105 tumors

1 Randomly partition into thirds

2 • Use 2/3 to rank PTEN status discriminating genes by MW *P*-value and train 1 SVM

3 • Use 1/3 as blind test set for PTEN status prediction by trained SVM

4 Repeat 2X (such that each third is predicted once)

5 Repeat 9X (with new random partition into thirds)

Thus, in total there are 30 MW-ranked gene lists, 30 SVMs trained and each tumor is classified by 10 SVMs