Linear kernel learning

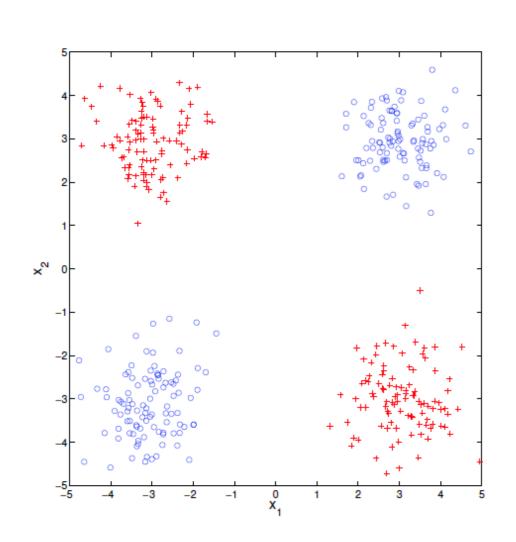
Prof. Samir Bhatt

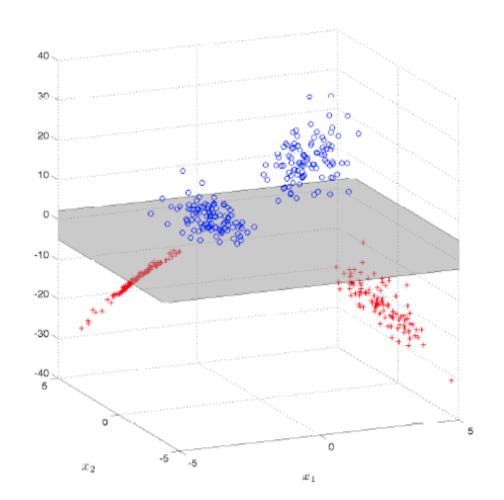
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The feature map





No linear classifier

$$(x_1, x_2) \rightarrow (x_1, x_2, x_1, x_2)$$

But we can linearise!



How to create high dimensional spaces?

Kernel methods allow construction of non linear methods after mapping to a higher dimensional feature space $\phi: X \to \mathbb{R}^D$

Typically rely on inner products $\phi(x_i)^T \phi(x_i)$

Its implicit - explicit features need not be computed

$$k(x_i, x_j) = \phi(x_i)^T \phi(x_j) = (1 + x_i^T x_j)^q$$
 with dimension $x \in \mathbb{R}^p$

Explicit construction would require
$$\begin{pmatrix} p+q\\q \end{pmatrix}$$

A 2d quadratic example
$$p = 2, q = 2$$
 is $\phi(x) = (1, \sqrt{2}x_1, \sqrt{2}x_2, \sqrt{2}x_1, x_2, x_1^2, x_2^2) \in \mathbb{R}^6$



Kernel as an inner product between feature maps

Let \mathcal{X} be a non empty set. A function $k:\mathcal{X}\times\mathcal{X}\to\mathbb{R}$ is a kernel if there exists a Hilbert space and a map $\phi:\mathcal{X}\to H$ such that for all $x,x'\in\mathcal{X},$

$$k(x, x') := \langle \phi(x), \phi(x') \rangle_H$$

This inner product can be thought of as a similarity between the various inputs

In a vector space the standard inner product is given by $\langle x, z \rangle = \sum_{i=1}^{n} x_i z_i$



If we are given a "measure of similarity" with two arguments, x, x', how can we determine if it is a valid kernel? What is the feature map? What if we want it to be infinite dimensional?

$$\Phi: \mathbf{x}_i \in \mathbb{R}^d \mapsto \Phi(\mathbf{x}_i) \in \mathbb{R}^D$$

$$\Phi: X \in \mathbb{R}^{N \times d} \mapsto \Phi(X) \in \mathbb{R}^{N \times D}$$

or

$$\Phi: \mathbf{x}_i \in \mathbb{R}^d \mapsto \Phi(\mathbf{x}_i) \in \mathbb{R}^\infty$$

$$\Phi: X \in \mathbb{R}^{N \times d} \mapsto \Phi(X) \in \mathbb{R}^{N \times \infty}$$

We can use a direct property - positive semidefiniteness (PSD)



