



MACHINE LEARNING TOOLBOX

Reusing a `trainControl`



A real-world example

- The data: customer churn at telecom company
- Fit different models and choose the best
- Models must use the same training/test splits
- Create a shared `trainControl` object

Example: customer churn data

```
# Summarize the target variables  
> library(caret)  
> library(C50)  
> data(churn)  
> table(churnTrain$churn) / nrow(churnTrain)
```

```
      yes      no  
0.1449145 0.8550855
```

```
# Create train/test indexes  
> set.seed(42)  
> myFolds <- createFolds(churnTrain$churn, k = 5)
```

```
# Compare class distribution  
> i <- myFolds$Fold1  
> table(churnTrain$churn[i]) / length(i)
```

```
      yes      no  
0.1441441 0.8558559
```

Example: customer churn data

```
> myControl <- trainControl(  
  summaryFunction = twoClassSummary,  
  classProbs = TRUE,  
  verboseIter = TRUE,  
  savePredictions = TRUE,  
  index = myFolds  
)
```

- Use folds to create a `trainControl` object
- Exact same cross-validation folds for each model



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Let's practice!



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Reintroduce `glmnet`



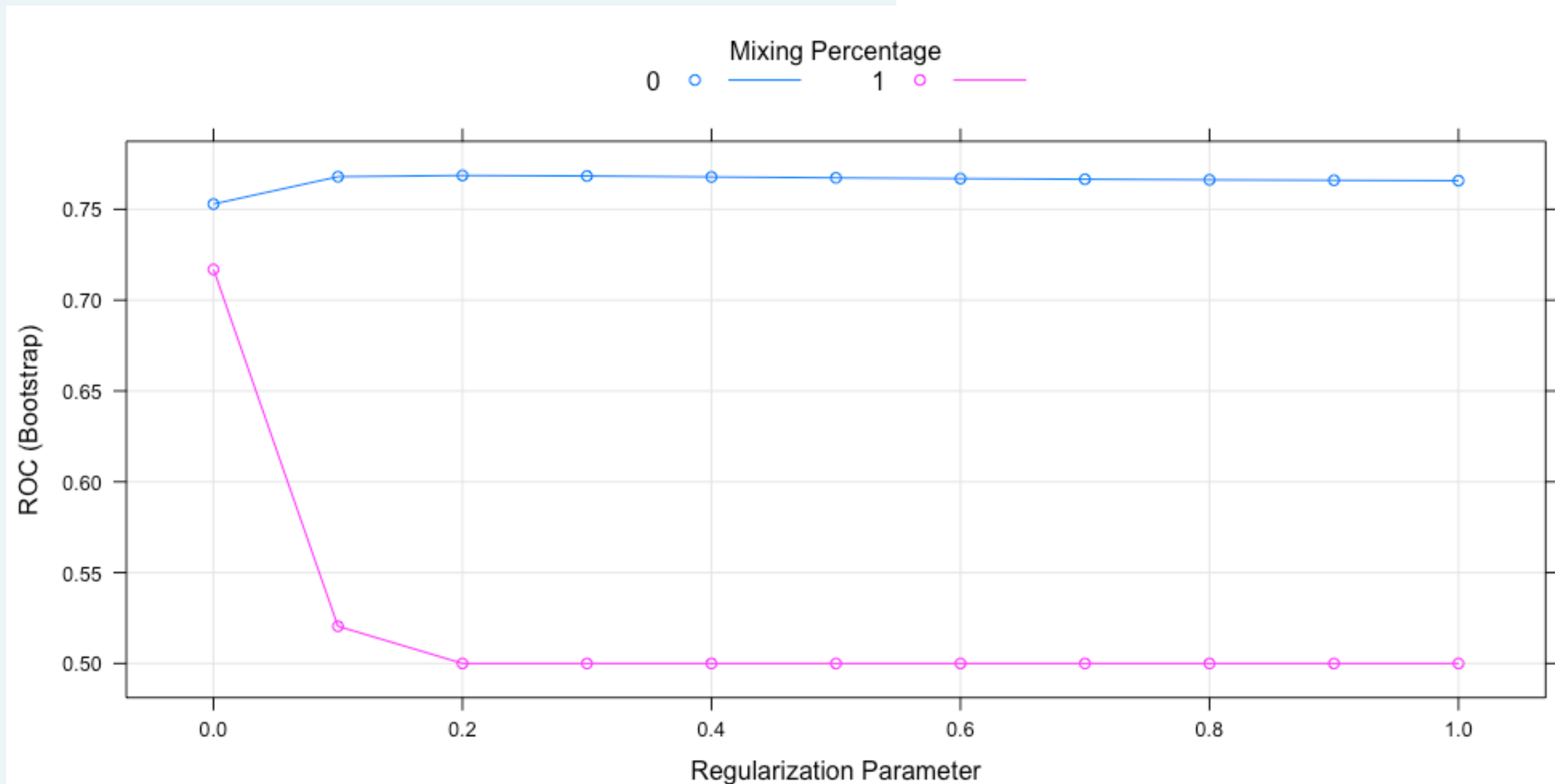
glmnet review

- Linear model with built-in variable selection
- Great baseline model
- Advantages
 - Fits quickly
 - Ignores noisy variables
 - Provides interpretable coefficients

Example: glmnet on churn data

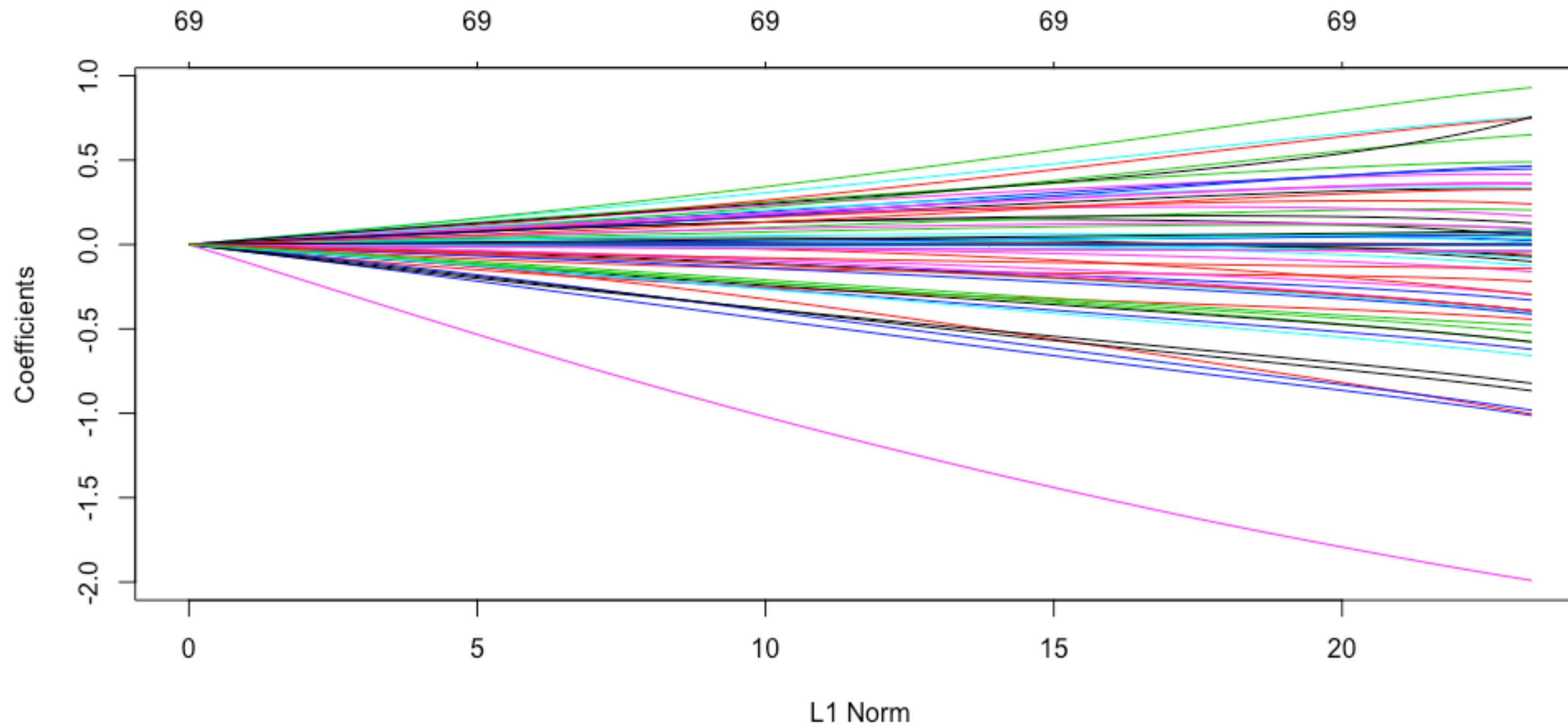
```
> # Fit the model
> set.seed(42)
> model_glmnet <- train(
  churn ~ ., churnTrain,
  metric = "ROC",
  method = "glmnet",
  tuneGrid = expand.grid(
    alpha = 0:1,
    lambda = 0:10/10
  ),
  trControl = myControl
)

> # Plot the results
> plot(model_glmnet)
```



