USAGE NOTES FOR PACKAGE CHEBPOL

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ABSTRACT. Here are some examples of how package chebpol is used, with description of differences between the methods.

1. Introduction

Interpolating a function f means to find a reasonably behaved function, an interpolant g which agrees with f on some points. I.e. given $N \in \mathbb{N}$ we have $\{x_i\}_{i=1}^N$ in the domain of f, and the interpolant g satisfies $g(x_i) = f(x_i)$ for $1 \le i \le N$. The points x_i are often referred to as knots.

We limit ourselves to functions $\mathbb{R}^n \to \mathbb{R}$, or even compactly supported functions $[-1,1]^n \to \mathbb{R}$.

Which method to use depends first and foremost on one thing. Can we choose the points x_i freely? If n > 1, can we choose the points as a grid, i.e. a Cartesian product of single-dimension points?

Say we have a function f(x, y) defined in a square, i.e. for $\{(x, y) \mid -1 \le x \le 1 \text{ and } -1 \le y \le 1\}$. Can we pick our knots exactly as we wish, e.g. as a Cartesian grid $X \times Y = \{(x, y) \mid x \in X \text{ and } y \in Y\}$ for some X and Y?

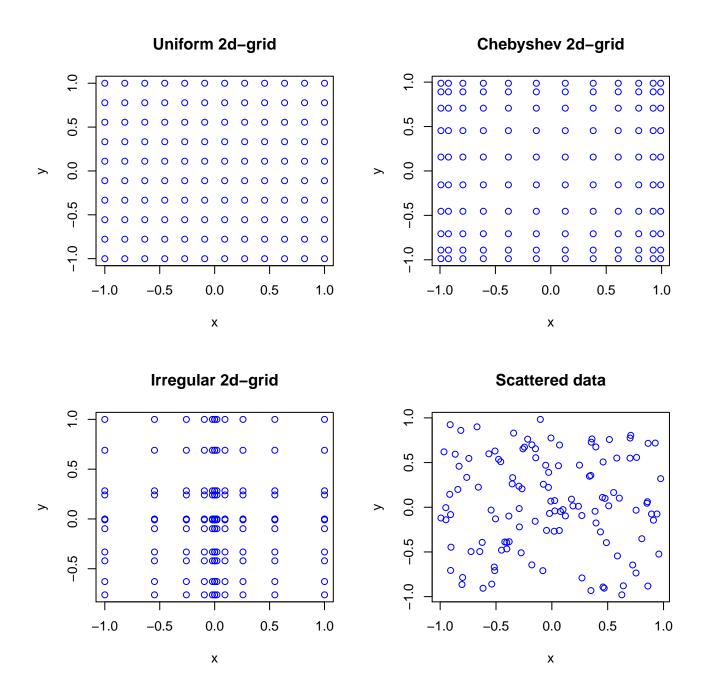
If we want our interpolant g to be a polynomial, there is a canonical way to do this. The knots should preferably be the Cartesian product of Chebyshev knots. The Chebyshev knots are not uniform, but a special set of knots which cluster near the end points.

If we are not free to choose the knots as we wish, but say we must stick to uniformly spaced knots, say $X = \{-1, -0.9, -0.8, \dots, 0.8, 0.9, 1.0\}$, it is not particularly wise to interpolate with a polynomial. The reason is that polynomial interpolants can get gradually worse as the number of knots increases.

Some functions can, for some reason or the other, only be evaluated in irregular grids, or should have a denser grid near certain areas. Some functions can't be evaluated at all, they only come as values in some scattered points. Here are typical examples in two dimensions.

```
library(chebpol)
par(mfrow=c(2,2))
grid <- list(x=seq(-1,1,length.out=12), y=seq(-1,1,length.out=10))
plot(y~x,data=expand.grid(grid),typ='p',col='blue',main='Uniform 2d-grid')
grid <- chebknots(c(x=12,y=10))
plot(y~x, data=expand.grid(grid), typ='p',col='blue',main='Chebyshev 2d-grid')
grid <- list(x=seq(-1,1,length.out=12)^3L, y=sort(c(0,runif(10,-1,1),1)))
plot(y~x, data=expand.grid(grid), typ='p',col='blue',main='Irregular 2d-grid')
data <- cbind(x=runif(120,-1,1),y=runif(120,-1,1))
plot(y~x, data=data, typ='p', col='blue', main='Scattered data')</pre>
```

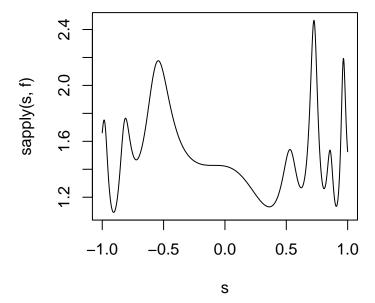
Date: July 6, 2018.