Example of kernels

Kernel	Kernel Function, $k(\mathbf{x}_i, \mathbf{x}_j)$	Power Spectral Density, $\mathbb{P}(\boldsymbol{\omega})$
Squared Exponential [†] ◊	$\sigma^2 \exp\left(-\frac{(\mathbf{x}_i - \mathbf{x}_j)^2}{2\ell^2}\right)$	$(2\pi) - \frac{D}{2}\sigma^D \exp\left(\frac{\sigma^D \ \boldsymbol{\omega}\ _2^2}{2}\right)$
Matén* ^{†‡}	$\frac{2^{1-\lambda}}{\Gamma(\lambda)} \left(\frac{\sqrt{2\lambda \ \mathbf{x}_i - \mathbf{x}_j\ _2}}{\sigma} \right)^{\lambda} k_{\lambda} \left(\frac{\sqrt{2\lambda \ \mathbf{x}_i - \mathbf{x}_j\ _2}}{\sigma} \right)$	$\frac{2^{D+\lambda}\pi^{\frac{D}{2}}\Gamma(\lambda+\frac{D}{2})\lambda^{\lambda}}{\Gamma(\lambda)\sigma^{2\lambda}}\left(\frac{2\lambda}{\sigma^{2}}+4\pi^{2}\left\ \boldsymbol{\omega}\right\ _{2}^{2}\right)^{-(\lambda+\frac{D}{2})}$
Cauchy	$\prod_{i=1}^{D} \frac{2}{1+(\mathbf{x}_i-\mathbf{x}_j)^2}$	$\exp(-\ \boldsymbol{\omega}\ _1)$
Laplacian [†]	$\exp\left(-\sigma \left\ \mathbf{x}_i - \mathbf{x}_j\right\ _1\right)$	$\left(\frac{2}{\pi}\right)^{\frac{D}{2}} \prod_{i=1}^{D} \frac{\sigma}{\sigma^2 + \omega_i^2}$

^{*} $\Gamma(\cdot)$ is the gamma function and $k_{\lambda}(\cdot)$ is the modified Bessel function of the second kind. † Parameter $\sigma > 0$. ‡ *Parameter* $\lambda > 0$. \diamond *Parameter* $\ell > 0$.



Plot a kernel function

```
n = 100
x=seq(0,1,0.001) # distances
k1=function(x,alpha) exp(-x/alpha)
k2=function(x,alpha) exp(-x^2/alpha)
alpha=0.2
par(mfrow=c(1,2))
plot(x, k1(x, alpha), type='l')
plot(x, k2(x, alpha), type='l')
```

A PSD function

A given symmetric function $k: \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ is PSD is for all $n \geq 1$ and a set of coefficients $a_1, \ldots, a_n \in \mathbb{R}^n$ and $x_1, \ldots, x_n \in \mathcal{X}^n$

$$\sum_{i=1}^{n} \sum_{j=1}^{n} a_i a_j k(x_i, x_j) \ge 0 = \sum_{i=1}^{n} \sum_{j=1}^{n} \langle a_i \phi(x_i), a_j \phi(x_j) \rangle_H$$

Here the distance intuition is clear



Example of polynomial in a finite space

Our model is a linear function of x_1, x_2, x_1x_2 i.e $f(x) = f_1x_1 + f_2x_2 + f_3x_1x_2$

Therefore our mapping is $\mathcal{X}: \mathbb{R}^2 \to \mathbb{R}$, and we say $f(\cdot) = [f_1, f_2, f_3]$

 $f(\cdot)$ is a function object (vector in \mathbb{R}^3) and f(x) is an evaluation (real number)

So $f(x) = f(\cdot)^T \phi(x) = \langle f(\cdot), \phi(x) \rangle_H$ which means evaluations of fat x are an inner product space. And $H:\mathbb{R}^2\to\mathbb{R}$

Reproducing kernel Hilbert Spaces

H is a Hilbert space of $\mathbb R$ valued functions on a non empty $\mathcal X$. A function $k:\mathcal X\times\mathcal X\to\mathbb R$ is a reproducing kernel of H and H is a RKHS if

- 1) For all $x \in \mathcal{X}, k(\cdot, x) \in H$
- 2) For all $x \in \mathcal{X}$ and for all $f \in H$, $\langle f(\cdot), k(\cdot, x) \rangle_H = f(x)$

Why does this matter? Members of say L_2 spaces can't be evaluated because you can change the value at one point without changing the value of the integral (Lebesgue).

An RKHS is associated with a kernel that reproduces every function in the space in the sense that for any x in the set on which the functions are defined, and evaluation at x can be performed by taking an inner product with a function determined by the kernel.