Plot the coefficients

```
> plot(model_glmnet$finalModel)
```





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Reintroduce random forest



Random forest review

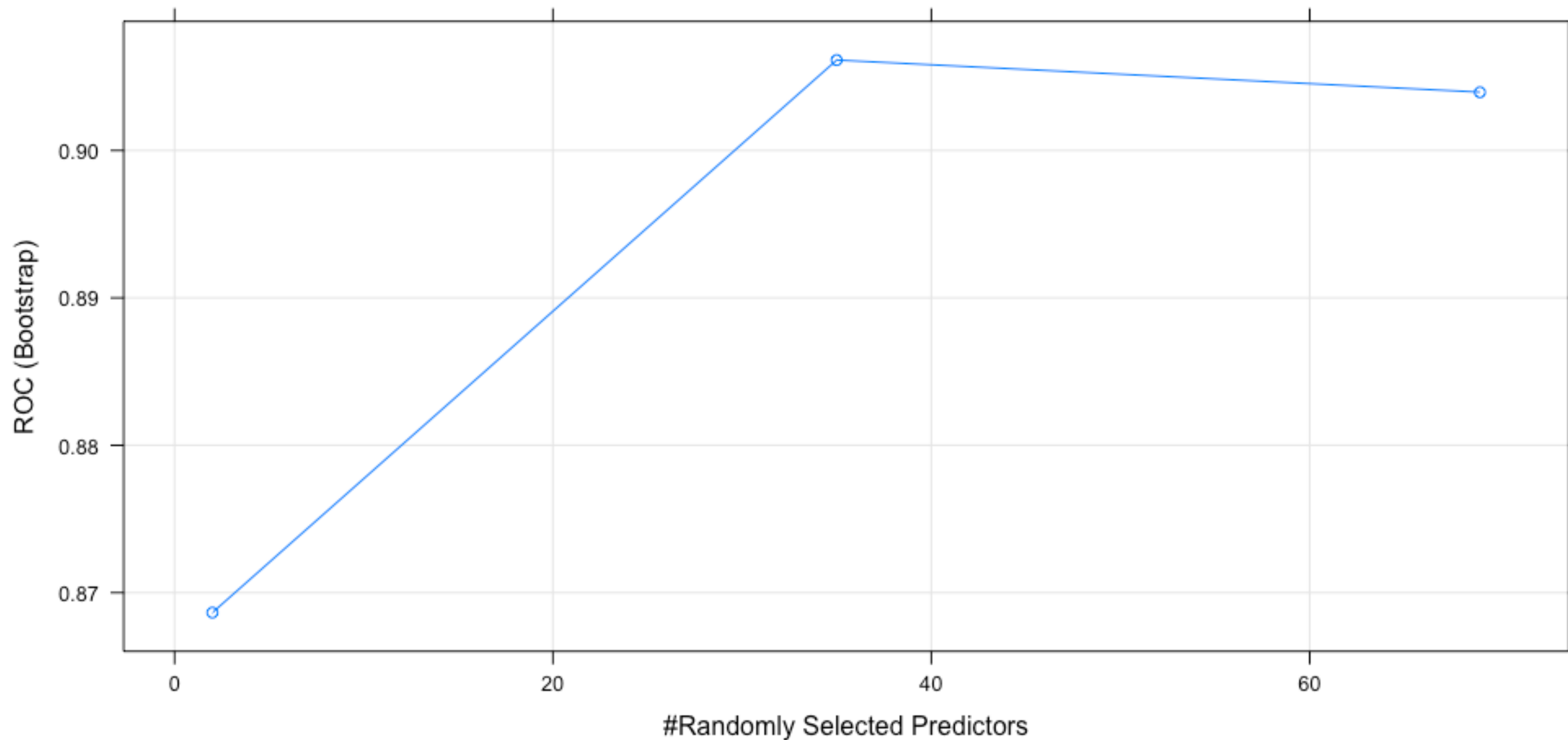
- Slower to fit than `glmnet`
- Less interpretable
- Often (but not always) more accurate than `glmnet`
- Easier to tune
- Require little preprocessing
- Capture threshold effects and variable interactions

