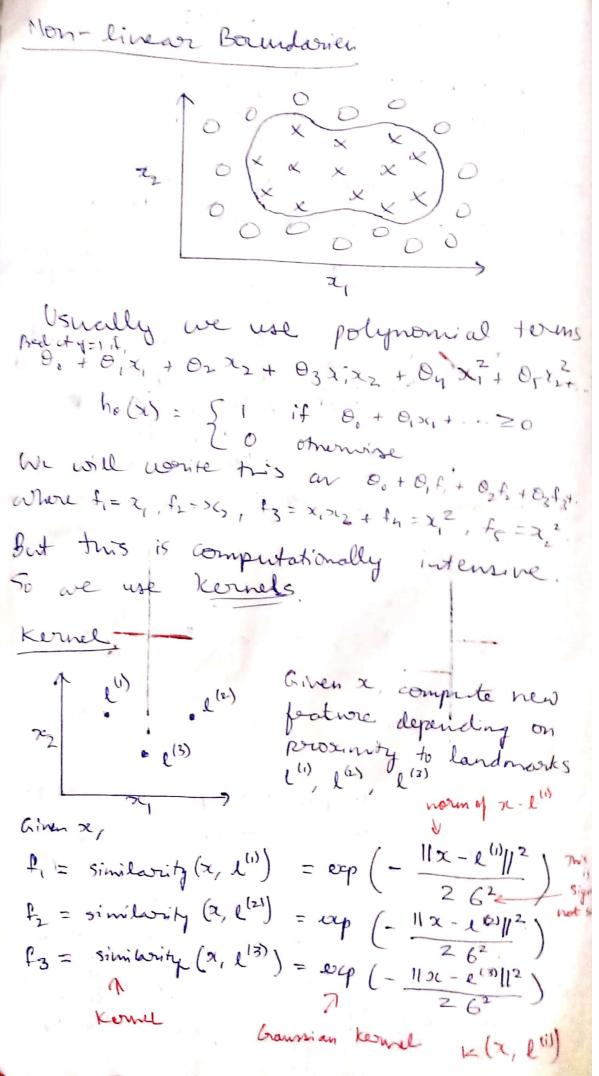


my= == = = = 101125 (i). ||o|| ≥ 1 if y(i) = 1 p(i) . 11011 ≤-1 it y(i) = 0 counder 2 cases to see how sun fuch the decision boundary: Let 0 = 0 This mean line needs to pan This is always nipio apparent 10 perpendular Let this O We can see here that bex(1) -> Co pli) and p(2) are small XE cet this :. For p(1). 11011 ≥ 61 L p(2) 1011 ≤ -1 : 1011 needs to be large but we we minimizing II Sime, so this line isn't good Notice here that p(1) and 1(2) are large : For p(1)-11011 21 11011 can be small p(2). || O|| ≤ -1, . This is what SVM! will choose Pl), p(2), p(3)... is brainally the margin Hore we have set of =0 and confined it to the origin, but its the same logic even it O to and the live doesn't pan twrough orpo.



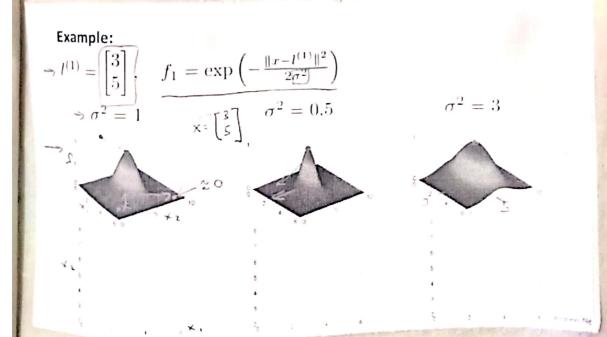
Lits take 
$$f_1$$
,

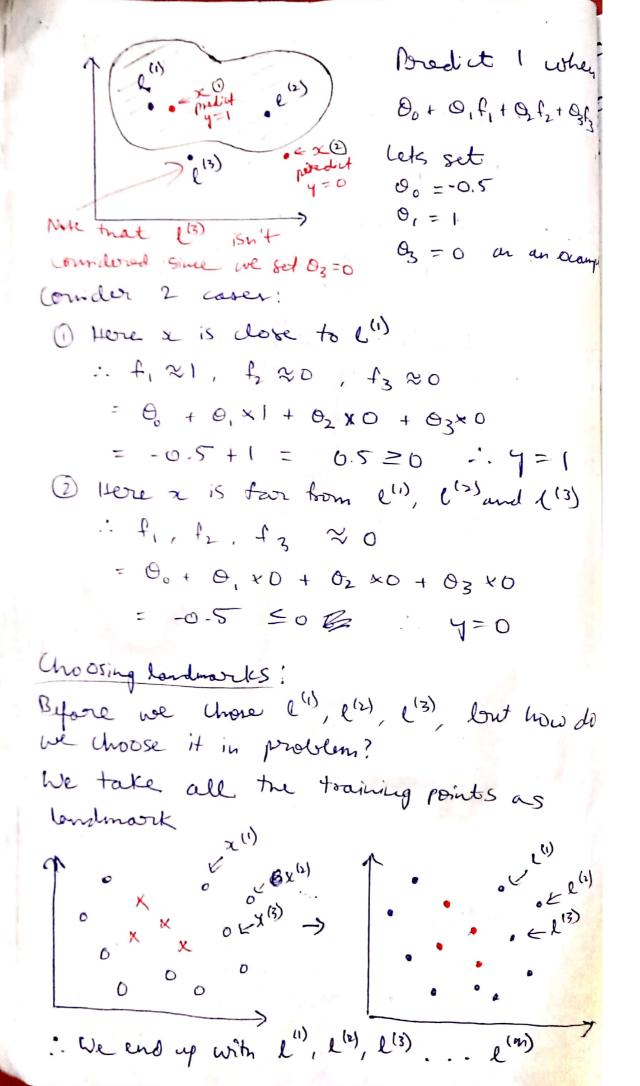
 $f_1$  similarity  $(x, \ell'') = \exp\left(-\frac{\|x - \ell''\|^2}{2\sigma^2}\right) = \exp\left(-\frac{\sum_{i=1}^{n}(2i - \ell_i')^2}{2\sigma^2}\right)$ 
 $f_1 \approx \exp\left(-\frac{O^2}{2G^2}\right) \approx 1$ 

