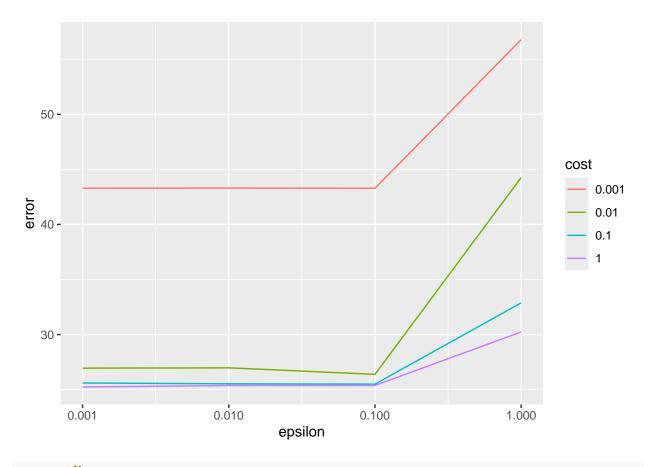
Weekly Assignment 8

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2025-04-22



```
tune.out$best.parameters
```

```
## cost epsilon
## 4 1 0.001
```

tune.out \$performances

```
##
       cost epsilon
                       error dispersion
## 1
     0.001
             0.001 43.29272 13.078277
## 2 0.010
             0.001 26.95169
                             10.557590
## 3 0.100
             0.001 25.60440
                             10.829528
## 4 1.000
             0.001 25.23923
                              10.888296
## 5 0.001
             0.010 43.30323
                              13.053327
## 6 0.010
             0.010 26.97850
                              10.603666
## 7 0.100
             0.010 25.52956
                             10.792008
## 8 1.000
             0.010 25.37565
                             11.032447
## 9 0.001
             0.100 43.28249
                             12.821787
## 10 0.010
             0.100 26.38916 10.443813
## 11 0.100
             0.100 25.50024
                             10.688779
## 12 1.000
             0.100 25.37825
                             10.714303
## 13 0.001
             1.000 56.78102
                              9.859582
## 14 0.010
             1.000 44.25528
                              7.906997
## 15 0.100
             1.000 32.87987
                               8.671015
## 16 1.000
             1.000 30.23543
                               6.873077
```

The best parameter tuple (cost, epsilon) from this 10-fold cross-validation training is (1, 0.001). The corresponding purple curve is convex throughout the entire range. The detailed performance results were also printed: we see that $(cost = 1, \epsilon = 0.001)$ indeed has the lowest error, closely followed by $(cost = 1, \epsilon = 0.01)$. The $(cost = 0.001, \epsilon = 0.001)$ pair produced the highest error. Ultimately, a very low margin of tolerance and a high cost yields the lowest cross-validation MSE.

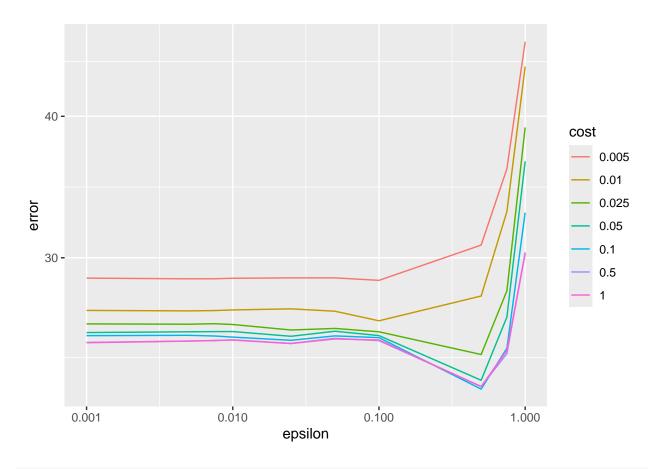
```
best.model = tune.out$best.model
ypred = predict(best.model, Boston[test,])
ytrue = Boston$medv[test]

mse = mean((ytrue - ypred)^2)
mae = mean(abs(ytrue - ypred))
cat('MSE:', round(mse, 4), '\n')

## MSE: 26.2043

cat('MAE:', round(mae, 4))
## MAE: 3.5494
```

Let us now try a grid search with smaller resolution and compare the test MSEs:



tune.out\$best.parameters

```
## cost epsilon
## 54 0.1 0.5
```

After carrying out a more elaborate grid search, the best parameter tuple turns out to be (cost = 0.1, $\epsilon = 0.5$). This is a much more balanced pair, with a much higher margin of tolerance and softer penalisation outside of the margin. The cross-validation MSE obtained with this pair is 22.96781, which is lower than the previous best training MSE from the higher resolution grid.

```
best.model = tune.out$best.model
ypred = predict(best.model, Boston[test,])
ytrue = Boston$medv[test]

mse = mean((ytrue - ypred)^2)
mae = mean(abs(ytrue - ypred))
cat('MSE:', round(mse, 4), '\n')

## MSE: 25.0509

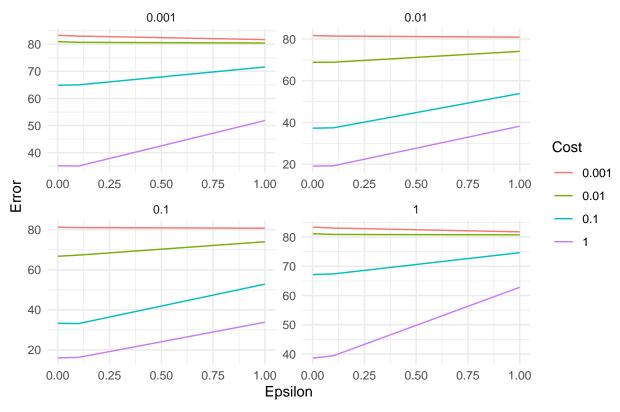
cat('MAE:', round(mae, 4))
```