Random forest on churn data

```
> set.seed(42)
> churnTrain$churn <- factor(churnTrain$churn, levels = c("no", "yes"))
> model_rf <- train(
  churn ~ ., churnTrain,
  metric = "ROC",
  method = "ranger",
  trControl = myControl
)
```

Random forest on churn data

```
> plot(model_rf)
```





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Comparing models



Comparing models

- Make sure they were fit on the same data!
- Selection criteria
 - Highest average AUC
 - Lowest standard deviation in AUC
- The `resamples()` function is your friend

Example: `resamples()` on churn data

```
# Make a list
> model_list <- list(
  glmnet = model_glmnet,
  rf = model_rf
)

# Collect resamples from the CV folds
> resamps <- resamples(model_list)
> resamps

Call:
resamples.default(x = model_list)

Models: glmnet, rf
Number of resamples: 5
Performance metrics: ROC, Sens, Spec
Time estimates for: everything, final model fit
```



Summarize the results

```
# Summarize the results
> summary(resamps)
Call:
summary.resamples(object = resamps)
```

```
Models: glmnet, rf
Number of resamples: 5
```

```
ROC
```

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	NA's
glmnet	0.7526	0.7624	0.7719	0.7686	0.7722	0.7840	0
rf	0.8984	0.9028	0.9077	0.9061	0.9093	0.9125	0