If  $x = \exp\left(-\frac{(\log_2 number)^2}{2G^2}\right) \approx 0$ 

So each landmark proper a new feature  $f_1$ 
 $f_1 = \exp\left(-\frac{(\log_2 number)^2}{2G^2}\right) \approx 0$ 
 $f_2 = \exp\left(-\frac{(\log_2 number)^2}{2G^2}\right) \approx 0$ 

STICK PRINTOUT





en (x(1), y(1)), (x(2), y(2)), ... (x(m), y(m)) (hoose (ii) = x(i), (2) = x(2), ..., (x, y) liver example or, f, = similarity(21, l(1)) .. f= fz = similarity (x, l(2)) where fo = 1 (we set this like) . For toxing example (x(i), y(i)) + (i) = sim (x(i), (1))  $f_2(i) = \sin(\chi(i), L^{(2)})$   $f_2(i) = \sin(\chi(i), L^{(2)})$   $f_2(i) = \sin(\chi(i), L^{(2)}) = \exp(\frac{-U}{26^2}) = 1$ fm(i) = sim (x(i), l(m))  $f(i) = \begin{cases} f_0(i) \\ f_0(i) \end{cases}$  where  $f_0(i) = 1$ Sw with kernell Typotrais: Criver 2, compute features f EIRm+1 "Bredict "y=1" of Ott = 0 ( ) ( ) (ost, (0 + (i)) + (1 - y (i)) costo (0 + (i1)) + 1 > (0) Mathematically 7 this terms
of justend of is different. They use o'MO
is an abready built software but theats only to decrease
to minimize justed in it. Scanned with CamScanner o C (= 1) Small 6: Higher bias, low variance

Large a Features f; vary more smoothly

Thigher bias, lower variance

Small a Features f; vary len smoothly

I lower bicer, higher variance

## SVM in Practice

1 vse SVM software package (eg. 116/1 inear, 116 svm, ...) to solve for parameters O.

Ned to specify:

- Choice of parameter C - Choice of Kernel (similarity function):

G.g. O. No kernel (linear kernel):

bredict y=1 if ot> < >0

we use tris when n (no. of features)
is large but m (de no. of training
examples) is less.

D'Ganssian Kernel:

 $f_i = \exp\left(-\frac{||x-l(i)||^2}{2a^2}\right)$ , where  $l(i)=x^0$ 

- Ned to choose a2

and/or is large.

The SUM software package will ask you to implement the revuel (similarity) function for the kernel (x1, x2)

for exp(-\frac{11x1-x211^2}{2\alpha^2})

netwin

note: Perform poature scaling before using the human kernel.

The SVM Software package will then use this trunction to generate all the featwier f., fz, fs... fin and town the SVM.

Some SVM software packager will have inbuilt kernal functions and will allow you to choose

Ther choices of kernels:

- All available Kernels need to Satisfy

"Mercer's Theorem" to make Sure SVM

packages optimization run correctly

Polynomial Kernel: K(X, e) = (xTe + constant)

= (xTe)<sup>2</sup>, (xTe)<sup>3</sup>, (xTe+1)<sup>3</sup>.

More esotoric Kernel: Storing Kernel,

Un-Square Kernel, histogram intersection Kernel,...

Multidan clonification

- Marry SVM packager already have brilt in multi-clan classification functionality.
- Otherwise we one-vs-all method (Train K SW15, one to distinguish y=i bom the rest, for i= 1,2,..., A) get 0(1), 0(2), ..., O(K). Pick don i with y=1 y=2 y=x largest (00) 1x

Logistic regren on is. SUMs.

n= no. of feature, m= no. of training example

-) If is large (relative to in):

- Vie to l'egistic regrenion or sur without

of h=10,000 m=10 or 1000

The is small, in is intermediate:

- USE SUM with Grownian kornel -) If h is shall, m is large:

They boy - Then SVM will be slow.

perform so create add more peatures, they we almost by logistic regresion or SVM without a terme

-) Henral networks likely to work well for most of these settings, but may be Slower to train,