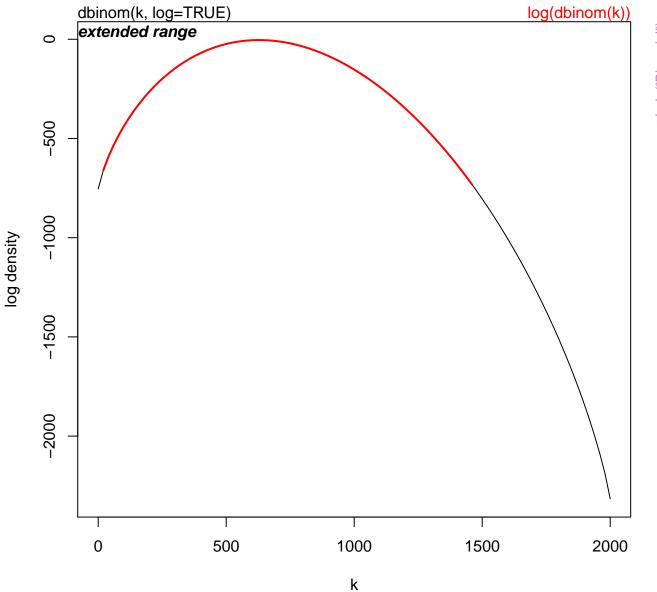
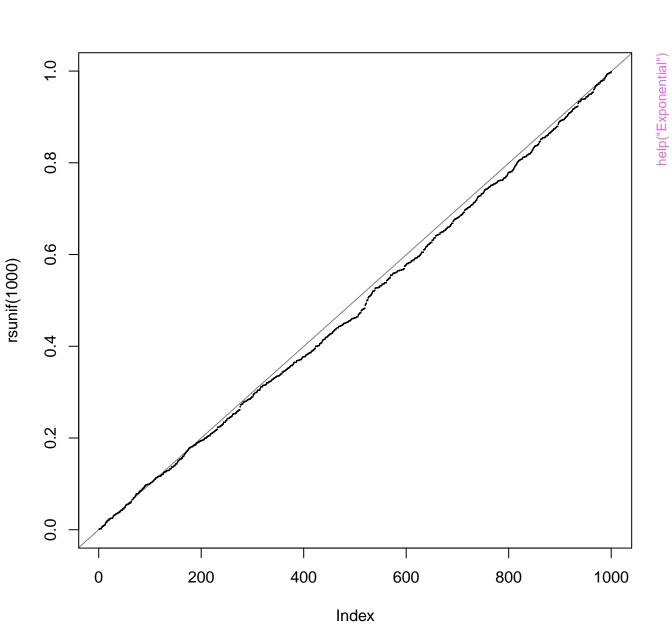
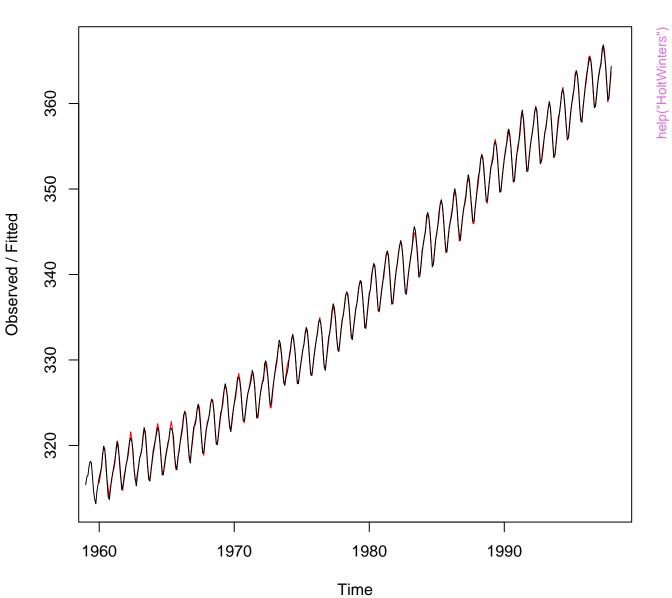
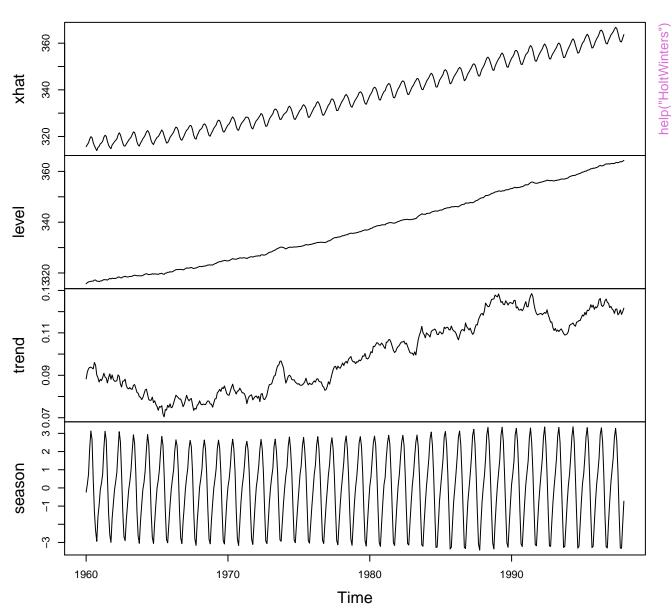
dbinom(*, log=TRUE) is better than log(dbinom(*))