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More on resamples

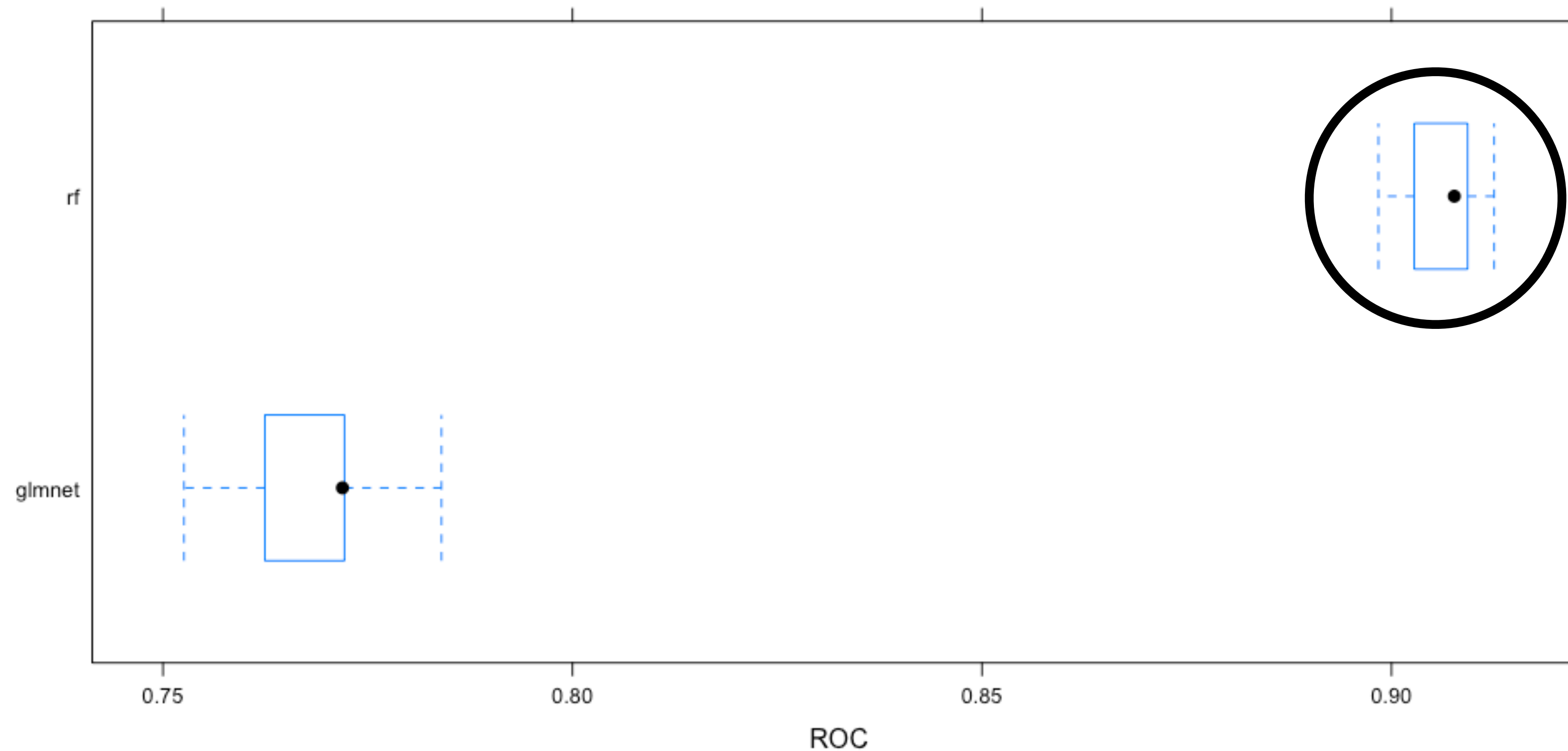


Comparing models

- Resamples has tons of cool methods
- One of my favorite functions (thanks Max!)
- Inspired the `caretEnsemble` package

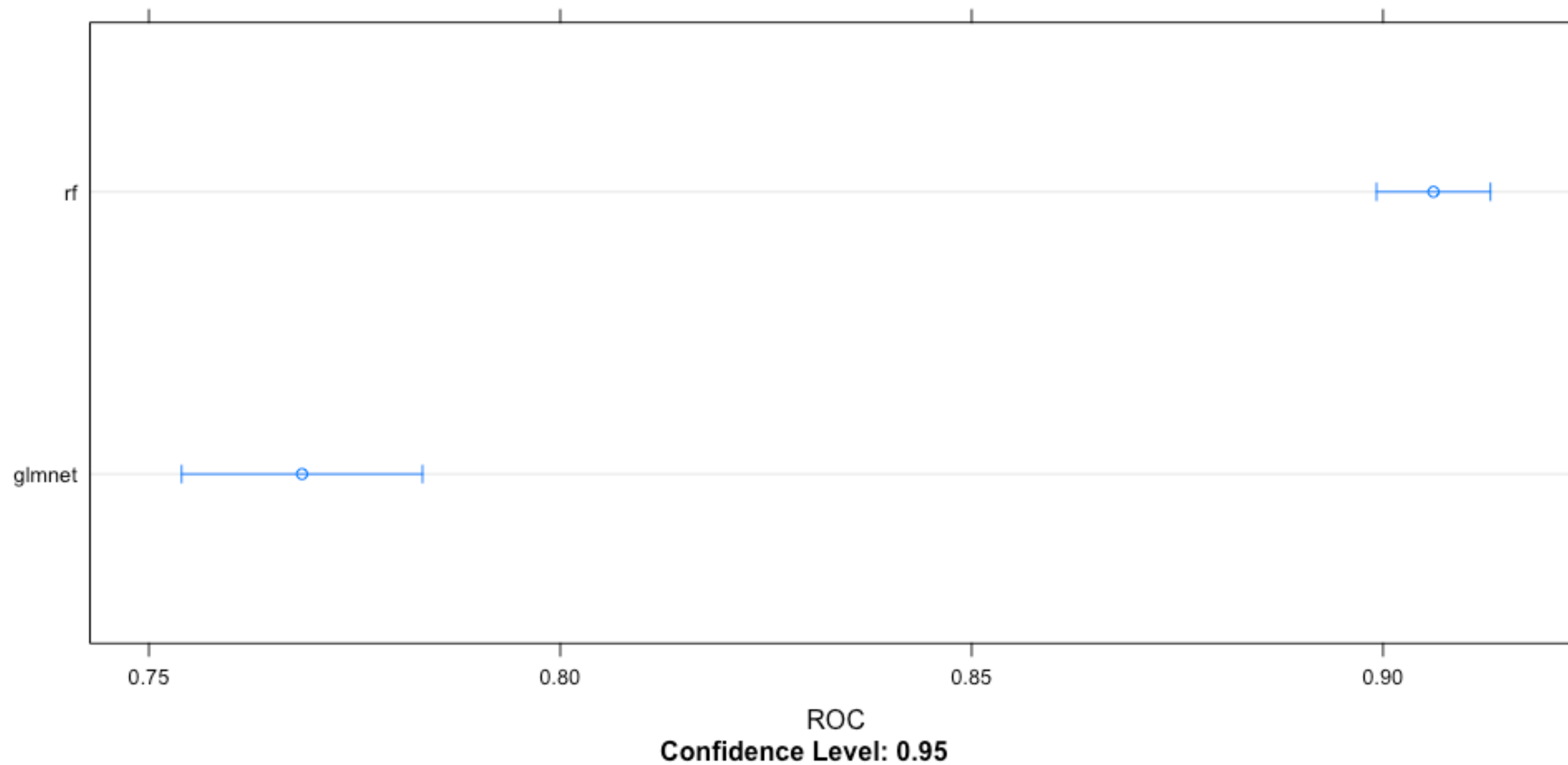
Box-and-whisker

```
> bwplot(resamps, metric = "ROC")
```