MAE: 3.6301

And with this pair we also obtain a lower test MSE - almost 5% lower, which is not a huge gain. Nevertheless, it is interesting to note that the mean absolute error is slightly higher for this pair than for the previous best pair - 2% higher. The previous higher cost parameter likely lead to a little bit of overfitting.

```
cost = c(.001, .01, .1, 1)
epsilon = c(.001, .01, .1, 1)
gamma = c(.001, .01, .1, 1)
set.seed(4645251)
tune.out = tune(svm, medv ~ ., data=Boston[train,], kernel='radial'
                , ranges=list(cost=cost, epsilon=epsilon, gamma=gamma))
perf = tune.out$performances
perf$cost = factor(perf$cost)
res <- tune.out$performances</pre>
# Plot: one for each gamma
ggplot(perf, aes(epsilon, error, group=cost, color=cost)) +
  geom_line() +
  facet_wrap(~ gamma, scales = "free") +
 labs(
    title = "MSE across epsilon and cost for each gamma",
    x = "Epsilon",
    y = "Error",
    color = "Cost"
  ) +
  theme_minimal()
```

MSE across epsilon and cost for each gamma



```
tune.out$best.parameters
```

After performing cross-validation with an SVR with a radial basis kernel, we find that the best parameter triplet is (cost = 1, $\epsilon = 0.001$, $\gamma = 0.1$). The first two values are familiar, since they are the exact same obtained with a linear kernel and the same resolution grid in our first experiment.

```
best.model = tune.out$best.model
ypred = predict(best.model, Boston[test,])
ytrue = Boston$medv[test]

mse = mean((ytrue - ypred)^2)
mae = mean(abs(ytrue - ypred))
cat('MSE:', round(mse, 4), '\n')

## MSE: 17.1149

cat('MAE:', round(mae, 4))

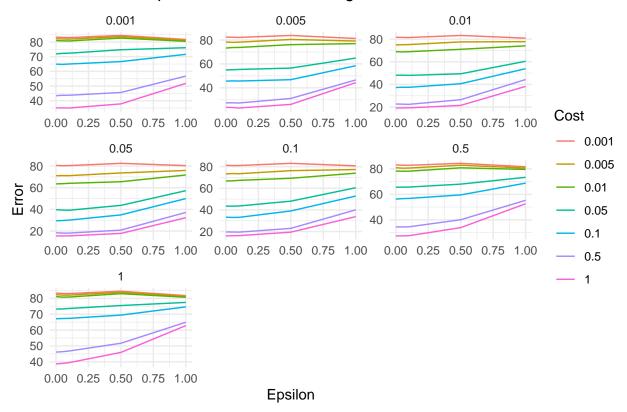
## MAE: 2.442
```

Both the MSE and MAE are significantly lower than what our best parameter pair yielded with a linear kernel.

Let us now repeat this experiment with a higher resolution grid:

```
cost = c(.001, .005, .01, .05, .1, .5, 1)
epsilon = c(.001, .005, .01, .05, .1, .5, 1)
gamma = c(.001, .005, .01, .05, .1, .5, 1)
set.seed(4645251)
tune.out = tune(svm, medv ~ ., data=Boston[train,], kernel='radial'
                , ranges=list(cost=cost, epsilon=epsilon, gamma=gamma))
perf = tune.out$performances
perf$cost = factor(perf$cost)
res <- tune.out$performances
# Plot: one for each gamma
ggplot(perf, aes(epsilon, error, group=cost, color=cost)) +
  geom_line() +
  facet_wrap(~ gamma, scales = "free") +
   title = "MSE across epsilon and cost for each gamma",
   x = "Epsilon",
   y = "Error",
   color = "Cost"
  theme_minimal()
```

MSE across epsilon and cost for each gamma



tune.out\$best.parameters

```
## cost epsilon gamma
## 175 1 0.05 0.05
```

```
best.model = tune.out$best.model
ypred = predict(best.model, Boston[test,])
ytrue = Boston$medv[test]

mse = mean((ytrue - ypred)^2)
mae = mean(abs(ytrue - ypred))
cat('MSE:', round(mse, 4), '\n')
```

MSE: 16.8298

```
cat('MAE:', round(mae, 4))
```

MAE: 2.4546

It turns out that the $(cost=1,\,\epsilon=0.05,\,\gamma=0.05)$ triplet gives an even lower MSE - though the gain is really minimal, only 1.5% -, but offers no improvement in terms of MAE. The fact that the best gamma is relatively high means that each point influences a rather small region, so bias is lower here. Our training set is relatively big in statistical terms, so it is no surprise that lowering the bias yields better predictions.

```
lm.model <- lm(medv ~ ., data = Boston[train, ])
lm.pred <- predict(lm.model, newdata = Boston[test, ])

mse = mean((ytrue - lm.pred)^2)
mae = mean(abs(ytrue - lm.pred))
cat('MSE:', round(mse, 4), '\n')

## MSE: 26.151

cat('MAE:', round(mae, 4))

## MAE: 3.7217</pre>
```

The error reduction compared to a simple multivariate regression is indeed significant both in terms of MSE and MAE.