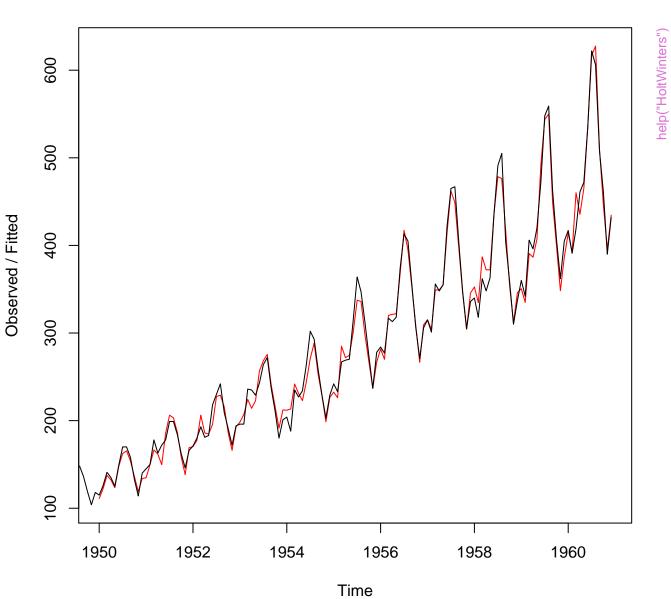




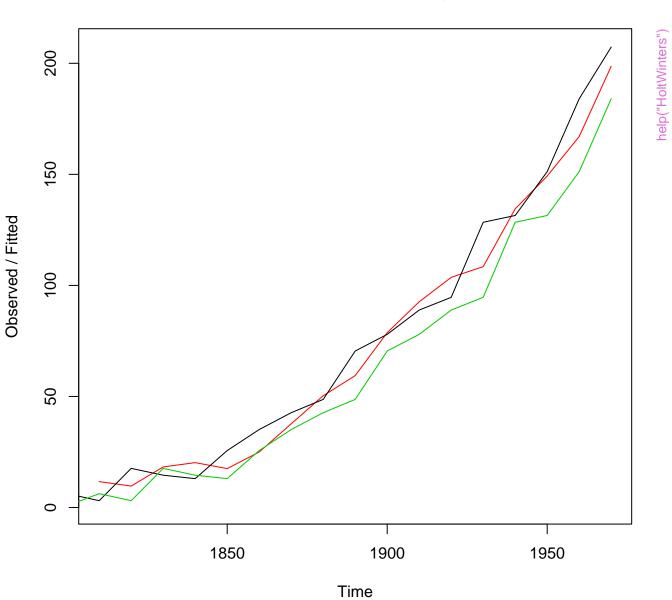




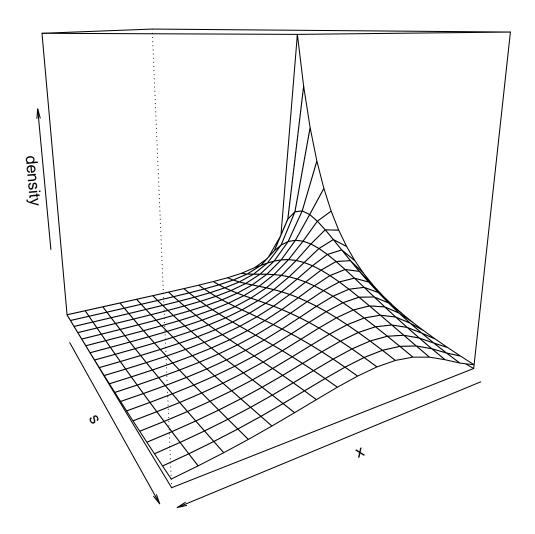




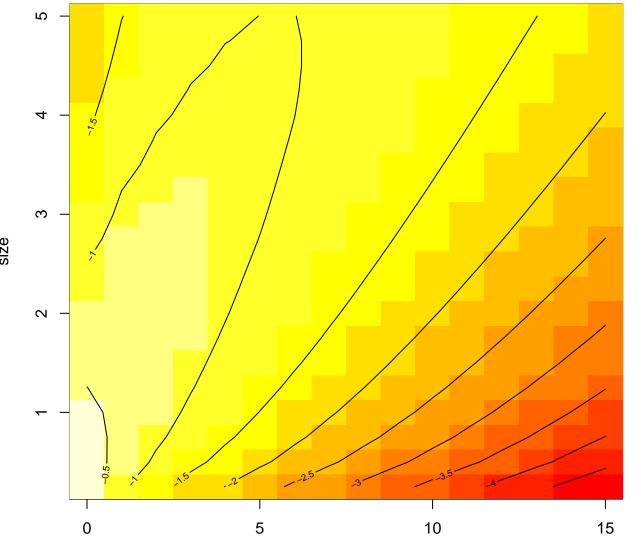
Holt-Winters filtering

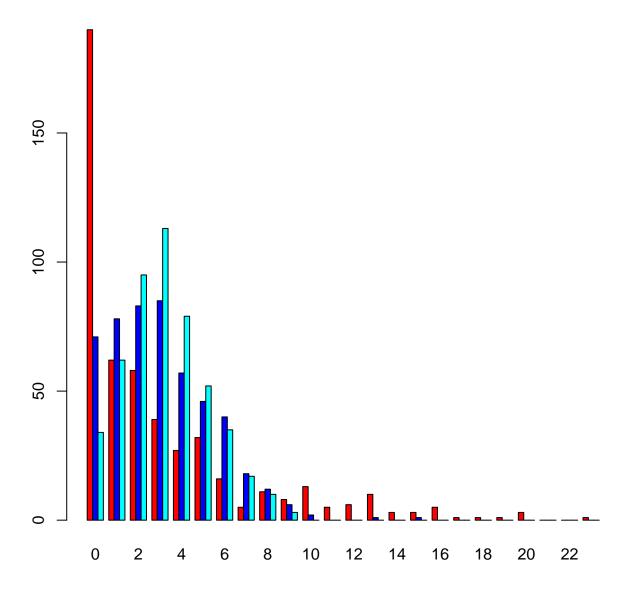


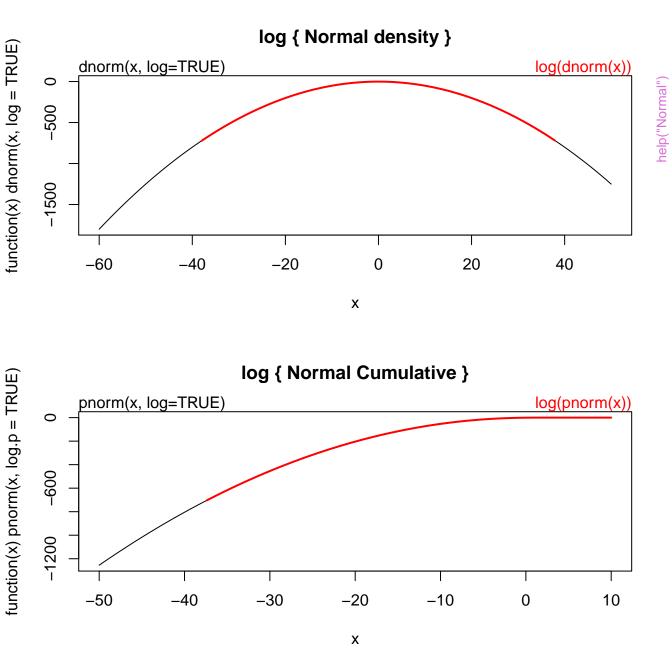
negative binomial density(x,s, pr = 0.4) vs. x & s

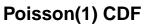


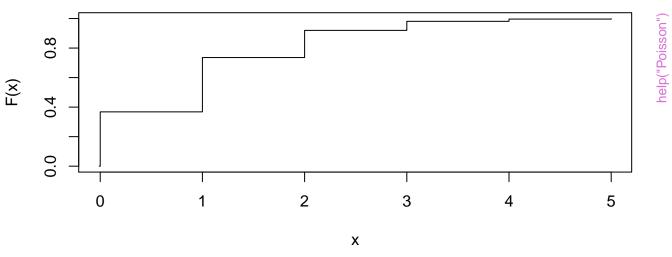
log [negative binomial density(x,s, pr = 0.4) vs. x & s]

