

```
functionForDifferentModels <- function(modelname1, modelname2, groupname, foldername1, foldername2, realtimecomponentmodelspace,
iskdetest) {
```

```

kdeWeek3pointPredictionforILI[[i]] <- USNationalXWeeksAhead(kdeFiles[[i]],
  "week three", "yes")
kdeWeek4pointPredictionforILI[[i]] <- USNationalXWeeksAhead(kdeFiles[[i]],
  "week four", "yes")
}

```

```

#####length(kdeWeek1)

```

```

removeStuff <- function(list, iskdetest) {
  # Removes elements 19, 20, 21, and 22.
  if (iskdetest=="iskdetest") {
    return(list.remove(list, c(19, 20, 21, 22)))
  }
  if (iskdetest=="isnotkdetest") {
    return(list)
  }
}

```

```

# Since R works this way we can just do it all at once
}

```

```

kdeWeek1Reduced <- removeStuff(kdeWeek1, iskdetest)
kdeWeek2Reduced <- removeStuff(kdeWeek2, iskdetest)
kdeWeek3Reduced <- removeStuff(kdeWeek3, iskdetest)
kdeWeek4Reduced <- removeStuff(kdeWeek4, iskdetest)

```

```

kdeWeek1pointPredictionforILIReduced <- removeStuff(kdeWeek1pointPredictionforILI, iskdetest)
kdeWeek2pointPredictionforILIReduced <- removeStuff(kdeWeek2pointPredictionforILI, iskdetest)
kdeWeek3pointPredictionforILIReduced <- removeStuff(kdeWeek3pointPredictionforILI, iskdetest)
kdeWeek4pointPredictionforILIReduced <- removeStuff(kdeWeek4pointPredictionforILI, iskdetest)
# Here, all but the rows we want to look at are removed.

```

#There is one crucial thing that we need to do as well before generating the plots: Because our files only share their respective data until week 18 of #2018 and then refer back to the last few weeks of 2017, we need to rotate the order of our files within each of our 8 files.

```

# We will just do another for-loop to replace these values and
# achieve what is functionally a rotation.

```

```

temp1 <- kcdeWeek1
temp2 <- kcdeWeek2
temp3 <- kcdeWeek3
temp4 <- kcdeWeek4
temp5 <- kdeWeek1Reduced
temp6 <- kdeWeek2Reduced
temp7 <- kdeWeek3Reduced
temp8 <- kdeWeek4Reduced

```

```

# We also need to do this for our ILI values.

```

```

temp9 <- kcdeWeek1pointPredictionforILI
temp10 <- kcdeWeek2pointPredictionforILI
temp11 <- kcdeWeek3pointPredictionforILI
temp12 <- kcdeWeek4pointPredictionforILI
temp13 <- kdeWeek1pointPredictionforILIReduced
temp14 <- kdeWeek2pointPredictionforILIReduced
temp15 <- kdeWeek3pointPredictionforILIReduced
temp16 <- kdeWeek4pointPredictionforILIReduced

```

```

for (i in 1:28) {
  # The modular portion (taking the remainder)
  # ensures that we do not go outside the bounds of
  # possible indices.

```

```

  kcdeWeek1[[i]] <- temp1[(((i+17)%28)+1)]
  kcdeWeek2[[i]] <- temp2[(((i+17)%28)+1)]
  kcdeWeek3[[i]] <- temp3[(((i+17)%28)+1)]
  kcdeWeek4[[i]] <- temp4[(((i+17)%28)+1)]

```

```

  kdeWeek1Reduced[[i]] <- temp5[(((i+17)%28)+1)]
  kdeWeek2Reduced[[i]] <- temp6[(((i+17)%28)+1)]
  kdeWeek3Reduced[[i]] <- temp7[(((i+17)%28)+1)]
  kdeWeek4Reduced[[i]] <- temp8[(((i+17)%28)+1)]
}