2. The Chebpol Package

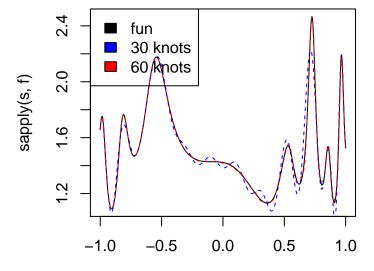
Since the generalization to multiple dimensions is fairly straightforward, we focus on the one-dimensional case. I.e. functions $[-1,1] \mapsto \mathbb{R}$. We use the following function:

```
f <- function(x) 1/mean(log1p(0.5 + sin(0.8+2.3*pi*c(0.6,1.4)*(x+0.09)^3)^2))
s <- seq(-1, 1, length.out=1000)
plot(s, sapply(s,f), typ='l')</pre>
```



Since this is a toy function, we can illustrate all sorts of knots. The first one out is the traditional Chebyshev interpolation. The oscillations are typical for polynomial interpolation with too few knots, that's how polynomials are.

```
ch30 <- ipol(f, dims=30, method='cheb')
ch60 <- ipol(f, dims=60, method='cheb')
plot(s, sapply(s,f), typ='l')
lines(s, ch30(s), col='blue', lty=2)
lines(s, ch60(s), col='red', lty=2)
legend('topleft',c('fun','30 knots','60 knots'),fill=c('black','blue','red'))</pre>
```

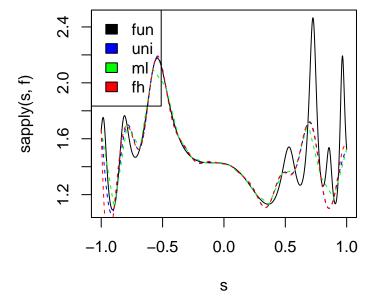


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We see that with 60 knots we follow the peaks quite well. There is another thing to note also. We can't evaluate the function f in all points at once, with f(s), we must sapply(s, f). However, the interpolants produced by package **chebpol** can evaluate in a series of points. It is even able to parallelize the evaluation, if you have more than one CPU in your computing contraption. It's just to feed it a threads= argument.

Then, what if we can't pick Chebyshev knots, but for some reason have to stick to uniformly spaced knots, say 20 or 40 of them. We then have a couple of methods to choose from. One is called "uniform" and is a trigonometric mapping of the uniformly spaced knots into Chebyshev knots. Another is piecewise linear. Then there is the Floater-Hormann rational interpolation from [1]. The F.H. interpolation comes with an integer parameter which is the degree of certain blending polynomials. In theory it works with any degree below the number of knots, but in practice high degrees introduce round off errors, so we better keep it small. We see some of the methods here:

```
plot(s, sapply(s,f), typ='l')
uni <- ipol(f, dims=20, method='uniform')
grid <- seq(-1,1,len=20)
ml <- ipol(f, grid=grid, method='multilinear')
fh <- ipol(f, grid=grid, method='fh', k=3)
lines(s, uni(s), col='blue', lty=2)
lines(s, ml(s), col='green', lty=2)
lines(s, fh(s, threads=4), col='red', lty=2)
legend('topleft',c('fun','uni','ml','fh'),fill=c('black','blue','green','red'))</pre>
```