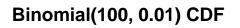


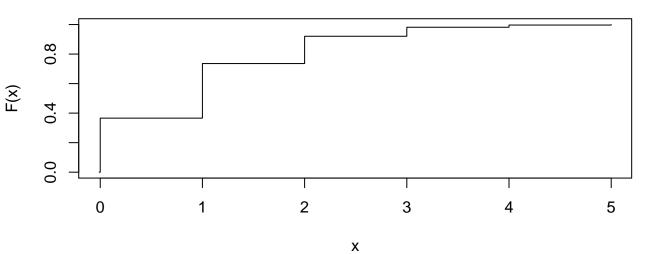




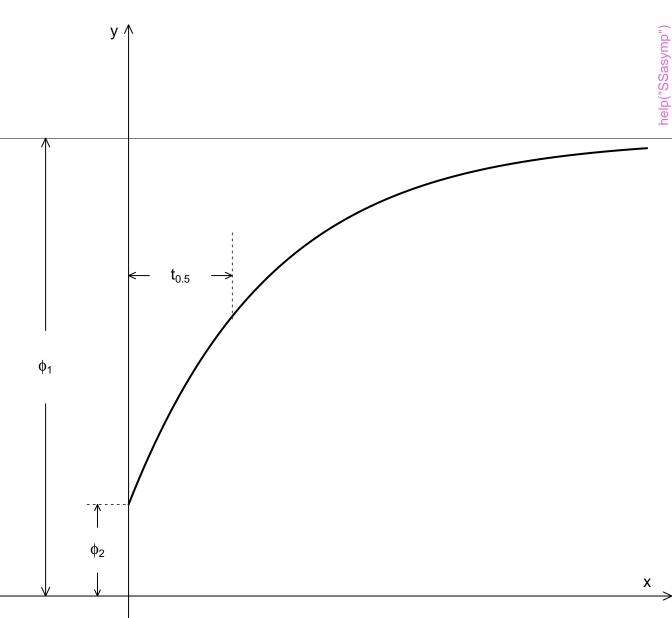




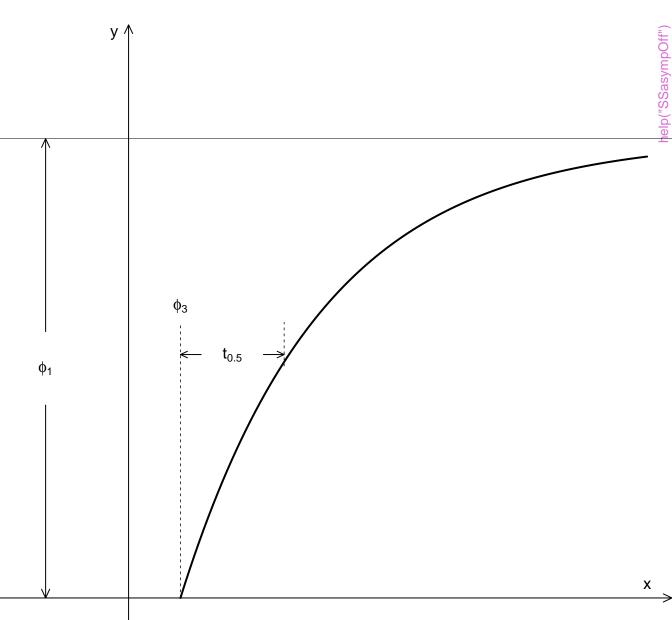




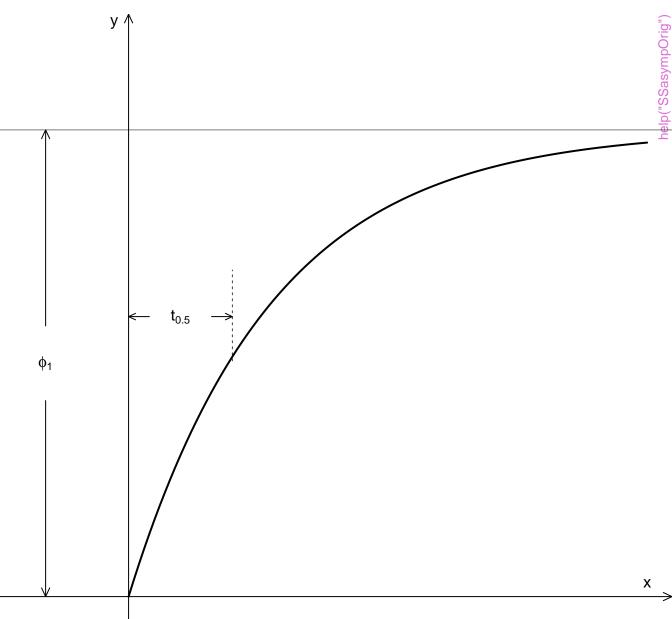
Parameters in the SSasymp model