```

```

kcdeWeek1pointPredictionforILI[[i]] <- temp9[((i+17)%%28)+1]]
kcdeWeek2pointPredictionforILI[[i]] <- temp10[((i+17)%%28)+1]]
kcdeWeek3pointPredictionforILI[[i]] <- temp11[((i+17)%%28)+1]]
kcdeWeek4pointPredictionforILI[[i]] <- temp12[((i+17)%%28)+1]]

kcdeWeek1pointPredictionforLIReduced[[i]] <- temp13[((i+17)%%28)+1]]
kcdeWeek2pointPredictionforLIReduced[[i]] <- temp14[((i+17)%%28)+1]]
kcdeWeek3pointPredictionforLIReduced[[i]] <- temp15[((i+17)%%28)+1]]
kcdeWeek4pointPredictionforLIReduced[[i]] <- temp16[((i+17)%%28)+1]]
}

# Now after refining the data files we need to
# look at the predictions for one week, two weeks,
# three weeks, and four weeks ahead and determine the shape of the
# Kolmogorov-Smirnov test statistics plots.

#####sum(is.na(kcdeWeek1Reduced[[1]]))
#####sum(is.na(kcdeWeek1pointPredictionforLIReduced[[1]]))

# Both sets of data files have length of 28 now.
# So there are two groups of 28 discrete distributions.

# The testStats variable has now been turned into a
# list of four vectors, each of which serves the function
# of the original testStats variable.
# This is done in order that we can generate four graphs
# corresponding to one week, two weeks, three weeks, and
# four weeks ahead.

testStats <- vector("list", 4)

for (i in 1:4) {
  testStats[[i]] <- numeric(28)
}

# In order to generate the CDF for the true K-S
# stats, first we're going to need to generate
# a vector corresponding to the actual CDF.
trueTestStats <- vector("list", 4)
for (i in 1:4) {
  trueTestStats[[i]] <- numeric(28)
}

kcdeWeek1CDF <- kcdeWeek1
kcdeWeek2CDF <- kcdeWeek2
kcdeWeek3CDF <- kcdeWeek3
kcdeWeek4CDF <- kcdeWeek4

kcdeWeek1CDF <- kcdeWeek1Reduced
kcdeWeek2CDF <- kcdeWeek2Reduced
kcdeWeek3CDF <- kcdeWeek3Reduced
kcdeWeek4CDF <- kcdeWeek4Reduced

for (i in 1:28) {
  for (j in 1:131) {
    kcdeWeek1CDF[[i]]$Value[j] <- sum(kcdeWeek1[[i]]$Value[0:j])
    kcdeWeek2CDF[[i]]$Value[j] <- sum(kcdeWeek2[[i]]$Value[0:j])
    kcdeWeek3CDF[[i]]$Value[j] <- sum(kcdeWeek3[[i]]$Value[0:j])
    kcdeWeek4CDF[[i]]$Value[j] <- sum(kcdeWeek4[[i]]$Value[0:j])

    kcdeWeek1CDF[[i]]$Value[j] <- sum(kcdeWeek1Reduced[[i]]$Value[0:j])
    kcdeWeek2CDF[[i]]$Value[j] <- sum(kcdeWeek2Reduced[[i]]$Value[0:j])
    kcdeWeek3CDF[[i]]$Value[j] <- sum(kcdeWeek3Reduced[[i]]$Value[0:j])
    kcdeWeek4CDF[[i]]$Value[j] <- sum(kcdeWeek4Reduced[[i]]$Value[0:j])
  }
}

for (i in 1:28) {
  testStats[[1]][i] <- max(abs(kcdeWeek1[[i]]$Value - kcdeWeek1Reduced[[i]]$Value))
  testStats[[2]][i] <- max(abs(kcdeWeek2[[i]]$Value - kcdeWeek2Reduced[[i]]$Value))
  testStats[[3]][i] <- max(abs(kcdeWeek3[[i]]$Value - kcdeWeek3Reduced[[i]]$Value))

```

```

testStats[[4]][i] <- max(abs(kcdeWeek4[[i]]$Value - kdeWeek4Reduced[[i]]$Value))
}
for (i in 1:28) {
  trueTestStats[[1]][i] <- max(abs(kcdeWeek1CDF[[i]]$Value - kdeWeek1CDF[[i]]$Value))
  trueTestStats[[2]][i] <- max(abs(kcdeWeek2CDF[[i]]$Value - kdeWeek2CDF[[i]]$Value))
  trueTestStats[[3]][i] <- max(abs(kcdeWeek3CDF[[i]]$Value - kdeWeek3CDF[[i]]$Value))
  trueTestStats[[4]][i] <- max(abs(kcdeWeek4CDF[[i]]$Value - kdeWeek4CDF[[i]]$Value))
}

# I also want to put the 5% significance threshold
# on the graph.

#####length(kcdeWeek1[[1]]$Value)

# This is 131, so we are basing each
# test statistic on a data set of size 131.

# Basically our thresholds should be

#####(1.62762)/(sqrt(131))

#####(1.3581)/(sqrt(131))

#####(1.22385)/(sqrt(131))

# based on this website http://www.real-statistics.com
# /statistics-tables/kolmogorov-smirnov-table/
# for significance levels of 0.1, 0.05, and 0.01.
# These are

m <- max(testStats[[1]], testStats[[2]], testStats[[3]], testStats[[4]])

ksStatsAllWeeksAhead <- gf_line(testStats[[1]] ~ seq_along(testStats[[1]]),
  xlab = "", ylab = "", title = "Distances for All Weeks",
  color = ~"1 Week Ahead") %>% gf_lims(y = c(0, m)) %>%
  gf_line(testStats[[2]] ~ seq_along(testStats[[2]]),
  xlab = "", ylab = "", title = "Distances for All Weeks",
  color = ~"2 Weeks Ahead") %>% gf_lims(y = c(0, m)) %>%
  gf_line(testStats[[3]] ~ seq_along(testStats[[3]]),
  xlab = "", ylab = "", title = "Distances for All Weeks",
  color = ~"3 Weeks Ahead") %>% gf_lims(y = c(0, m)) %>%
  gf_line(testStats[[4]] ~ seq_along(testStats[[4]]),
  xlab = "", ylab = "", title = "Distances for All Weeks",
  color = ~"4 Weeks Ahead") %>% gf_lims(y = c(0, m)) + scale_x_continuous(breaks = c(1, 3,
  5, 7, 9, 11, 13, 15, 17, 19, 21, 23, 25, 27),
  labels = c("1" = "w43", "3" = "w45", "5" = "w47", "7" = "w49",
  "9" = "w51", "11" = "w1", "13" = "w3", "15" = "w5", "17" = "w7",
  "19" = "w9", "21" = "w11", "23" = "w13", "25" = "w15", "27" = "w17")) +
  theme(text = element_text(size = 9))

# Having stored these plots we will now
# look at both distributions and compare them.

# We know that our re-ordering of the files in the folder was
# probably successful because our following first two values of
# 0.217 and 0.255 are matched by the first graph (for one week
# ahead) - in the real-time-component models folder,
# kcdeFiles[[19]] corresponds to EW43-2017-ReichLab_kcde
# and kdeFiles[[23]] corresponds to EW43-2017-ReichLab_kde
# in the ReichLab_kcde and ReichLab_kde folders respectively.

tmp <- USNationalXWeeksAhead(kcdeFiles[[19]], "week one", "no")
tmp2 <- USNationalXWeeksAhead(kdeFiles[[23]], "week one", "no")

