
```

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## [1477] FALSE FALSE
## [1489] FALSE FALSE
fnames <- paste0("dog.", 1501:2000, ".jpg")
file.copy(file.path(original_dataset_dir, fnames),
         file.path(validation_dogs_dir))
```

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[1] FALSE FALSE
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```

```
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
)
fnames <- list.files(file.path(train_dir,"cats"), full.names = T)</pre>
img_path <- fnames[[round(runif(1,1,length(fnames)))]]</pre>
img <- image_load(img_path, target_size = c(150,150))</pre>
img_array <- image_to_array(img)</pre>
img_array <- array_reshape(img_array, c(1,150,150,3))</pre>
augmentation_generator <- flow_images_from_data(</pre>
    img_array,
    generator = datagen,
    batch_size = 1
op \leftarrow par(mfrow=c(2,2), pty="s", mar=c(1,0,.1,0))
for (i in 1:4) {
  batch <- generator_next(augmentation_generator)</pre>
  plot(as.raster(batch[1,,,]))
}
```



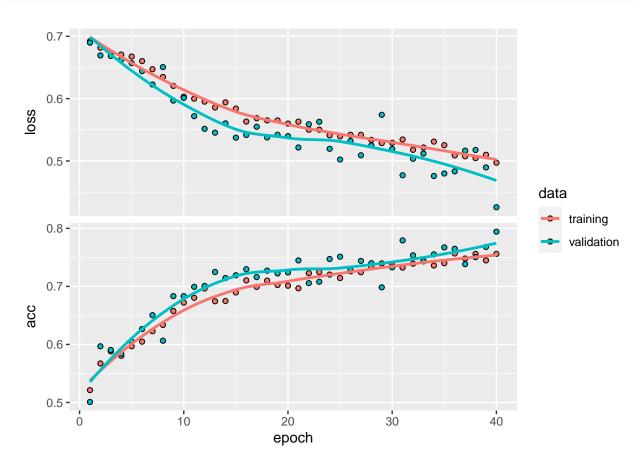






```
par(op)
model3 <- keras_model_sequential() %>%
  layer_conv_2d(filters = 32, kernel_size = c(3,3), activation = "relu", input_shape = c(150,150,3)) %>
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 64, kernel_size = c(3,3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 128, kernel_size = c(3,3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 128, 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")
model3 %>% compile(
  loss = "binary_crossentropy",
  optimizer = optimizer_rmsprop(learning_rate = 1e-4),
  metrics = c("acc")
test_datagen <- image_data_generator(rescale = 1/255)</pre>
train_generator <- flow_images_from_directory(</pre>
  train_dir,
  datagen,
  target_size = c(150, 150),
  batch_size = 25,
 class_mode = "binary"
```

```
validation_generator <- flow_images_from_directory(</pre>
  validation_dir,
  test_datagen,
  target_size = c(150, 150),
  batch_size = 25,
  class_mode = "binary"
history3 <- model3 %>% fit(
  train_generator,
  steps_per_epoch = 100,
 epochs = 40,
 validation_data = validation_generator,
 validation_steps = 50
test_generator <- flow_images_from_directory(</pre>
  test_dir,
 test_datagen,
 target_size = c(150, 150),
 batch_size = 20,
  class_mode = "binary"
plot(history3)
```



```
model3 %>% evaluate_generator(test_generator, steps = 50)
```

```
## loss acc
## 0.4895864 0.7570000
```

This model has a loss of .485 and an accuracy of .76 . Small improvement over the model with less training images. Slightly lower loss and accuracy than the previous model.

Question 3

The following model has the same number of training images, but increases by 500 of validation and testing images.

```
dir.create(base dir)
train_dir <- file.path(base_dir, "train")</pre>
dir.create(train_dir)
validation_dir <- file.path(base_dir, "validation")</pre>
dir.create(validation_dir)
test_dir <- file.path(base_dir, "test")</pre>
dir.create(test dir)
train_cats_dir <- file.path(train_dir, "cats")</pre>
dir.create(train_cats_dir)
train_dogs_dir <- file.path(train_dir, "dogs")</pre>
dir.create(train_dogs_dir)
validation cats dir <- file.path(validation dir, "cats")</pre>
dir.create(validation_cats_dir)
validation_dogs_dir <- file.path(validation_dir, "dogs")</pre>
dir.create(validation_dogs_dir)
test_cats_dir <- file.path(test_dir, "cats")</pre>
dir.create(test_cats_dir)
test_dogs_dir <- file.path(test_dir, "dogs")</pre>
dir.create(test_dogs_dir)
fnames <- paste0("cat.", 1:1500, ".jpg")</pre>
file.copy(file.path(original_dataset_dir, fnames),
          file.path(train_cats_dir))
```

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```

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[181] FALSE FALSE
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[829] FALSE FALSE
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```

```
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
fnames <- list.files(file.path(train_dir,"cats"), full.names = T)</pre>
img_path <- fnames[[round(runif(1,1,length(fnames)))]]</pre>
img <- image_load(img_path, target_size = c(150,150))</pre>
img_array <- image_to_array(img)</pre>
img_array <- array_reshape(img_array, c(1,150,150,3))</pre>
augmentation_generator <- flow_images_from_data(</pre>
    img_array,
    generator = datagen,
    batch size = 1
op \leftarrow par(mfrow=c(2,2), pty="s", mar=c(1,0,.1,0))
for (i in 1:4) {
  batch <- generator_next(augmentation_generator)</pre>
  plot(as.raster(batch[1,,,]))
```