Parameters in the SSasympOff model



Parameters in the SSasympOrig model

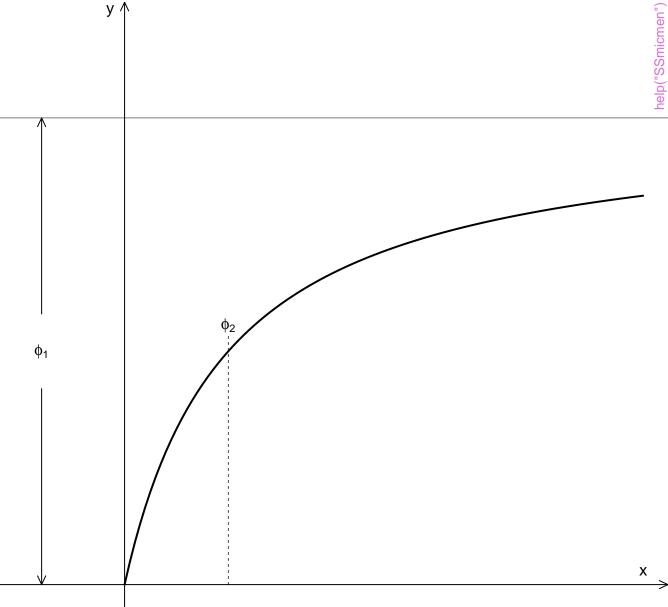


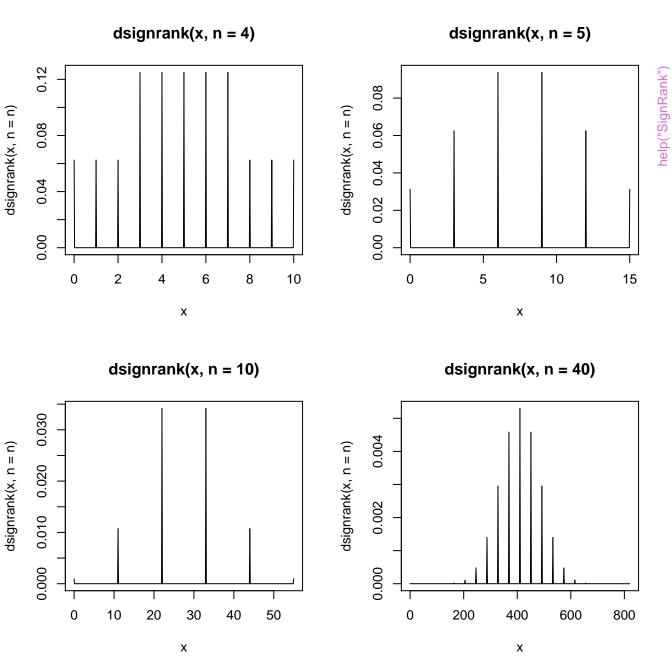
Components of the SSbiexp model 5 -

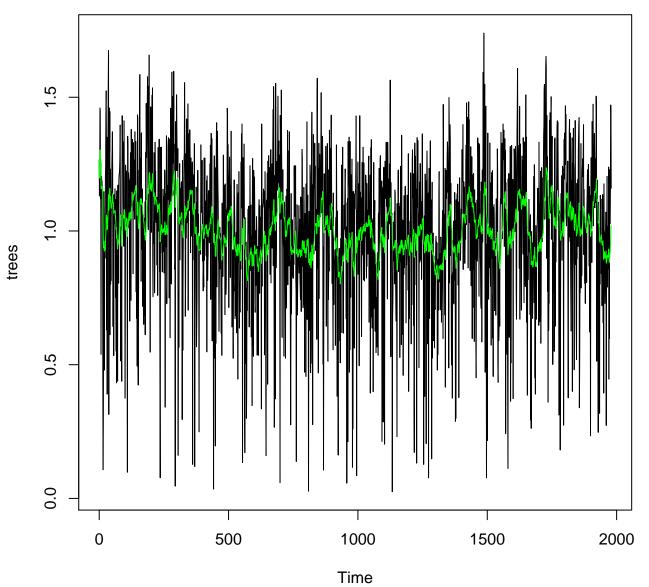
Parameters in the SSfpl model у↑ φ_1 ϕ_2