Let simulate some data from this

We can say that we have sample from a multivariate normal with zero mean and covariance = our kernel matrix

Then z = N(0,1)A where $AA^T = K$ via the Cholesky decomposition

Note this decomposition as it will be relevant later when we look at Fourier features



Let simulate some data from this

```
n=2.00
x = seq(0, 1, length.out = n)
alpha=0.1
k1=function(x,alpha) exp(-x/alpha)
k2=function(x,alpha) exp(-x<sup>2</sup>/alpha)
K1=k1 (as.matrix (dist(x)), alpha)
K2=k2 (as.matrix (dist(x)), alpha)
par(mfrow=c(2,2))
image (K1)
image (K2)
K1 = chol(K1 + diag(1e-8, n))
K2 = chol(K2 + diag(1e-8, n))
gamma = cbind(rnorm(n))
plot(x,K1%*%gamma,type='l',col='red',lwd=5)
plot(x, K2%*%gamma, type='l', col='blue', lwd=5)
```



Can we learn? The representer theorem

Standard supervised learning set up $(x_1, y_1), \dots, (x_n, y_n), x \in \mathbb{R}^d, y \in \mathbb{R}$

We want to find a function f^* in our RKHS H that solves this minimisation of risk

$$\min_{f \in H} \sum_{i=1}^{n} \mathcal{L}(y_i, f(x_i)) + \Omega(\|f\|_h^2)$$

Obvious example for regression is $\mathcal{L}(y, f(x)) = (y - f(x))^2$

The solution is
$$f^* = \sum_{i=1}^n \alpha_i k(\cdot, x_i)$$



Least squares!

by setting X=K(X,X) we can find f^* for a squared loss by least squares and identify coefficients γ

$$\mathcal{L}(y - \beta X^T) = \frac{1}{n} (\beta X^T - y)^2$$

$$\nabla \mathcal{L}(y - \beta X^T) = \frac{2}{n} X^T (\beta X^T - y) = 0$$

$$X^T X \beta = X^T y$$

$$\beta = (X^T X)^{-1} X^T y$$

Fitting on simulated data

```
n = 200
x = seq(0, 1, length.out=n)
alpha=0.1
k1=function(x,alpha) exp(-x/alpha)
K1=k1 (as.matrix (dist(x)), alpha)
K1 = chol(K1 + diag(1e-8, n))
gamma = cbind(rnorm(n))
z=K1%*%qamma
f=lm.fit(x=K1,y=z)
par(mfrow=c(2,1))
plot(z,f$fitted.values,pch=16)
abline (0, 1, col='red')
plot (gamma, f$coefficients, pch=16)
abline (0, 1, col='red')
```



Not full rank (collinearity)

$$(X^T X)^{-1}$$
 can be singular & non invertible

How do we prevent overfitting?



Ridge regression

$$\frac{1}{2}(y - \gamma X^{T})^{2} + \frac{1}{2}C \parallel \gamma \parallel^{2}$$

$$\parallel \gamma \parallel^{2} := \sqrt{\gamma_{1}^{2} + \gamma_{2}^{2} + \dots + \gamma_{d}^{2}}$$
Squared loss function

Favour small norms (preventing over complex functions)

Adding some bias (to reduce variance)

So least squares becomes

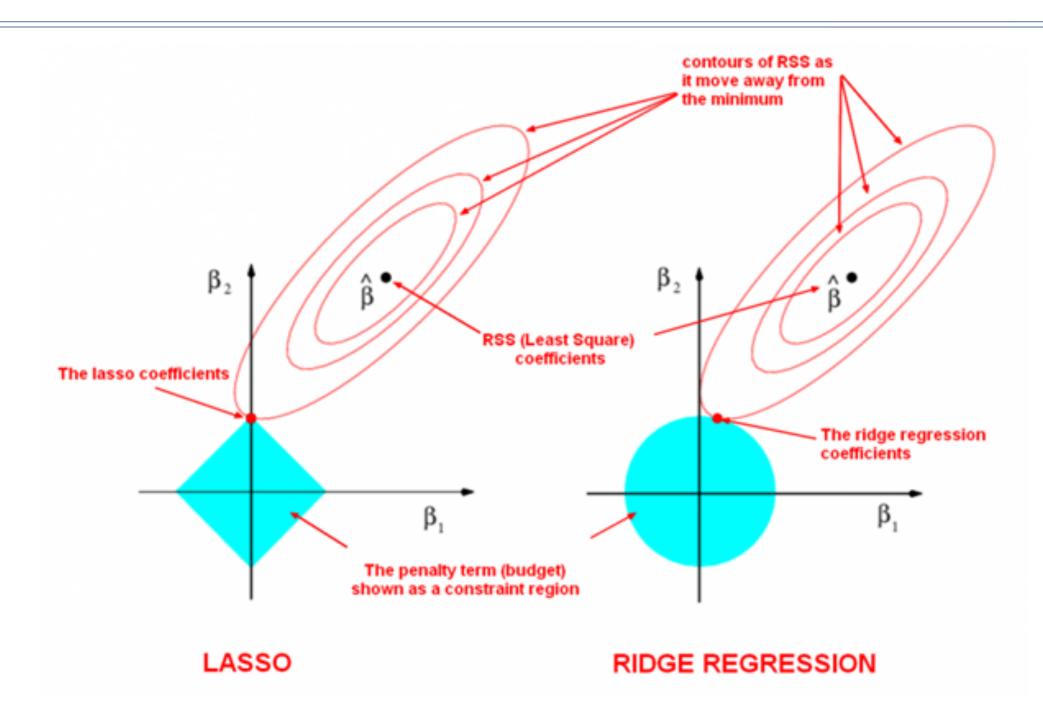
$$\nabla \mathcal{L}(y - \beta X^T) = X^T (\beta X^T - y) + \lambda \beta = 0$$