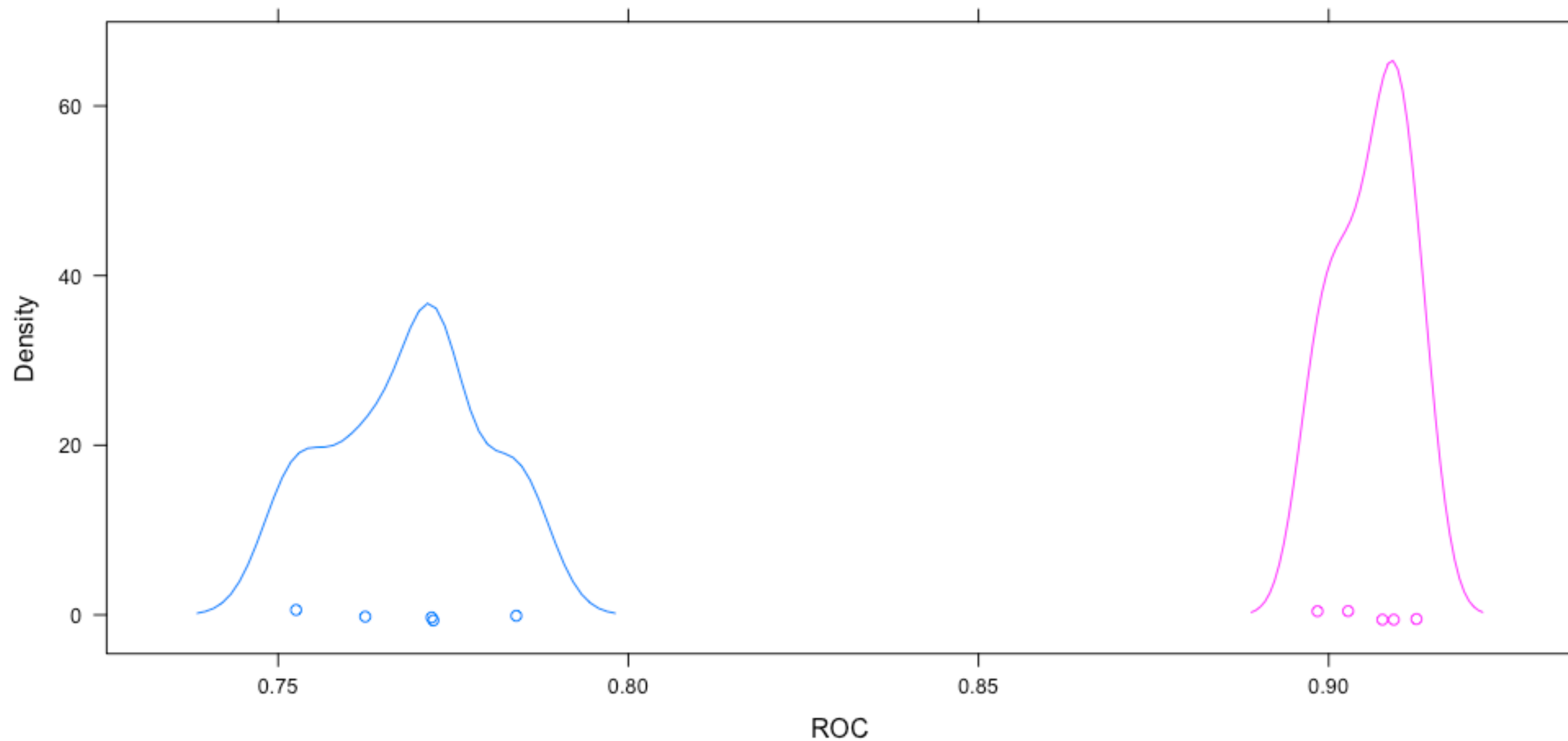
Dot plot

```
> dotplot(resamps, metric = "ROC")
```