Parameters in the SSlogis model у↑ φ_1

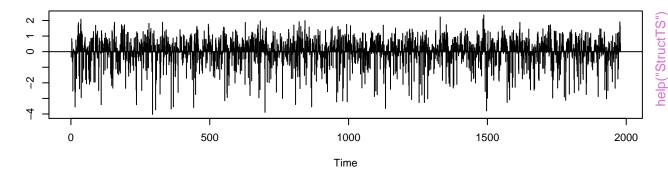
Parameters in the SSmicmen model



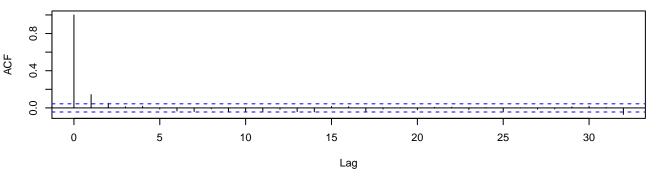




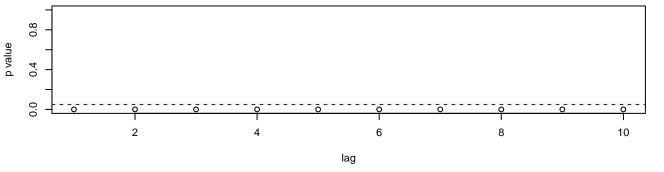
Standardized Residuals

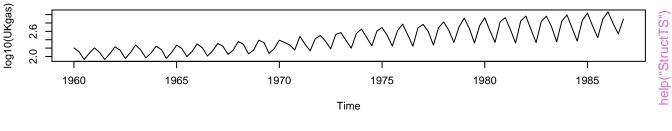


ACF of Residuals

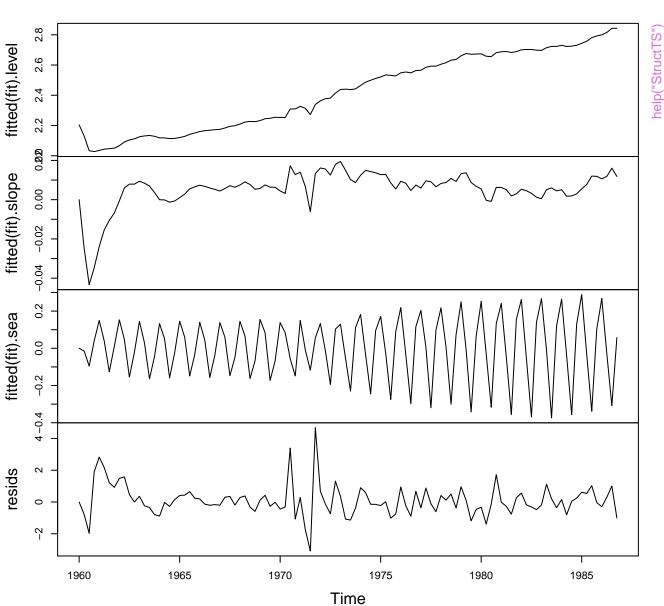


p values for Ljung-Box statistic

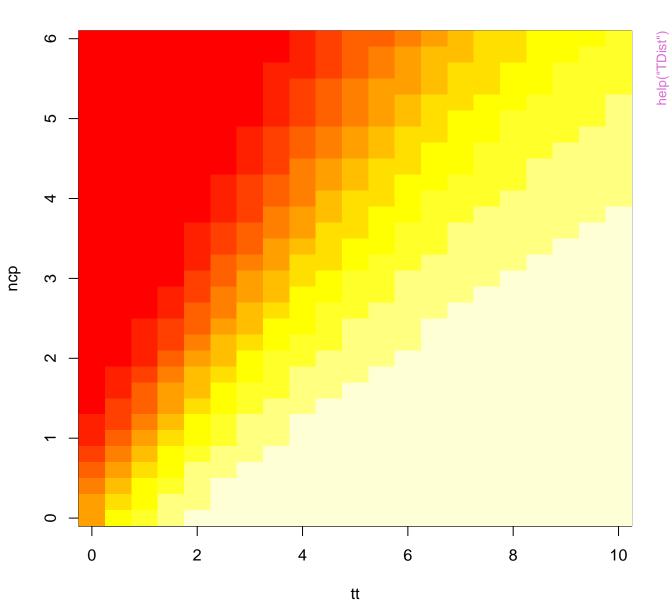




