Assignment 2:

D Possible to do slightly better than decreven trees for tuma, but not by much 2) Ovafitting with small K, <u>maybe</u> (depends on noise, intrinsic dimensionality of data) 3) KNN is grite slow if distance evaluation is alar u) In large dimensions, reduces start to look very outhogonal to one another. 5) Lower dimensionality makes it harder to overfit 6) Maybe, if you are concerned about overfitting or noise How do the answers to 2) and 4) connect to one another? What about 2) and 6), and 4) and 5) Agericus-Lepiota has higher dimensionality then primary tume, but higher accuracy as well ?!

Quiz

1) Why is leave-one-out cross validation practical For K-NN classifiers and impractical for other settings?

2) For any bootstrap sample, what's the expected number of times any element of the original sample will be chosen?

In the limit of an infinitely large sample, what is the probability that any given element of the sample is selected?

Probability that a sample ran't selected in one of the n drews: $\frac{n-1}{n}$

Probability it will alkated in any:

 $\lim_{n \to \infty} \left(\frac{n^{-1}}{n} \right)^{-1} = \left(\left(-\frac{1}{n} \right)^{n} = e^{-1} \right)^{-1}$

Probability it is soledal at least once: 1-1:0.63

Precision, Becall, Sensitivity, Oh my! Acc = AtD Confusion matrix ATBECTD $\frac{\hat{y}=0}{y=0} \quad \frac{\hat{y}=1}{B}$ $\frac{\hat{y}=0}{Y=1} \quad C \quad D$ T = C + DS = B + DI = D $\frac{Precision}{S} = \frac{I}{S} = \frac{D}{BtD}$ 7I = A 7 S= AtC $\frac{\text{Becall}}{T} = \frac{D}{C+D}$ Sensitivity = healt = D CHD Specificity = $\frac{\neg I}{\neg S} = \frac{A}{A+C}$

Cross Velidation - Validation data is used inofficiently - Let's partition training data mto K subsets then create K training - validation splits - The average of the validation error is a better estimate of test error - "K-fold cross validation" - Use this to decide on hyperparameters - What if K increases to ω^{γ} ĆV -leave-one-at -Only useful w not borribly methicia,t when Algorithm 9 KNN-TRAIN-LOO(D) - K NN $:: err_k \leftarrow o, \forall 1 \le k \le N - 1$ // err_k stores how well you do with $k \operatorname{NN}$ 2: for n = 1 to N do $S_m \leftarrow \langle || \mathbf{x}_n - \mathbf{x}_m ||, m \rangle, \forall m \neq n$ // compute distances to other points $S \leftarrow \text{sort}(S)$ // put lowest-distance objects first $\hat{\mathbf{y}} \leftarrow \mathbf{o}$ // current label prediction for k = 1 to N - 1 do $\langle dist, m \rangle \leftarrow S_k$ $\hat{y} \leftarrow \hat{y} + y_m$ // let kth closest point vote if $\hat{y} \neq y_m$ then $err_k \leftarrow err_k + 1$ // one more error for kNN end if end for 13: end for ^{14:} **return** $\operatorname{argmin}_k err_k$ // return the K that achieved lowest error

Bootstrap Besampling (Statistical Testing) Then is 8% really greater than 7.9%? -fundamental new idea: error estimation matters - When -Null hypothesis testing asks - How likely is it that we saw this difference because of chance? - If we could repeat the experiment infinited many times, how often would we have made a mistake? Instead of reporting "b% on our population", we report (an estimate of) the distribution by rerunning and experiment on (many estimates of, resamples of the population **Algorithm 10** BOOTSTRAPEVALUATE(y, \hat{y} , NumFolds) 1: $scores \leftarrow []$ 2: for k = 1 to NumFolds do $truth \leftarrow []$ // list of values we want to predict pred \leftarrow [] // list of values we actually predicted for n = 1 to N do 5: $m \leftarrow$ uniform random value from 1 to N // sample a test point 6: $truth \leftarrow truth \oplus y_m$ // add on the truth 7: pred \leftarrow pred $\oplus \hat{y}_m$ // add on our prediction 8: end for $scores \leftarrow scores \oplus \mathbf{F-score}(truth, pred)$ // evaluate 10. III: end for 12: **return** (MEAN(scores), STDDEV(scores))