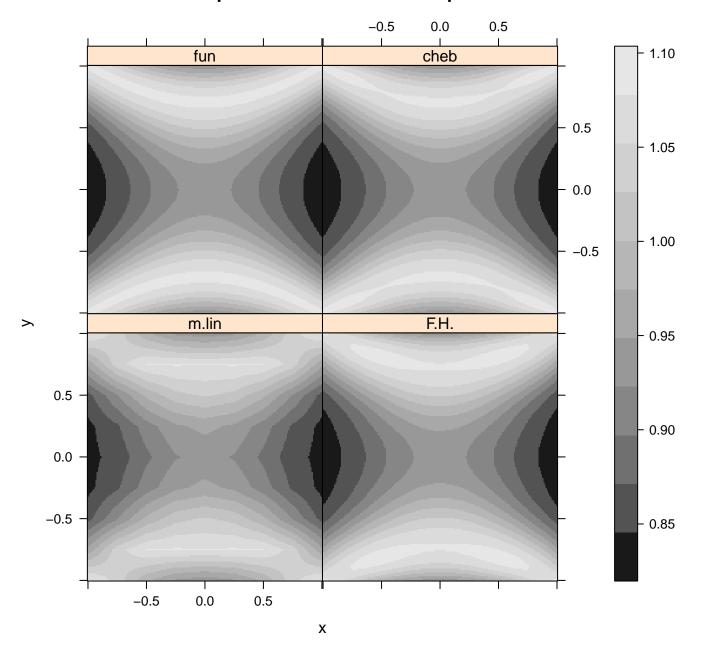
3. Multidimensional data

All of the above methods work with more than one dimension via a standard tensor product construction, but that presupposes that the function can be evaluated on a Cartesian grid. The Chebyshev and uniform methods use particular grids. The method "general" is similar to "uniform" in that it transforms an arbitrary Cartesian grid into a Chebyshev grid by means of a monotonic spline in each dimension. The multilinear and Floater-Hormann methods also work on arbitrary Cartesian grids.

To illustrate how this works, here is a quite nice function on $[-1,1]^2$ and level plots of it and the 9×9 Chebyshev interpolation, and the multilinear and Floater-Hormann methods on a uniform 9×9 grid.

```
f1 <- function(x) 1.5/log(5+sin(pi/2*(x[1]^2-2*x[2]^2)))
ch1 <- ipol(f1, dims=c(9,9), method='cheb')
igrid <- list(x=seq(-1,1,len=9), y=seq(-1,1,len=9))
ml1 <- ipol(f1, grid=igrid, method='multilinear')
fh1 <- ipol(f1, grid=igrid, method='fh', k=3)</pre>
```

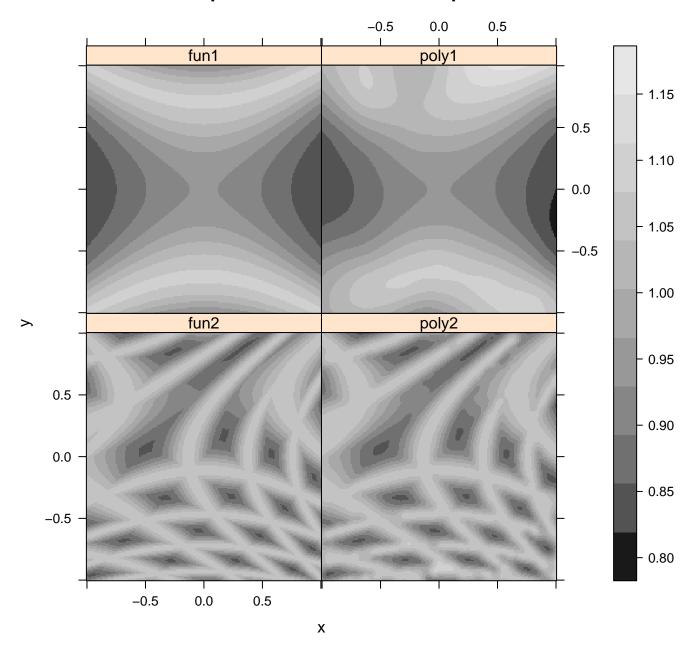
Level plots of function and interpolations



It is not so easy to see in a picture, but the multilinear interpolant is not differentiable in the grid points, it is piecewise linear. The other interpolants are typically twice continuously differentiable.

There is one interpolation method we have not mentioned yet. If the data are scattered in two or more dimensions, none of the methods above will work. They need a Cartesian product grid. The **chebpol** package contains one method for scattered data also, the polyharmonic spline (see [2]). It is parametric with an integer $k \in \mathbb{N}$. For free we also throw in a Gaussian radial basis function interpolation for negative $k \in \mathbb{R}^-$. Here we create two functions and plot level plots of them and their polyharmonic interpolations. One of the functions is quite smooth, the other is quite wobbly and requires a lot of scattered points.