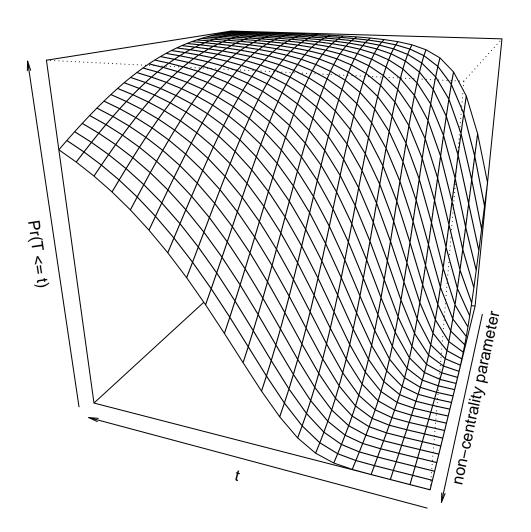
UK gas consumption



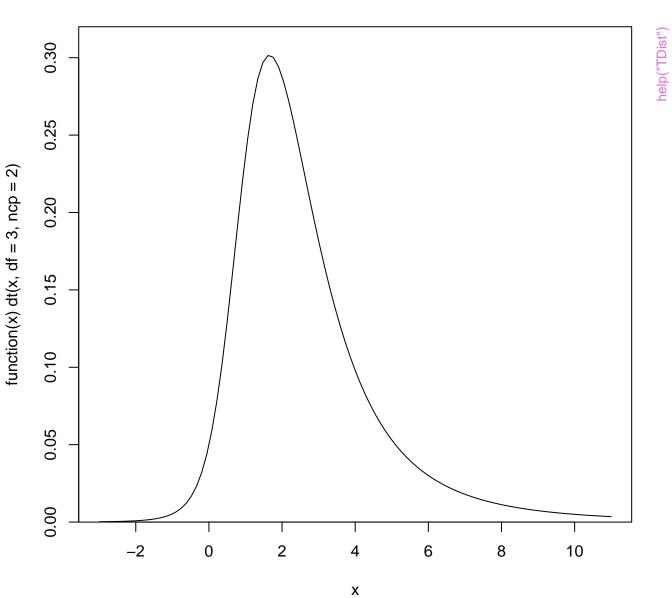
Non-central t - Probabilities

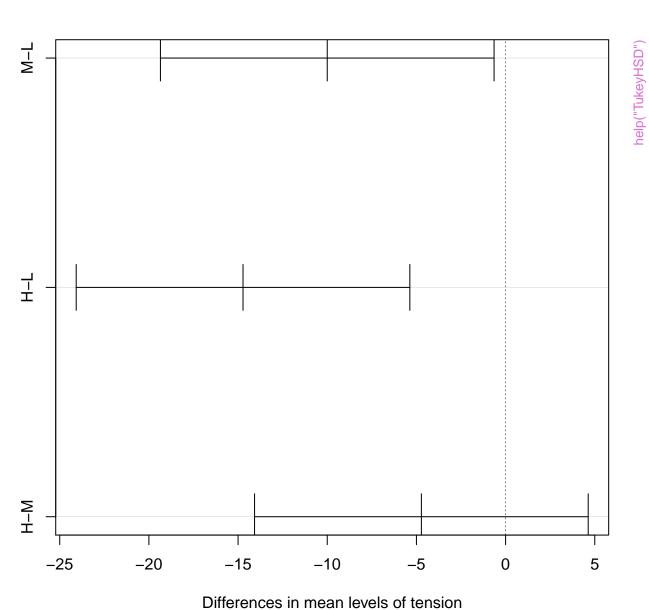


Non-central t - Probabilities

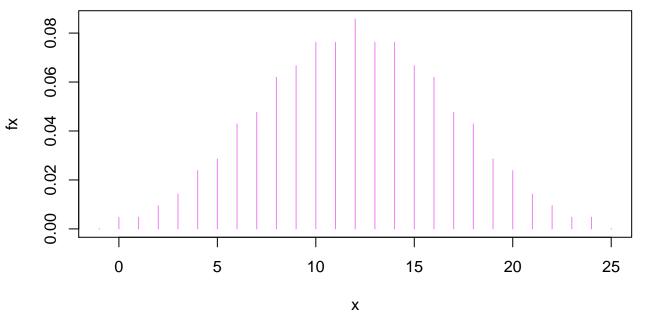


Non-central t - Density



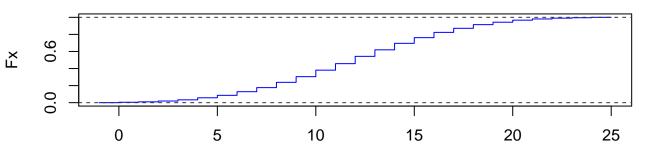




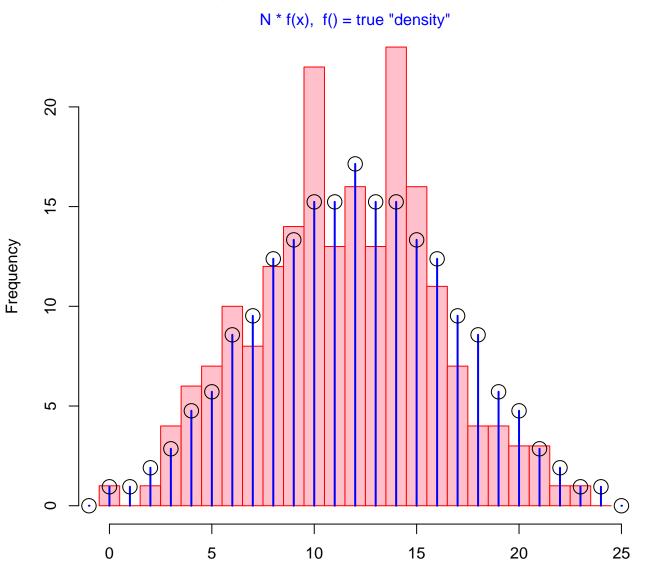


Distribution of Wilcoxon-Statist.(n=6, m=4)