$$\beta = (X^T X + \lambda I_n)^{-1} X^T y$$

Basically the same but with this term added on the diagonal



What this is doing geometrically





Lets fit ridge regression

```
n = 200
x = seq(0, 1, length.out=n)
alpha=0.1
k1=function(x,alpha) exp(-x/alpha)
K1=k1 (as.matrix (dist(x)), alpha)
K1 = chol(K1 + diag(1e-8, n))
gamma = cbind(rnorm(n))
z=K1%*%gamma
z = z + rnorm(n, 0, 1)
f=lm.fit(x=K1,y=z)
plot(x,z,pch=16)
lines (x, f$fitted.values, col='red')
lines(x,z,col='blue')
library(glmnet)
lambdas \leftarrow 10^seq(5, -5, length.out=100)
cv fit <- cv.glmnet(K1, z, alpha = 0, lambda = lambdas)
opt lambda <- cv fit$lambda.min
fit \leftarrow glmnet(K1, z , alpha = 0, lambda = opt lambda)
f2 \leftarrow predict(fit, s = opt lambda, newx = K1)
lines (x, f2, col='green', lwd=2)
```



What about Bayesian models?

Ridge regression penalty = N(0,C) prior

$$y \sim N(\mu, \sigma^2)$$

$$\mu = \gamma X^T$$

$$\gamma \sim N(0, C\mathbb{I})$$

$$C \sim \pi()$$

$$\lambda \sim \pi()$$

Or something better....

Sparsity information and regularization in the horseshoe and other shrinkage priors

Juho Piironen and Aki Vehtari

A Bayesian example

```
set.seed(123)
n = 50
x = seq(0, 1, length.out=n)
alpha=0.1
k1=function(x,alpha) exp(-x/alpha)
K1=k1 (as.matrix (dist(x)), alpha)
K1 = chol(K1 + diag(1e-8, n))
gamma = cbind(rnorm(n))
z=K1%*%gamma
z = z + rnorm(n, 0, 1)
f=lm.fit(x=K1,y=z)
plot(x,z,pch=16)
lines(x, f$fitted.values, col='red')
lines(x,z,col='blue')
library(glmnet)
lambdas \leftarrow 10^{seq}(5, -5, length.out=100)
cv fit <- cv.glmnet(K1, z , alpha = 0, lambda = lambdas)</pre>
opt lambda <- cv fit$lambda.min
fit \leftarrow glmnet(K1, z , alpha = 0, lambda = opt lambda)
f2 <- predict(fit, s = opt lambda, newx = K1)
lines (x, f2, col='green', lwd=2)
```

A Bayesian example



Lets take stock

- We have learned what kernels are and why they are amazing
- We can solve the risk minimisation problem via the representer theorem
- We realise that overfitting is a possibility and control for that via the ridge penalty

Whats the problem then?

- Complexity for linear regression $\mathcal{O}(nm^2)$ and therefore for kernels it becomes $\mathcal{O}(nm^2) \to \mathcal{O}(n^3)$
- Can we go back to $\mathcal{O}(nm^3)$ while still having an infinitely implied space?
- The answer amazingly is yes, but also kind of....

Random Fourier features (Rahimi and Recht 2009)

Theorem 1. (Bochner's Theorem) A stationary continuous kernel $k(x_i, x_j) = \kappa(x_i - x_j)$ on \mathbb{R}^d is positive definite if and only if $\kappa(\delta)$ is the Fourier transform of a non-negative measure.