```

```

tmp3 <- USNationalXWeeksAhead(kcdeFiles[[20]], "week one", "no")
tmp4 <- USNationalXWeeksAhead(kcdeFiles[[24]], "week one", "no")

#####max(abs(tmp$Value - tmp2$Value))
#####max(abs(tmp3$Value - tmp4$Value))

# http://reichlab.io/assets/images/logo/nav-logo.png
#####img <- readPicture("C:/Users/gladi/Downloads/simple.svg")
#####imgGrob <- gTree(children=gList(pictureGrob(img)))

#####

# We're also going to try to see if the statistics
# are linear.
# We're going to include all points and graph the maximums for one week ahead.
# The gray dots are points before the log transformation.

# We are going to graph the maximums
# for one week ahead. A log transformation
# might be needed so we're going to use that
# and include the original points in orange.

linearityTestMaximums <- gf_point(log(testStats[[1]][1:28]) ~ seq_along(testStats[[1]][1:28]),
  title = "Maximums for One Week Ahead",
  xlab = "Index", ylab = "log(max_distance_between_pmfs)") %>% gf_lims(y = c(0, max(testStats[[1]]))) %>%
  gf_point(testStats[[1]][1:28] ~ seq_along(testStats[[1]][1:28]),
    xlab = "Index",
    color = ~"before log transformation") %>% gf_lims(y = c(0, max(testStats[[1]]))) %>%
  gf_theme(legend.position = "bottom") +
  geom_smooth(method = 'lm', formula = y ~ x)

lineOfBestFit <- glm(log(testStats[[1]][1:28]) ~ seq_along(testStats[[1]][1:28]))
#####stepAIC(lineOfBestFit)

actualILlagainstMaximums <-
  gf_line(testStats[[1]] ~ weightedILIRange,
    xlab = "True ILI", ylab = "Maximum Distances", title = "Maximum Distances Against Weighted %ILI",
    color = ~"1 Week Ahead") %>% gf_theme(legend.position = "top") %>% gf_lims(y = c(0, m)) %>%

  gf_line(testStats[[2]] ~ weightedILIRange,
    xlab = "", ylab = "", title = "Maximum Distances Against Weighted %ILI",
    color = ~"2 Weeks Ahead") %>% gf_lims(y = c(0, m)) %>%
  gf_line(testStats[[3]] ~ weightedILIRange,
    xlab = "", ylab = "", title = "Maximum Distances Against Weighted %ILI",
    color = ~"3 Weeks Ahead") %>% gf_lims(y = c(0, m)) %>%
  gf_line(testStats[[4]] ~ weightedILIRange,
    xlab = "True ILI", ylab = "Maximum Distances", title = "Maximum Distances Against Weighted %ILI",
    color = ~"4 Weeks Ahead") %>%
  gf_lims(y = c(0, m)) + scale_x_continuous(breaks = c(1.52071, 1.53749, 1.55644, 1.62950, 1.74765, 1.79983, 1.88999, 2.06099, 2.09090,
2.28197, 2.28786, 2.42157, 2.49534, 2.58516, 2.77608, 3.19245, 3.37109, 3.69947, 4.73950, 4.96989, 5.71576, 5.75997, 5.90718, 6.45058,
6.52457, 7.17131, 7.39126, 7.52959), labels = c("1.5", "1.5", "1.6", "1.6", "1.7", "1.8", "1.9", "2.1", "2.1", "2.3", "2.3", "2.4", "2.5", "2.6", "2.8",
"3.2", "3.4", "3.7", "4.7", "5.0", "5.7", "5.8", "5.9", "6.5", "6.5", "7.2", "7.4", "7.5")) +
  theme(text = element_text(size = 9))

# It looks like the maximum distance is much
# more for the first three starting weeks
# between these two models.

#####plot
#####typeof(plot)

# By the same method we're going to derive the
# true, standard K-S test statistics from the
# cumulative distribution functions.

m2 <- max(trueTestStats[[1]], trueTestStats[[2]], trueTestStats[[3]], trueTestStats[[4]])