Χ

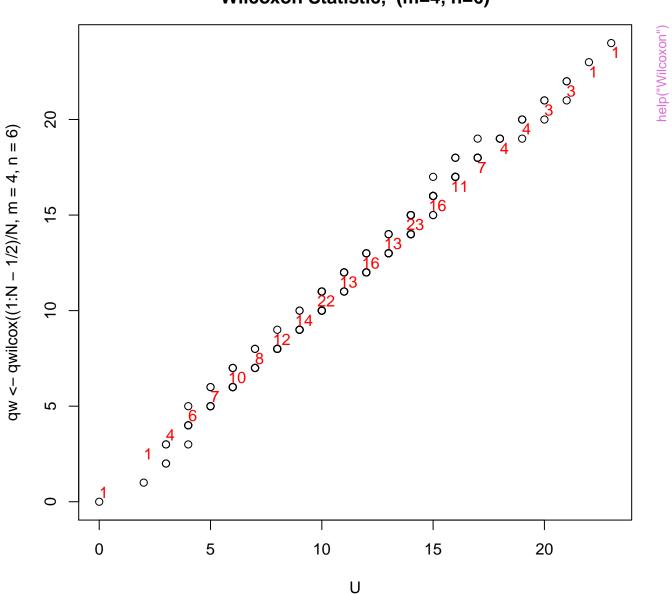


Histogram of $U \leftarrow rwilcox(N, m = 4, n = 6)$

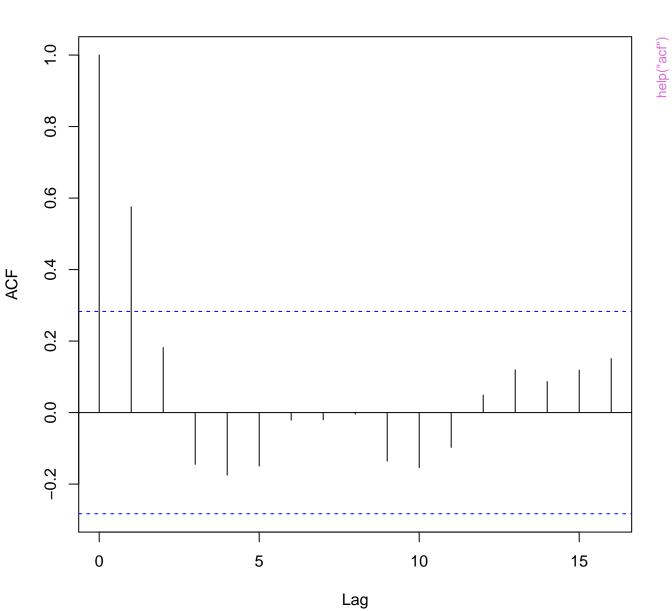


U <- rwilcox(N, m = 4, n = 6)N = 200

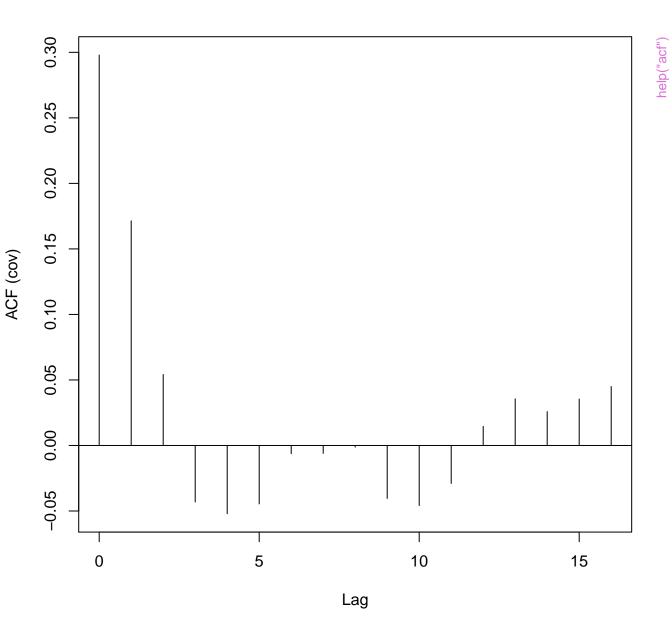
Q-Q-Plot of empirical and theoretical quantiles Wilcoxon Statistic, (m=4, n=6)



