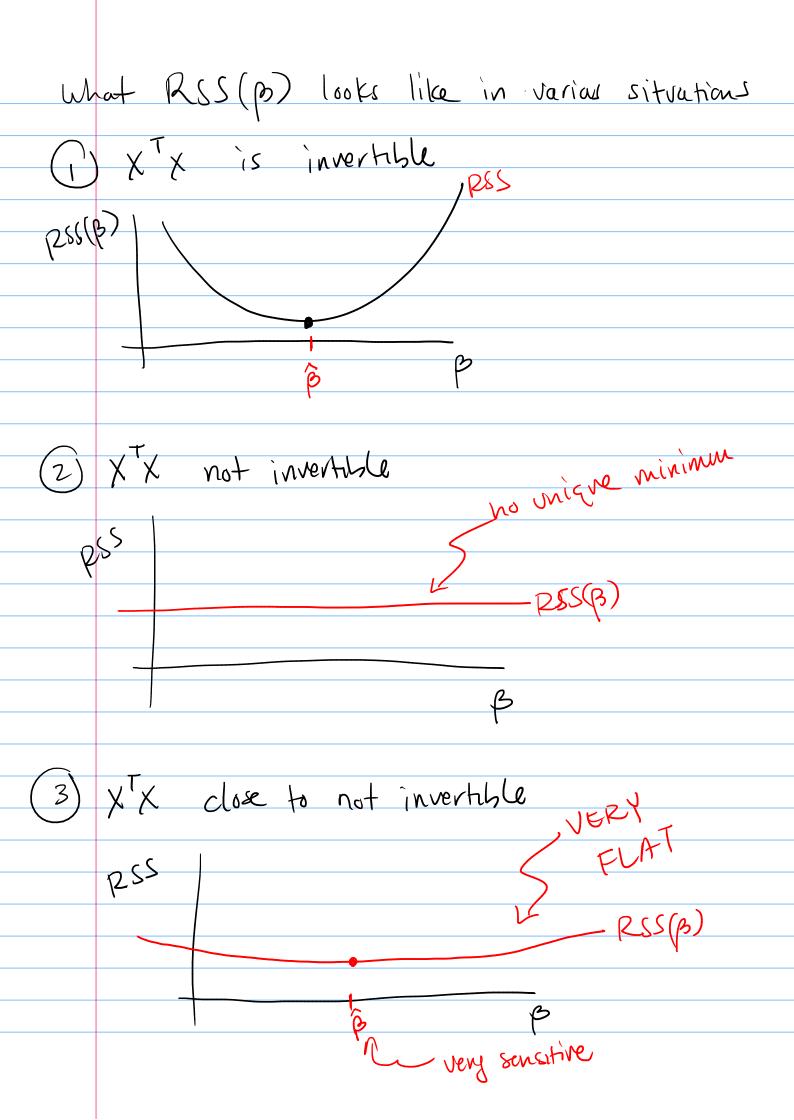
	Lecture 4: More Linear Regression + KNN
Gene	crically I can encode a K-terel factor ving K-1 dynny vars,
	12-1 acomong acrs,
£x,	Variable = Hogwarts House = SG, R, H, S3
	$data = \begin{array}{c} H \\ C \\$
Haw	do I interpret coefs?
	Coef assoc. v) H dummy is
d	holding other vars constant, what is predicted diff botwn H and S"
Filti	ng issues:
	all that I get is by setting aRSS = 0
	which yields my normal egus:
	$X^T X B = X Y$
-	F XTX is invertible, ther $\beta = (x^T x)^T Y$
	when can this fail?

XTX isn't invertible (=> rank(X) <# cols of X Pf. = Acome rank(X) < # cols. Then JU + O where XU = O. Thus $X^TXv=0$. So JU +0 when (XTX) U=0 i.l. ronk(XX) < # cals of XX i.l. its not invertible. => If XX isnt invertible then JU to when $\chi^T \chi \nu = 0$. for $\alpha / \|\alpha\|^2 = \alpha^T \alpha$ thus $0 = V^TO = V^TX^TXV = (XV)'(XV)$ $= \|Xv\|^2$ $= \|xv\|^2$ 30 po where XV = 0, i.e. X is rank deficient.

When does this happen in reality! 1) Accidentially include a var. twice in my design $X = \begin{bmatrix} 1 & 2 & 2 \\ 1 & 3 & 3 \\ 1 & 4 & 4 \end{bmatrix}$ If one of my vars is a LC of the $X = \begin{bmatrix} 1 & 0 & 1 \\ 1 & 1 & 0 \\ 1 & 0 & 1 \end{bmatrix}$ Intercept = M+F If # cols of X 7 # ohs. Ex. I measure 50,000 genes for each of 30 putier 4) This can approx! happen if some vars are almost a LC of each other Ex. two vers. are highly correlated



KNN Regression (K nearest neighbors regression) For LR we have a really strong global assurption about the form of f f(1) = x b 1 f(x)= /3+/3,X ers. in I-D Predictions follow linear for in entire space Training data affects fit very far away Benefit! strong global assumption makes f practial to find (opt. B over a p-dimil space) KNN makes a weaker local assumption abot the form of f where f(x) only depend on nearby training pts In particular $f(x) = \frac{1}{K} \sum_{n \in N_K(x)} ay$, yn over K as $f(x) = \frac{1}{K} \sum_{n \in N_K(x)} ay$, yn over K