Hence, for an appropriately scaled shift invariant complex kernel $\kappa(\delta)$, i.e. for $\kappa(0) = 1$, Bochner's Theorem ensures that its inverse Fourier Transform is a probability measure:

$$k(x_1, x_2) = \int_{\mathbb{R}^d} e^{i\omega^T (x_1 - x_2)} \mathbb{P}(d\omega). \tag{7}$$

Theorem 2. (Yaglom, 1987 [12,22]) A nonstationary kernel $k(x_1, x_2)$ is positive definite in \mathbb{R}^d if and only if it has the form:

$$k(x_1, x_2) = \int_{\mathbb{R}^D \times \mathbb{R}^D} e^{i(w_1^T x_1 - w_2^T x_2)} \mu(dw_1, dw_2)$$
(19)

where $\mu(dw_1, dw_2)$ is the Lebesgue-Stieltjes measure associated to some positive semi-definite function $f(w_1, w_2)$ with bounded variation.



Kernel	Kernel Function, $k(\mathbf{x}_i, \mathbf{x}_j)$	Power Spectral Density, $\mathbb{P}(\boldsymbol{\omega})$
Squared Exponential [†] ◊	$\sigma^2 \exp\left(-\frac{(\mathbf{x}_i - \mathbf{x}_j)^2}{2\ell^2}\right)$	$(2\pi) - \frac{D}{2}\sigma^D \exp\left(\frac{\sigma^D \ \boldsymbol{\omega}\ _2^2}{2}\right)$
Matén* ^{†‡}	$\frac{2^{1-\lambda}}{\Gamma(\lambda)} \left(\frac{\sqrt{2\lambda \ \mathbf{x}_i - \mathbf{x}_j\ _2}}{\sigma} \right)^{\lambda} k_{\lambda} \left(\frac{\sqrt{2\lambda \ \mathbf{x}_i - \mathbf{x}_j\ _2}}{\sigma} \right)$	$\frac{2^{D+\lambda}\pi^{\frac{D}{2}}\Gamma(\lambda+\frac{D}{2})\lambda^{\lambda}}{\Gamma(\lambda)\sigma^{2\lambda}}\left(\frac{2\lambda}{\sigma^{2}}+4\pi^{2}\left\ \boldsymbol{\omega}\right\ _{2}^{2}\right)^{-(\lambda+\frac{D}{2})}$
Cauchy	$\prod_{i=1}^{D} \frac{2}{1 + (\mathbf{x}_i - \mathbf{x}_j)^2}$	$\exp(-\left\ \boldsymbol{\omega}\right\ _1)$
Laplacian [†]	$\exp\left(-\sigma \left\ \mathbf{x}_i - \mathbf{x}_j\right\ _1\right)$	$\left(\frac{2}{\pi}\right)^{\frac{D}{2}} \prod_{i=1}^{D} \frac{\sigma}{\sigma^2 + \omega_i^2}$

^{*} $\Gamma(\cdot)$ is the gamma function and $k_{\lambda}(\cdot)$ is the modified Bessel function of the second kind. † Parameter $\sigma > 0$. † Parameter $\lambda > 0$. \diamond Parameter $\ell > 0$.

Gaussian
Student T
exponential
Cauchy



Random Fourier Features (Bochner)

$$k(x_1, x_2) = \int_{\mathbb{R}^D} e^{i\omega^T (x_1 - x_2)} \mathbb{P}(d\omega)$$

$$= \mathbb{E}_{\omega \sim \mathbb{P}} \left[e^{i\omega^T (x_1 - x_2)} \right],$$

$$= \mathbb{E}_{\omega \sim \mathbb{P}} \left[\cos(\omega^T (x_1 - x_2)) + i \sin(\omega^T (x_1 - x_2)) \right]$$

$$= \mathbb{E}_{\omega \sim \mathbb{P}} \left[\cos(\omega^T (x_1 - x_2)) \right]$$

$$= \mathbb{E}_{\omega \sim \mathbb{P}} \left[\cos(\omega^T (x_1 - x_2)) \right]$$

$$= \mathbb{E}_{\omega \sim \mathbb{P}} \left[\cos(\omega^T x_1) \cos(\omega^T x_2) + \sin(\omega^T x_1) \sin(\omega^T x_2) \right]$$

$$\approx \frac{1}{m} \sum_{k=1}^m \left(\cos(\omega_k^T x_1) \cos(\omega_k^T x_2) + \sin(\omega_k^T x_1) \sin(\omega_k^T x_2) \right)$$

$$= \frac{1}{m} \sum_{k=1}^m \Phi_k(x_1)^T \Phi_k(x_2)$$

where $\{\omega_k\}_{k=1}^m \stackrel{i.i.d.}{\sim} \mathbb{P}$ and we denoted

$$\Phi_k(x_l) = \begin{pmatrix} \cos(\omega_k^T x_l) \\ \sin(\omega_k^T x_l) \end{pmatrix}.$$