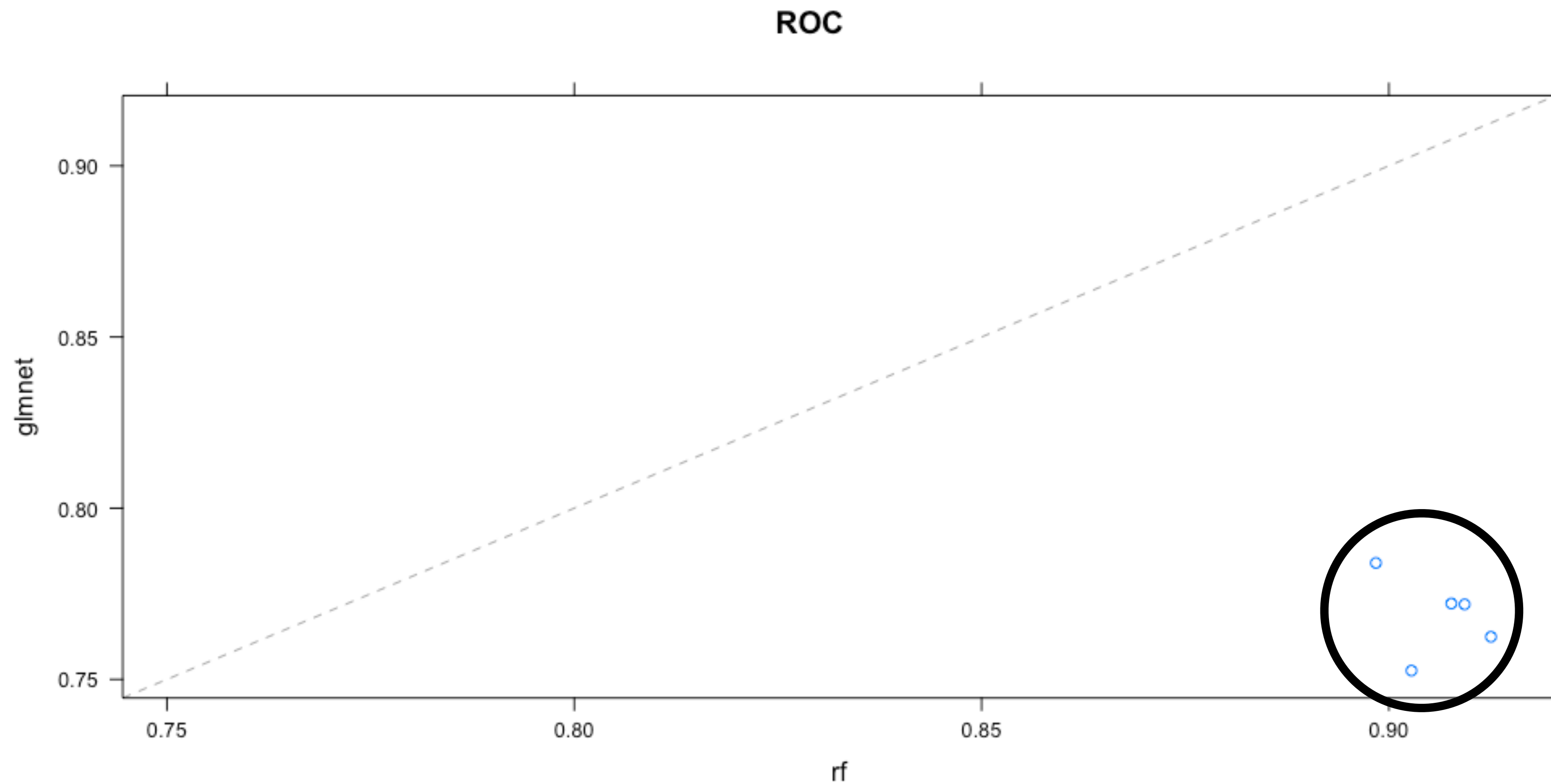
Density plot

```
> densityplot(resamps, metric = "ROC")
```