```

```

trueKSSStatsAllWeeksAhead <- gf_line(trueTestStats[[1]] ~ seq_along(trueTestStats[[1]]),
  xlab = "", ylab = "", title = "K-S Stats for All Weeks",
  color = ~"1 Week Ahead") %>% gf_lims(y = c(0, m2)) %>%
  gf_line(trueTestStats[[2]] ~ seq_along(trueTestStats[[2]]),
  xlab = "", ylab = "", title = "K-S Stats for All Weeks",
  color = ~"2 Weeks Ahead") %>% gf_lims(y = c(0, m2)) %>%
  gf_line(trueTestStats[[3]] ~ seq_along(trueTestStats[[3]]),
  xlab = "", ylab = "", title = "K-S Stats for All Weeks",
  color = ~"3 Weeks Ahead") %>% gf_lims(y = c(0, m2)) %>%
  gf_line(trueTestStats[[4]] ~ seq_along(trueTestStats[[4]]),
  xlab = "", ylab = "", title = "K-S Stats for All Weeks",
  color = ~"4 Weeks Ahead") %>% gf_lims(y = c(0, m2)) + scale_x_continuous(breaks = c(1, 3,
    5, 7, 9, 11, 13, 15, 17, 19, 21, 23, 25, 27),
    labels = c("1" = "w43", "3" = "w45", "5" = "w47", "7" = "w49",
      "9" = "w51", "11" = "w1", "13" = "w3", "15" = "w5", "17" = "w7",
      "19" = "w9", "21" = "w11", "23" = "w13", "25" = "w15", "27" = "w17")) +
  theme(text = element_text(size = 9)) + geom_hline(aes(yintercept = 0.1422058)) + geom_text(aes(0.01422058, label = 0.01, vjust = -1)) +
  geom_hline(aes(yintercept = 0.1186577)) + geom_text(aes(0.01186577, label = 0.05, vjust = -0.5)) + geom_hline(aes(yintercept = 0.1069283)) +
  geom_text(aes(0.01069283, label = 0.1, vjust = 1))

```

```

# We are going to graph the K-S test
# statistics
# for one week ahead. A log transformat-
# ion might be needed so we're going to use that
# and include the original points in orange.

```

```

linearityTestKSSStatistics <- gf_point(log(trueTestStats[[1]][1:28]) ~ seq_along(trueTestStats[[1]][1:28]),
  title = "K-S Statistics for One Week Ahead",
  xlab = "Index", ylab = "log(K-S Statistics)") %>%
  gf_point(testStats[[1]][1:28] ~ seq_along(testStats[[1]][1:28]),
  xlab = "Index",
  color = ~"before log transformation") %>% gf_theme(legend.position = "bottom") +
  geom_smooth(method = 'lm', formula = y ~ x)

```

```

lineOfBestFit2 <- glm((log(trueTestStats[[1]][1:28]) ~ seq_along(trueTestStats[[1]][1:28])))
#####stepAIC(lineOfBestFit2)