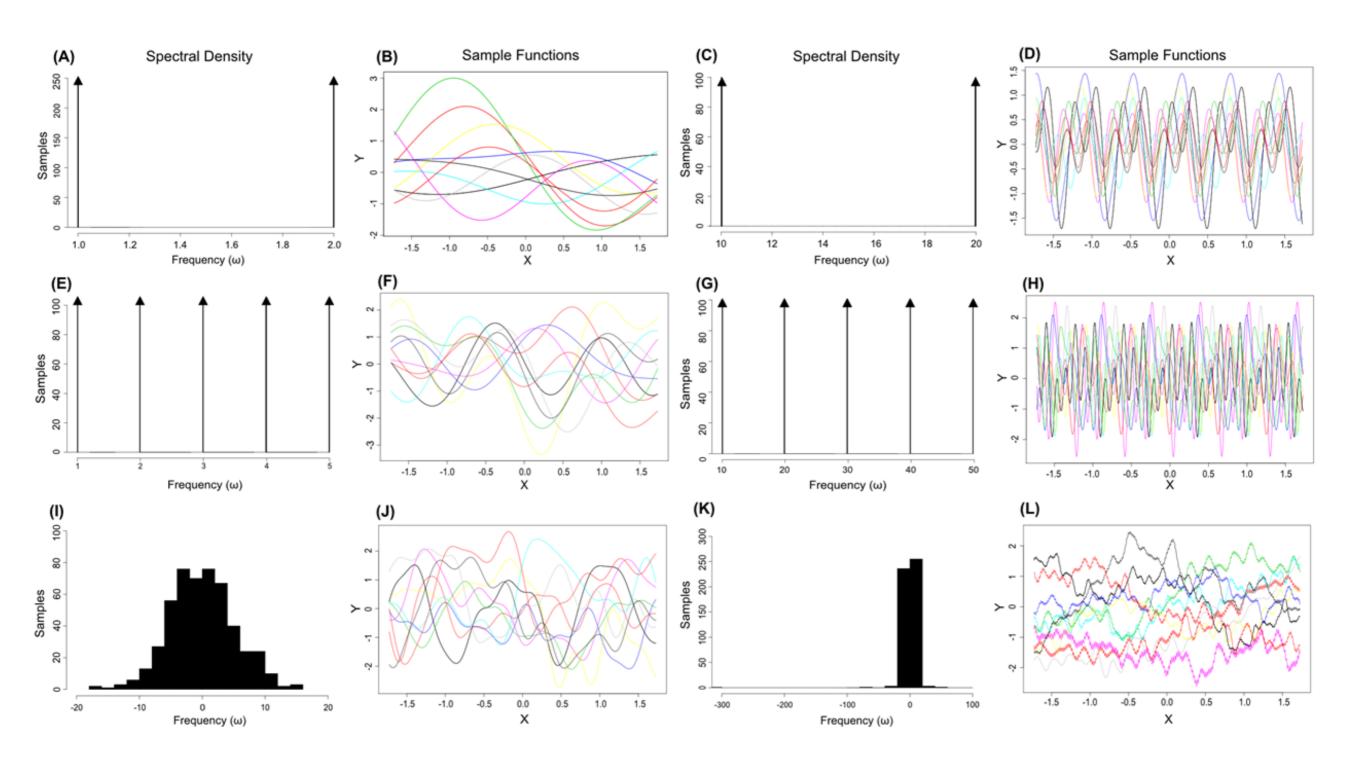
$$\begin{split} k(x_1,x_2) &= \frac{1}{4} \int_{\mathbb{R}^D \times \mathbb{R}^D} \left(e^{i(\omega_1^T x_1 - \omega_2^T x_2)} + e^{i(\omega_2^T x_1 - \omega_1^T x_2)} + e^{i(\omega_1^T x_1 - \omega_1^T x_2)} + e^{i(\omega_2^T x_1 - \omega_2^T x_2)} \right) \mu(d\omega_1 d\omega_2) \\ &= \frac{1}{4} \mathbb{E}_{\mu} \left(e^{i(\omega_1^T x_1 - \omega_2^T x_2)} + e^{i(\omega_2^T x_1 - \omega_1^T x_2)} + e^{i(\omega_1^T x_1 - \omega_1^T x_2)} + e^{i(\omega_2^T x_1 - \omega_2^T x_2)} \right) \\ &\approx \frac{1}{4m} \sum_{k=1}^m \left(e^{i(x_1^T \omega_k^1 - x_2^T \omega_k^2)} + e^{i(x_1^T \omega_k^2 - x_2^T \omega_k^1)} + e^{i(x_1^T \omega_k^1 - x_2^T \omega_k^1)} + e^{i(x_1^T \omega_k^2 - x_2^T \omega_k^2)} \right) \\ &= \frac{1}{4m} \sum_{k=1}^m \left\{ \cos(x_1^T \omega_k^1) \cos(x_2^T \omega_k^1) + \cos(x_1^T \omega_k^1) \cos(x_2^T \omega_k^2) \right. \\ &\quad + \cos(x_1^T \omega_k^2) \cos(x_2^T \omega_k^1) + \cos(x_1^T \omega_k^2) \cos(x_2^T \omega_k^2) \\ &\quad + \sin(x_1^T \omega_k^1) \sin(x_2^T \omega_k^1) + \sin(x_1^T \omega_k^1) \sin(x_2^T \omega_k^2) \\ &\quad + \sin(x_1^T \omega_k^2) \sin(x_2^T \omega_k^1) + \sin(x_1^T \omega_k^2) \sin(x_2^T \omega_k^2) \right\} \quad (taking \ the \ real \ part) \\ &= \frac{1}{4m} \sum_{k=1}^m \Phi_k(x_1)^T \Phi_k(x_2) \end{split}$$