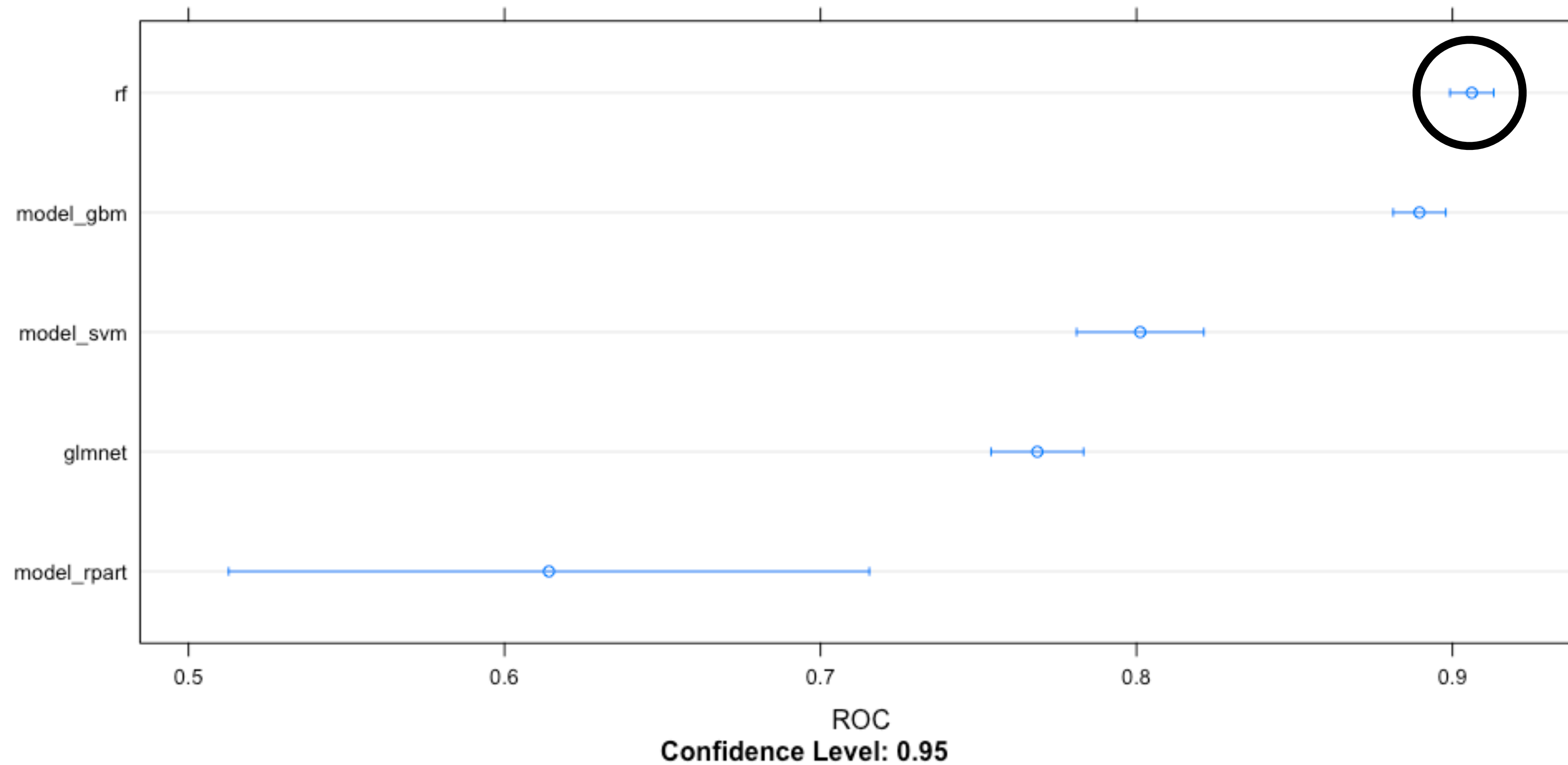
Scatter plot

```
> xyplot(resamps, metric = "ROC")
```



Another dot plot

```
> dotplot(lots_of_models, metric = "ROC")
```





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Summary

What you've learned

- How to use the caret package
- Model fitting and evaluation
- Parameter tuning for better results
- Data preprocessing

Goals of the caret package

- Simplify the predictive modeling process
- Make it easy to try many models and techniques
- Provide common interface to many useful packages



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Go build some models!