```

```

actualILlagainstKSSStatistics <-
  gf_line(trueTestStats[[1]] ~ weightedILIRange,
  xlab = "True ILI", ylab = "K-S Test Statistics", title = "Test Statistics Against Weighted %ILI",
  color = ~"1 Week Ahead") %>% gf_theme(legend.position = "top") %>% gf_lims(y = c(0, m2)) %>%

  gf_line(trueTestStats[[2]] ~ weightedILIRange,
  xlab = "", ylab = "", title = "Test Statistics Against Weighted %ILI",
  color = ~"2 Weeks Ahead") %>% gf_lims(y = c(0, m2)) %>%
  gf_line(trueTestStats[[3]] ~ weightedILIRange,
  xlab = "", ylab = "", title = "Test Statistics Against Weighted %ILI",
  color = ~"3 Weeks Ahead") %>% gf_lims(y = c(0, m2)) %>%
  gf_line(trueTestStats[[4]] ~ weightedILIRange,
  xlab = "True ILI", ylab = "K-S Test Statistics", title = "Test Statistics Against Weighted %ILI",
  color = ~"4 Weeks Ahead") %>%
  gf_lims(y = c(0, m2)) + scale_x_continuous(breaks = c(1.52071, 1.53749, 1.55644, 1.62950, 1.74765, 1.79983, 1.88999, 2.06099, 2.09090,
    2.28197, 2.28786, 2.42157, 2.49534, 2.58516, 2.77608, 3.19245, 3.37109, 3.69947, 4.73950, 4.96989, 5.71576, 5.75997, 5.90718, 6.45058,
    6.52457, 7.17131, 7.39126, 7.52959), labels = c("1.5", "1.5", "1.6", "1.6", "1.7", "1.8", "1.9", "2.1", "2.1", "2.3", "2.3", "2.4", "2.5", "2.6", "2.8",
    "3.2", "3.4", "3.7", "4.7", "5.0", "5.7", "5.8", "5.9", "6.5", "6.5", "7.2", "7.4", "7.5")) +
  theme(text = element_text(size = 9)) + geom_hline(aes(yintercept = 0.1422058)) + geom_text(aes(0.01422058, label = 0.01, vjust = -1)) +
  geom_hline(aes(yintercept = 0.1186577)) + geom_text(aes(0.01186577, label = 0.05, vjust = -0.5)) + geom_hline(aes(yintercept = 0.1069283)) +
  geom_text(aes(0.01069283, label = 0.1, vjust = 1))

```

```

#The test statistics seems to oscillate to some degree, and the initial spike in their values seems to indicate that a linear regression still is most
#likely not the best fit.

```

```

#The true ILI of course takes on a positive value that both models try to predict but with fairly significant differences. P-Values should be
extracted #from these test statistics to determine the magnitude of this significance.

```

```

#It makes sense that our models tend to differ more as they try to predict further into the future.

```

#Before we look for possible linearity we should create graphs of each individual distribution in order to determine how the models are responsible for #the larger values of the first three points on the graphs.

# The following code will generate graphs of the probabilities assigned to each bin for one week ahead, two weeks ahead, and three weeks ahead.

```
kcde1week <- kde1week <- kcde2weeks <- kde2weeks <-
  kcde3weeks <- kde3weeks <- kcde4weeks <- kde4weeks <-
  vector("list", 6)
for (i in 1:6) {
  kcde1week[[i]] <- kde1week[[i]] <- kcde2weeks[[i]] <-
    kde2weeks[[i]] <- kcde3weeks[[i]] <- kde3weeks[[i]] <-
    kcde4weeks[[i]] <- kde4weeks[[i]] <-
    numeric(131)
}
```

# Because kcdeweek1 is a data frame of size 131  
# and we basically want to explain the fact that  
# the first three K-S test statistics between the  
# kcde and kde models are much larger than the rest,  
# we use the first for-loop to populate the kcde1week  
# list with four different vectors that contain all  
# probability predictions for each bin for that week ahead.

# We arbitrarily chose to have four vectors in each  
# list because this would show the probability  
# distributions that correspond to the first  
# four K-S test statistics on the graph and might  
# provide some insight into which model is the  
# culprit in the observed increased difference.

# Creating new vectors isn't actually necessary but will  
# make the code for the actual graphs slightly smaller,  
# which is what we want. It will just allow us to focus  
# on what we want.

```
for (j in 1:6) {
  for (i in 1:131) {
    kcde1week[[j]][[i]] <- kcdeWeek1[[j]]$Value[[i]]
    kde1week[[j]][[i]] <- kdeWeek1[[j]]$Value[[i]]
    kcde2weeks[[j]][[i]] <- kcdeWeek2[[j]]$Value[[i]]
    kde2weeks[[j]][[i]] <- kdeWeek2[[j]]$Value[[i]]
    kcde3weeks[[j]][[i]] <- kcdeWeek3[[j]]$Value[[i]]
    kde3weeks[[j]][[i]] <- kdeWeek3[[j]]$Value[[i]]
    kcde4weeks[[j]][[i]] <- kcdeWeek4[[j]]$Value[[i]]
    kde4weeks[[j]][[i]] <- kdeWeek4[[j]]$Value[[i]]
  }
}
```

# Now we must also define the plots.

```
max1 <- max(kcde1week[[1]], kcde1week[[2]], kcde1week[[3]], kcde1week[[4]], kcde1week[[5]], kcde1week[[6]])
```