Level plots of functions and interpolations



The polyharmonic spline method is quite slow. It requires solving a $K \times K$ dense linear system, where K is the number of knots. It helps a lot to use a tuned and parallel BLAS with R, such as Intel's MKL.

4. Speed considerations

There are two parts to interpolation. Creating the interpolant, and using it. Both of these take some time. The time depends on the method, as well as the dimension r of the problem, and the number of grid points n along a dimension, or the number of knots K. For the Cartesian grids, the number of knots is n^r .

Traditionally, Chebyshev interpolation has been favoured for two reasons. The Chebyshev interpolation is the right way to do polynomial interpolation, due to certain analytical facts. The Chebyshev coefficients can also be calculated by a (possibly multi dimensional) Fast Fourier Transform (FFT). This is much faster than solving a large linear system which is the pedestrian approach for finding polynomial interpolants. **chebpol** uses the FFT from the software package FFTW, if it was available during compilation. Otherwise a slower matrix method is used. Creation with FFT typically takes of the order $O(n^r \log n)$. The evaluation of Chebyshev interpolants is not particularly fast, **chebpol** uses the Clenshaw-Curtis algorithm, which is a variant of Horner's method for evaluating polynomials. It typically goes in time n for each dimension, but they must be combined, so $O(n^r)$.

The multilinear interpolation is the fastest in **chebpol**. Creating the interpolant takes little time at all, the grid and function values are saved, as is, that's all. Evaluation consists of computing a convex combination of the function values in the corners of the surrounding hypercube. Typically a $O(\log n)$ task for each dimension, but there are 2^r corners for an r-dimensional problem, so $O(r2^r + r \log n)$.

Creation of the Floater-Hormann interpolation requires calculating the weights, roughly an $O(rn^2)$ task. Evaluation is $O(n^r)$. Quite similar to the Chebyshev method.

The polyharmonic spline method is the most demanding. Creation requires calculating some weights which amounts to solving a dense $K \times K$ linear system, a $O(K^3)$ task (or $O(n^{3r})$ to compare with the grid methods). Evaluation is faster, of the order O(Kr) (or $O(rn^r)$).

To alleviate speed problems, parts of **chebpol** has been parallelized with OpenMP. The number of threads is taken from getOption("chebpol.threads") which is initialized from the environment variable CHEBPOL_THREADS. It defaults to 1. Some of the creation routines use parallelization, however the polyharmonic spline does not. It depends on the underlying BLAS for solving the system, of course it helps if it is optimized and parallel.

Evaluation of the interpolants are parallelized if one evaluates more than one point. I.e. the interpolants can take a matrix of column vectors, and may calculate values in parallel:

```
m <- matrix(runif(2*6,-1,1),2)
print(m)
## [,1] [,2] [,3] [,4] [,5] [,6]
## [1,] 0.4906175  0.2094953  0.3710621  0.2709684  -0.9915001  -0.08214581
## [2,] 0.7909871  -0.1995721  -0.2714803  0.4300307  0.4532603  0.94648520
print(ph2(m, threads=3))
## [1] 1.0040382  0.9846754  1.0055827  0.9107847  0.9306019  0.8980599</pre>
```

This means that even if one has a function available in R, it might be faster to use an interpolant since it's automatically parallelized. Even if one does not have more than one CPU, it is worthwile to use the matrix interface if more than one point is to be evaluated (such as in the vector interface of package **cubature**), it is a lot faster with ph2(m) than with apply(m,2,ph2).

```
f <- ipol(sin, grid=seq(0,1,length.out=1000), method='multilinear')
a <- runif(1e6)
system.time(sin(a))
##
      user
            system elapsed
##
     0.067
             0.000
                     0.068
system.time(f(a,threads=4))
##
            system elapsed
      user
##
     0.155
            0.001 0.045
```

REFERENCES

- Michael S. Floater and Kai Hormann, Barycentric rational interpolation with no poles and high rates of approximation, Numerische Mathematik 107 (2007), no. 2, 315–331.
- 2. Thomas Hangelbroek and Jeremy Levesley, On the density of polyharmonic splines, Journal of Approximation Theory 167 (2013), 94 108.

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