where $\{(\omega_k^1, \omega_k^2)\}_{k=1}^m \overset{i.i.d.}{\sim} \mu$ and

$$\Phi_k(x_l) = \begin{pmatrix} \cos(x_l^T \omega_k^1) + \cos(x_l^T \omega_k^2) \\ \sin(x_l^T \omega_k^1) + \sin(x_l^T \omega_k^2) \end{pmatrix}.$$



What do the functions look like?



Lets have a look

```
set.seed(1234)
n=500
x=as.matrix(seq(-2,2,length.out=n))
x=scale(x) # remember to always scale
k = 2.00
# change this (the spectral density)
omega1<-t(cbind(c(rep(3,k))))
omega2 < -t(rnorm(k, 0, 5))
omega3<-t(rcauchy(k, 0, 2))
par(mfrow=c(3,2))
nsamp < -10
for(i in 1:4) {
   eval(parse(text=paste0("omega<-omega",i)))
   xxprojected <- x%*% (omega) # project
   f<- sqrt(1/k)*cbind(cos(xxprojected), sin(xxprojected)) # monte carlo bit
   K<-(f) %*%t(f) # recreate kernel matrix
   cK = chol(K + diag(1e - 6, n))
   samp<- matrix(rnorm(n*nsamp), nrow=nsamp, ncol=n) %*% cK</pre>
   samp<-t(samp)</pre>
   hist (omega, 20, col='black', main="Spectral Density")
   plot(x, samp[,1], type='l', ylim=c(min(samp), max(samp)), main="Samples")
   for (j in 1:nsamp) {
       lines(x, samp[, j], type='l', col=j)
```



Algorithm 1 Random Fourier features for nonstationary kernels

Input: spectral measure μ , dataset X, number of frequencies m

Output: Approximation to K_{xx}

Start Algorithm:

Sample m pairs of frequencies $\{(\omega_k^1, \omega_k^2)\}_{k=1}^m \stackrel{i.i.d.}{\sim} \mu$ giving Ω^1 and Ω^2

Compute
$$\Phi_x = \left[\cos(\mathbf{X}(\Omega^1)^T) + \cos(\mathbf{X}(\Omega^2)^T) \mid \sin(\mathbf{X}(\Omega^1)^T) + \sin(\mathbf{X}(\Omega^2)^T)\right] \in \mathbb{R}^{n \times 2m}$$

$$\widehat{K_{\mathbf{XX}}} = \frac{1}{4m} \Phi_x \Phi_x^T$$

End Algorithm

Spatial Mapping with Gaussian Processes and Nonstationary Fourier Features

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Or in code

```
set.seed(123)
n=500
x= cbind(runif(n),runif(n))
x=scale(x)
alpha=0.5
m=100
Omega = (cbind(rnorm(m),rnorm(m))) # Squared exponential kernel
Proj = x %*% t(Omega) # Projection - combine data with sample frequencies
Phi = sqrt(1/m) * cbind(cos(Proj/alpha), sin(Proj/alpha)) # Fourier feature
matrix
K = Phi %*% t(Phi) # approximation of Kernel matrix
d=as.matrix(dist(x))
K1<-exp(-0.5*d*d/(alpha^2))
plot(K1,K)
abline(0,1,col='red')</pre>
```