# For some reason we have to define the legend not  
# directly using the paste() command.

```
Legend <- paste("1w ", modelName1, sep = "")
```

```
plotw43AllWeeksAhead <- gf_line(kcde1week[[1]] ~ seq_along(kcde1week[[1]]), color = ~Legend, xlab = "Week 43", ylab = "") %>%
  gf_line(kde1week[[1]] ~ seq_along(kde1week[[1]]), color = ~paste("1w ", modelName2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kcde2weeks[[1]] ~ seq_along(kcde2weeks[[1]]), color = ~paste("2w ", modelName1, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kde2weeks[[1]] ~ seq_along(kde2weeks[[1]]), color = ~paste("2w ", modelName2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kcde3weeks[[1]] ~ seq_along(kcde3weeks[[1]]), color = ~paste("3w ", modelName1, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kde3weeks[[1]] ~ seq_along(kde3weeks[[1]]), color = ~paste("3w ", modelName2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kcde4weeks[[1]] ~ seq_along(kcde4weeks[[1]]), color = ~paste("4w ", modelName1, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  Bins", title = "PMFs") %>%
```

```
gf_line(kde4weeks[[1]] ~ seq_along(kde4weeks[[1]]), color = ~paste("4w ", modelName2, sep = "")) %>% gf_lims(y = c(0, max1)) +
  scale_x_continuous(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130), labels = c("0%", "1%", "2%", "3%", "4%", "5%", "6%",
    "7%", "8%", "9%", "10%", "11%", "12%", "13%")) + theme(text = element_text(size = 9))
```

```
Legend <- paste("1w ", modelName1, sep = "")
```

```
plotw44AllWeeksAhead <- gf_line(kcde1week[[2]] ~ seq_along(kcde1week[[2]]), color = ~Legend, xlab = "Week 44", ylab = "") %>%
  gf_line(kde1week[[2]] ~ seq_along(kde1week[[2]]), color = ~paste("1w ", modelName2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kcde2weeks[[2]] ~ seq_along(kcde2weeks[[2]]), color = ~paste("2w ", modelName1, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
  gf_line(kde2weeks[[2]] ~ seq_along(kde2weeks[[2]]), color = ~paste("2w ", modelName2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
```

```

gf_line(kcde3weeks[[2]] ~ seq_along(kcde3weeks[[2]]), color = ~paste("3w ", modelname1, sep = ""), xlab = "Week 44", ylab = "") %>%
gf_line(kcde3weeks[[2]] ~ seq_along(kcde3weeks[[2]]), color = ~paste("3w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde4weeks[[2]] ~ seq_along(kcde4weeks[[2]]), color = ~paste("4w ", modelname1, sep = ""), xlab = "Week 44", ylab = "Probability for
Bins", title = "PMFs") %>%
gf_line(kcde4weeks[[2]] ~ seq_along(kcde4weeks[[2]]), color = ~paste("4w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) +
scale_x_continuous(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130), labels = c("0%", "1%", "2%", "3%", "4%", "5%", "6%",
"7%", "8%", "9%", "10%", "11%", "12%", "13%")) + theme(text = element_text(size = 9))
Legend <- paste("1w ", modelname1, sep = "")
plotw45AllWeeksAhead <- gf_line(kcde1week[[3]] ~ seq_along(kcde1week[[3]]), color = ~Legend, xlab = "Week 45", ylab = "") %>%
gf_line(kcde1week[[3]] ~ seq_along(kcde1week[[3]]), color = ~paste("1w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde2weeks[[3]] ~ seq_along(kcde2weeks[[3]]), color = ~paste("2w ", modelname1, sep = ""), xlab = "Week 45", ylab = "") %>%
gf_line(kcde2weeks[[3]] ~ seq_along(kcde2weeks[[3]]), color = ~paste("2w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde3weeks[[3]] ~ seq_along(kcde3weeks[[3]]), color = ~paste("3w ", modelname1, sep = ""), xlab = "Week 45", ylab = "") %>%
gf_line(kcde3weeks[[3]] ~ seq_along(kcde3weeks[[3]]), color = ~paste("3w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde4weeks[[3]] ~ seq_along(kcde4weeks[[3]]), color = ~paste("4w ", modelname1, sep = ""), xlab = "Week 45", ylab = "Probability for
Bins", title = "PMFs") %>%
gf_line(kcde4weeks[[3]] ~ seq_along(kcde4weeks[[3]]), color = ~paste("4w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) +
scale_x_continuous(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130), labels = c("0%", "1%", "2%", "3%", "4%", "5%", "6%",