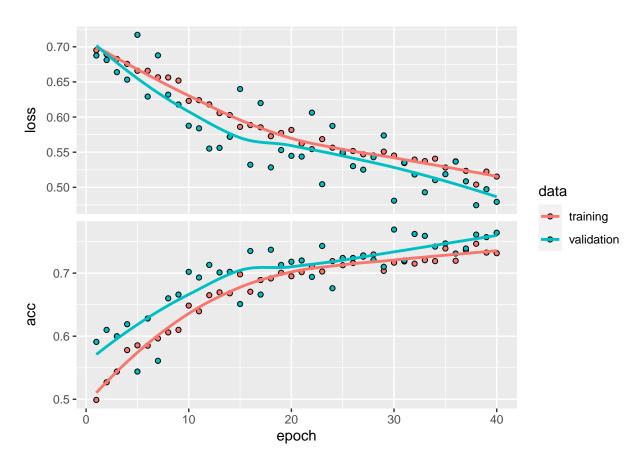






```
par(op)
model4 <- keras_model_sequential() %>%
  layer_conv_2d(filters = 32, kernel_size = c(3,3), activation = "relu", input_shape = c(150,150,3)) %>
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 64, kernel_size = c(3,3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 128, kernel_size = c(3,3), activation = "relu") %>%
  layer_max_pooling_2d(pool_size = c(2,2)) %>%
  layer_conv_2d(filters = 128, 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")
model4 %>% compile(
  loss = "binary_crossentropy",
  optimizer = optimizer_rmsprop(learning_rate = 1e-4),
  metrics = c("acc")
test_datagen <- image_data_generator(rescale = 1/255)</pre>
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"
history4 <- model4 %>% fit(
  train_generator,
  steps_per_epoch = 100,
  epochs = 40,
  validation_data = validation_generator,
 validation_steps = 50
test_datagen <- image_data_generator(rescale = 1/255)</pre>
test_generator <- flow_images_from_directory(</pre>
  test_dir,
  test_datagen,
  target_size = c(150, 150),
  batch_size = 20,
  class_mode = "binary"
plot(history4)
```



```
model4 %>% evaluate_generator(test_generator, steps = 50)
```

```
## loss acc
## 0.5248497 0.7420000
```

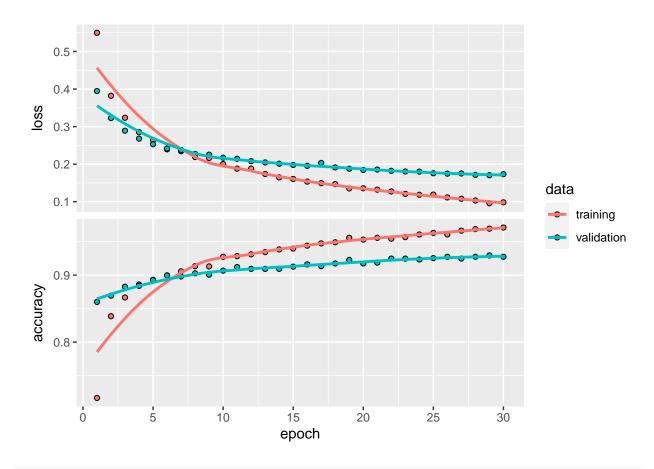
This model now has a loss of .515 and an accuracy of .747. This has a higher loss and slightly lower accuracy than the previous model This is worse than the model with less training and validation data.

Question 4:

The final model uses feature extraction

```
conv_base <- application_vgg16(</pre>
    weights = "imagenet",
    include_top = FALSE,
    input\_shape = c(150, 150, 3)
  base_dir <- "~/Downloads/cats_and_dogs_small"</pre>
  train_dir <- train_dir <- file.path(base_dir, "train")</pre>
  validation_dir <- validation_dir <- file.path(base_dir, "validation")</pre>
  test_dir <- test_dir <- file.path(base_dir, "test")</pre>
  datagen <- 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"
    )
    i <- 0
    while(TRUE) {
      batch <- generator_next(generator)</pre>
      inputs_batch <- batch[[1]]</pre>
      labels_batch <- batch[[2]]</pre>
      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)
    }
    list(
      features = features,
      labels = labels
    )
```

```
train <- extract_features(train_dir, 3000)</pre>
validation <- extract_features(validation_dir, 1500)</pre>
test <- extract_features(test_dir, 1500)</pre>
reshape_features <- function(features) {</pre>
  array_reshape(features, dim = c(nrow(features), 4 * 4 * 512))
train$features <-reshape_features(train$features)</pre>
validation$features <- reshape_features(validation$features)</pre>
test$features <- reshape_features(test$features)</pre>
model5 <- keras_model_sequential() %>%
  layer_dense(units = 256, activation = "relu", input_shape = 4 * 4 * 512) %>%
  layer_dropout(rate = 0.5) %>%
  layer_dense(units = 1, activation = "sigmoid")
model5 %>% compile(
  optimizer = optimizer_rmsprop(learning_rate = 2e-5),
  loss = "binary_crossentropy",
  metrics = c("accuracy")
History5 <- model5 %>% fit(
 train$features, train$labels,
 epochs = 30,
 batch_size = 20,
 validation_data = list(validation$features, validation$labels)
)
plot(History5)
```



model5 %>% evaluate(test\$features, test\$labels)

loss accuracy ## 0.2569493 0.8960000

This model has a loss of .244 and and an accuracy of .898. This has a lower loss and higher accuracy than model 4. This shows a higher level of overfitting than the previous models, but less overfitting than the first model.