- The approximation of the covariance matrix is not great, and theoretical bounds suggest that this approach needs thousands of features
- Empirical evidence suggests otherwise, we need just a few hundred
- But theory has shown the Φ creates a provide a dense basis set for for an l_2 ball of l_2 functions
- Similar to uniform approximation of Neural networks



What is the generalisation bound

$$R(f^*) - R(\hat{f})$$

•
$$\mathcal{O}(\frac{1}{\sqrt{n}} + \frac{1}{\sqrt{D}})$$
 Rahimi and Recht

• $\mathcal{O}(\frac{log(n)}{\sqrt{n}})$ Rudi (under certain conditions as good as full ridge)



We can use Φ in linear regression

```
set.seed(123)
n = 500
x = seq(0, 1, length.out=n)
alpha=0.1
k1=function(x,alpha) exp(-x/alpha)
K1=k1(as.matrix(dist(x)),alpha)
K1 = chol(K1 + diag(1e-8, n))
gamma = cbind(rnorm(n))
z=K1%*%qamma
z = z + rnorm(n, 0, 1)
f=lm.fit(x=K1,y=z)
plot(x,z,pch=16,cex=0.5)
lines(x,z,col='blue')
library(glmnet)
lambdas \leftarrow 10^{seq}(5, -5, length.out=100)
cv fit <- cv.glmnet(K1, z , alpha = 0, lambda = lambdas)</pre>
opt lambda <- cv fit$lambda.min
fit \leftarrow glmnet(K1, z , alpha = 0, lambda = opt lambda)
f2 <- predict(fit, s = opt lambda, newx = K1)
lines(x,f2,col='green',lwd=2)
m = 200
Omega = cbind(rnorm(m)) # Squared exponential kernel
Proj = scale(x) %*% t(Omega) # Projection - combine data with sample frequencies
Phi = sqrt(1/m) * cbind(cos(Proj/alpha), sin(Proj/alpha)) # Fourier feature matrix
lambdas \leftarrow 10^s eq(5, -5, length.out=100)
cv_fit <- cv.glmnet(Phi, z_, alpha = 0, lambda = lambdas)</pre>
opt lambda <- cv fit$lambda.min
fit <- glmnet(Phi, z , alpha = 0, lambda = opt lambda)</pre>
f2 <- predict(fit, s = opt lambda, newx = Phi)
lines(x, f2, col='brown', lwd=2)
```



Why use RFFs?

- Easy to program
- Can create high dimensional kernels without too much loss (its all about integration)
- Its approximates the whole kernel and not just a low rank one based on some data
- It casts everything as linear regression which makes it easy to customise

How can we improve things?

Quasi-Monte Carlo integration - Halton Sequence

```
set.seed(123)
n = 500
x= cbind(runif(n), runif(n))
x=scale(x)
alpha=0.5
m = 100
d=as.matrix(dist(x))
K1 < -\exp(-0.5*d*d/(alpha^2))
Omega = cbind(rnorm(m), rnorm(m)) # Squared exponential kernel
Proj = x %*% t(Omega) # Projection - combine data with sample frequencies
Phi = sqrt(1/m) * cbind(cos(Proj/alpha), sin(Proj/alpha)) # Fourier feature matrix
K = Phi %*% t(Phi) # approximation of Kernel matrix
par(mfrow=c(1,2))
plot(K1,K)
abline (0, 1, col='red')
library(randtoolbox)
Omega qmc<-t(qnorm(halton(m,2)))</pre>
Omega qmc = cbind(rnorm(m), rnorm(m)) # Squared exponential kernel
Proj = x %*% t(Omega) # Projection - combine data with sample frequencies
Phi = sqrt(1/m) * cbind(cos(Proj/alpha), sin(Proj/alpha)) # Fourier feature matrix
K2 = Phi %*% t(Phi) # approximation of Kernel matrix
plot (K2, K)
abline (0, 1, col='red')
```





- We can train these models using mini-batch gradient descent
- Dropout can be amazing
- GPUs can be very efficient