"7%", "8%", "9%", "10%", "11%", "12%", "13%")) + theme(text = element_text(size = 9))
Legend <- paste("1w ", modelname1, sep = "")
plotw46AllWeeksAhead <- gf_line(kcde1week[[4]] ~ seq_along(kcde1week[[4]]), color = ~Legend, xlab = "Week 46", ylab = "") %>%
gf_line(kcde1week[[4]] ~ seq_along(kcde1week[[4]]), color = ~paste("1w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde2weeks[[4]] ~ seq_along(kcde2weeks[[4]]), color = ~paste("2w ", modelname1, sep = ""), xlab = "Week 46", ylab = "") %>%
gf_line(kcde2weeks[[4]] ~ seq_along(kcde2weeks[[4]]), color = ~paste("2w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde3weeks[[4]] ~ seq_along(kcde3weeks[[4]]), color = ~paste("3w ", modelname1, sep = ""), xlab = "Week 46", ylab = "") %>%
gf_line(kcde3weeks[[4]] ~ seq_along(kcde3weeks[[4]]), color = ~paste("3w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde4weeks[[4]] ~ seq_along(kcde4weeks[[4]]), color = ~paste("4w ", modelname1, sep = ""), xlab = "Week 46", ylab = "Probability for
Bins", title = "PMFs") %>%
gf_line(kcde4weeks[[4]] ~ seq_along(kcde4weeks[[4]]), color = ~paste("4w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) +
scale_x_continuous(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130), labels = c("0%", "1%", "2%", "3%", "4%", "5%", "6%",
"7%", "8%", "9%", "10%", "11%", "12%", "13%")) + theme(text = element_text(size = 9))
Legend <- paste("1w ", modelname1, sep = "")
plotw47AllWeeksAhead <- gf_line(kcde1week[[5]] ~ seq_along(kcde1week[[5]]), color = ~Legend, xlab = "Week 47", ylab = "") %>%
gf_line(kcde1week[[5]] ~ seq_along(kcde1week[[5]]), color = ~paste("1w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde2weeks[[5]] ~ seq_along(kcde2weeks[[5]]), color = ~paste("2w ", modelname1, sep = ""), xlab = "Week 47", ylab = "") %>%
gf_line(kcde2weeks[[5]] ~ seq_along(kcde2weeks[[5]]), color = ~paste("2w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde3weeks[[5]] ~ seq_along(kcde3weeks[[5]]), color = ~paste("3w ", modelname1, sep = ""), xlab = "Week 47", ylab = "") %>%
gf_line(kcde3weeks[[5]] ~ seq_along(kcde3weeks[[5]]), color = ~paste("3w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde4weeks[[5]] ~ seq_along(kcde4weeks[[5]]), color = ~paste("4w ", modelname1, sep = ""), xlab = "Week 47", ylab = "Probability for
Bins", title = "PMFs") %>%
gf_line(kcde4weeks[[5]] ~ seq_along(kcde4weeks[[5]]), color = ~paste("4w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) +
scale_x_continuous(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130), labels = c("0%", "1%", "2%", "3%", "4%", "5%", "6%",
"7%", "8%", "9%", "10%", "11%", "12%", "13%")) + theme(text = element_text(size = 9))
Legend <- paste("1w ", modelname1, sep = "")
plotw48AllWeeksAhead <- gf_line(kcde1week[[6]] ~ seq_along(kcde1week[[6]]), color = ~Legend, xlab = "Week 48", ylab = "") %>%
gf_line(kcde1week[[6]] ~ seq_along(kcde1week[[6]]), color = ~paste("1w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde2weeks[[6]] ~ seq_along(kcde2weeks[[6]]), color = ~paste("2w ", modelname1, sep = ""), xlab = "Week 48", ylab = "") %>%
gf_line(kcde2weeks[[6]] ~ seq_along(kcde2weeks[[6]]), color = ~paste("2w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde3weeks[[6]] ~ seq_along(kcde3weeks[[6]]), color = ~paste("3w ", modelname1, sep = ""), xlab = "Week 48", ylab = "") %>%
gf_line(kcde3weeks[[6]] ~ seq_along(kcde3weeks[[6]]), color = ~paste("3w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) %>%
gf_line(kcde4weeks[[6]] ~ seq_along(kcde4weeks[[6]]), color = ~paste("4w ", modelname1, sep = ""), xlab = "Week 48", ylab = "Probability for
Bins", title = "PMFs") %>%
gf_line(kcde4weeks[[6]] ~ seq_along(kcde4weeks[[6]]), color = ~paste("4w ", modelname2, sep = "")) %>% gf_lims(y = c(0, max1)) +
scale_x_continuous(breaks = c(0, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, 120, 130), labels = c("0%", "1%", "2%", "3%", "4%", "5%", "6%",
"7%", "8%", "9%", "10%", "11%", "12%", "13%")) + theme(text = element_text(size = 9))

```

```

# These allow us to see that the KCDE model is primarily the one that is
# responsible for these extreme deviations in K-S test statistics.
# As we can see, the KDE model largely remains the same as far as its
# probabilities for each bin are presented across each week (43, 44, 45,
# 46, 47, 48)

```

```

# We can also plot the actual ILI against the K-S
# test statistics to find a correlation.
# The correspondence of each ILI - K-S Test statistic
# pair is preserved in these graphs.

```



```
# In order to appropriately label the graph we're going to
# have to look at the values for true weighted ILI in their
# proper order.
```

```
#####sort(weightedILIRange, decreasing = F)
#####round(c(1.52071, 1.53749, 1.55644, 1.62950, 1.74765, 1.79983, 1.88999, 2.06099, 2.09090, 2.28197, 2.28786,
2.42157, 2.49534, 2.58516, 2.77608, 3.19245, 3.37109, 3.69947, 4.73950, 4.96989, 5.71576, 5.75997, 5.90718, 6.45058, 6.52457, 7.17131,
7.39126, 7.52959), 1)
```

```
#####plot(lineOfBestFit)
```

```
#####for (i in c(1,2)) { # We want to do a Wald Test
##### print(1 - pchisq((coef(lineOfBestFit)[i] / sqrt(vcov(lineOfBestFit)[i,i]))^2, df = 1))
#####}
#####anova(lineOfBestFit, stepAIC(lineOfBestFit, test = "Chisq")
#####lineOfBestFit$residuals
#####residualPlots(lineOfBestFit, tests = TRUE)
#####inflIndexPlot(lineOfBestFit)
#####summary(lineOfBestFit)
#####qqnorm(lineOfBestFit$residuals)
```

#However, there is no definitive indication that the test statistics should come from a commonly known function such as an exponential one, so trying to find a linear function is probably not a high-yield path for most comparisons between models.

```
# Now, our second set of graphs shows the maximum
# distance between the point values for each week.
```

```
# This is different from the K-S test.
```

```
pointValues <- vector("list", 4)
```

```
for (i in 1:4) {
  pointValues[[i]] <- numeric(28)
} # Now that we have populated this list,
```

```
for (i in 1:28) {
  pointValues[[1]][i] <- max(abs(kcdeWeek1pointPredictionforILI[[i]]$Value -
    kcdeWeek1pointPredictionforLIReduced[[i]]$Value))
  pointValues[[2]][i] <- max(abs(kcdeWeek2pointPredictionforILI[[i]]$Value -
    kcdeWeek2pointPredictionforLIReduced[[i]]$Value))
  pointValues[[3]][i] <- max(abs(kcdeWeek3pointPredictionforILI[[i]]$Value -
    kcdeWeek3pointPredictionforLIReduced[[i]]$Value))
  pointValues[[4]][i] <- max(abs(kcdeWeek4pointPredictionforILI[[i]]$Value -
    kcdeWeek4pointPredictionforLIReduced[[i]]$Value))
}
```

```
max2 <- max(pointValues[[1]], pointValues[[2]], pointValues[[3]], pointValues[[4]])
```

```
allPointValueDifferences <- gf_line(pointValues[[1]] ~ seq_along(pointValues[[1]]), color = ~"1w ahead",
  xlab = "Index", ylab = "Difference in Predicted ILI") %>% gf_lims(y = c(0, max2)) %>%
  gf_line(pointValues[[2]] ~ seq_along(pointValues[[2]]), color = ~"2w ahead",
  xlab = "Index", ylab = "Difference in Predicted ILI") %>% gf_lims(y = c(0, max2)) %>%
  gf_line(pointValues[[3]] ~ seq_along(pointValues[[3]]), color = ~"3w ahead",
  xlab = "Index", ylab = "Difference in Predicted ILI") %>% gf_lims(y = c(0, max2)) %>%
  gf_line(pointValues[[4]] ~ seq_along(pointValues[[4]]), color = ~"4w ahead",
  xlab = "Index", ylab = "Difference in Predicted ILI", title = "Difference in Point Values") + theme(text = element_text(size
= 9))
```

```
grid.arrange(trueILIPlot, allPointValueDifferences)
```

```
grid.arrange(plotw43AllWeeksAhead, plotw44AllWeeksAhead, plotw45AllWeeksAhead, plotw46AllWeeksAhead, plotw47AllWeeksAhead,
plotw48AllWeeksAhead, top = paste(modelname1, " and ", modelname2, " PMFS for %ILI, All Weeks Ahead"))
```

```
grid.arrange(ksStatsAllWeeksAhead,
  linearityTestMaximums, actualILIagainstMaximums, top = paste("Comparing ", groupname, "'s ", modelname1,
" and ", modelname2, " Models \n for Selected Weeks of 2017-2018 (Week 43, 2017 to ",
"Week 18, 2018) \n Maximum Differences between the ", modelname1, " and ", modelname2, " Models' \n",
"Probability Mass Functions for All ILI Bins (increment 0.1)", sep = ""))
```

```
grid.arrange(trueKSStatsAllWeeksAhead, linearityTestKSStatistics, actualILIagainstKSStatistics, top = paste("Comparing ", groupname, "'s ",
modelName1,
" and ", modelName2, " Models \n for Selected Weeks of 2017-2018 (Week 43, 2017 to ",
"Week 18, 2018) \n Kolmogorov-Smirnov Test Statistics between the ", modelName1, " and ",
modelName2, " Models' \n Empirical CDF for All ILI Bins (increment 0.1)", sep = ""))
}
```