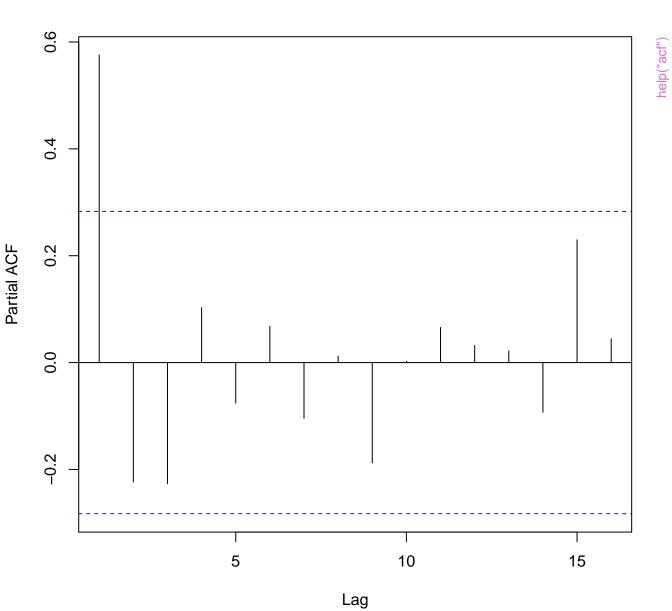
Series Ih



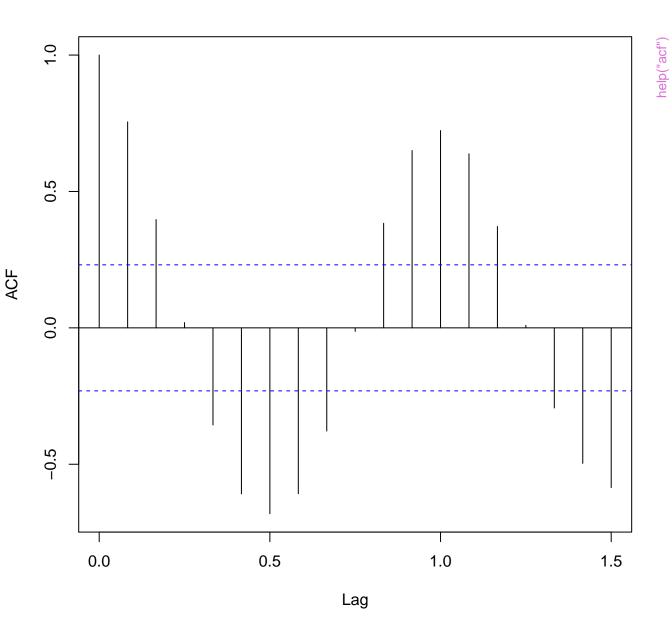
Series Ih



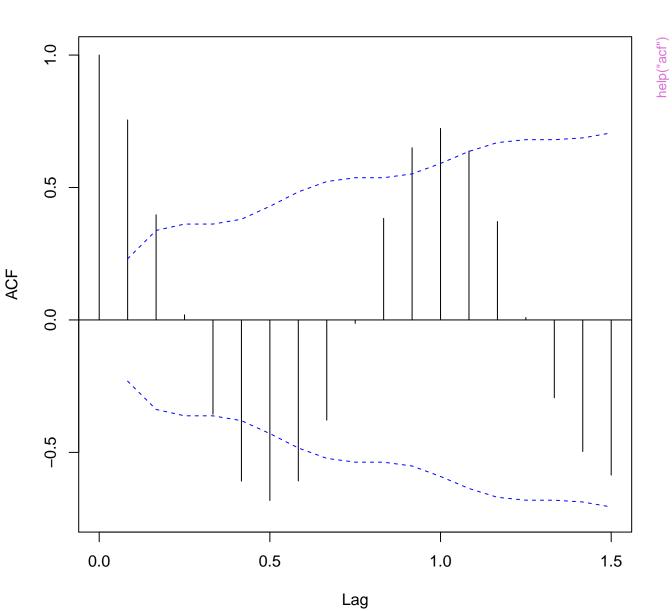
Series Ih

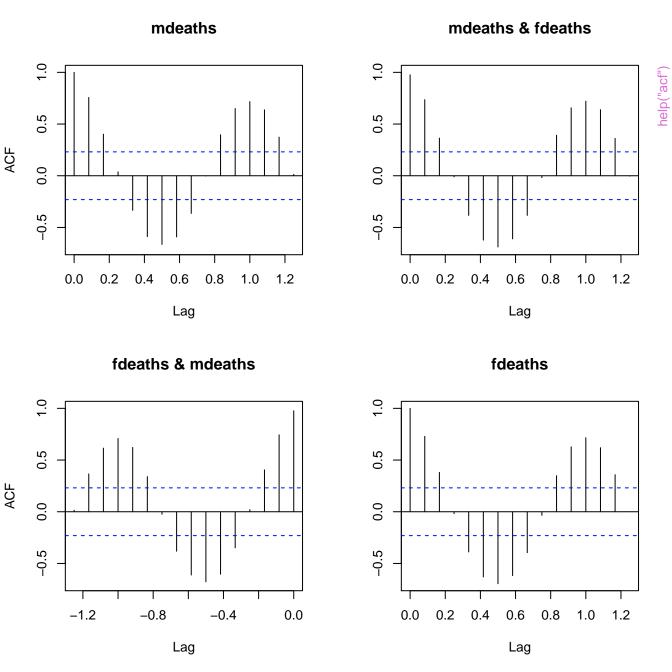


Series Ideaths

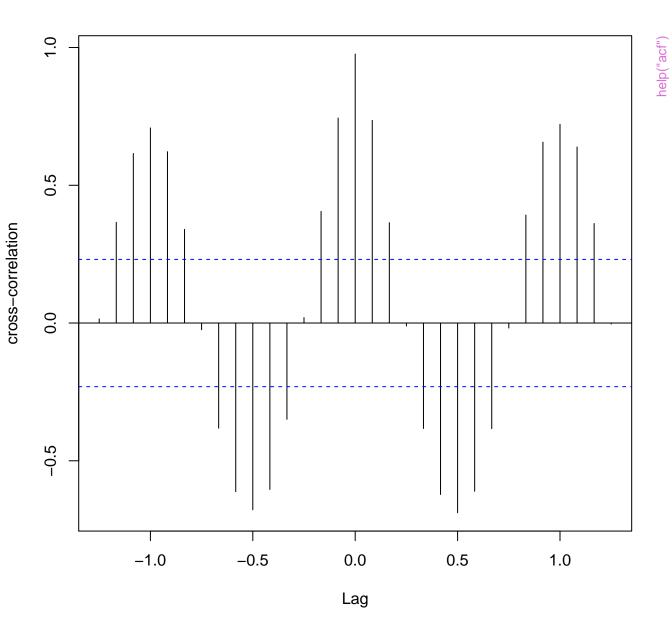


Series Ideaths

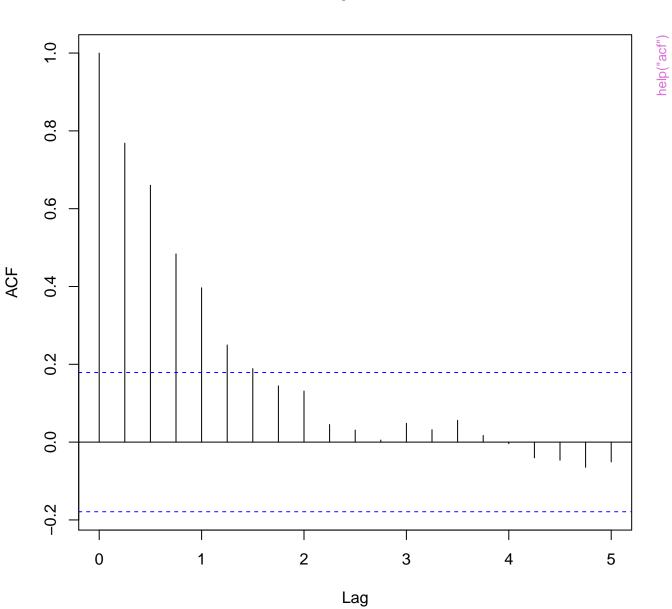




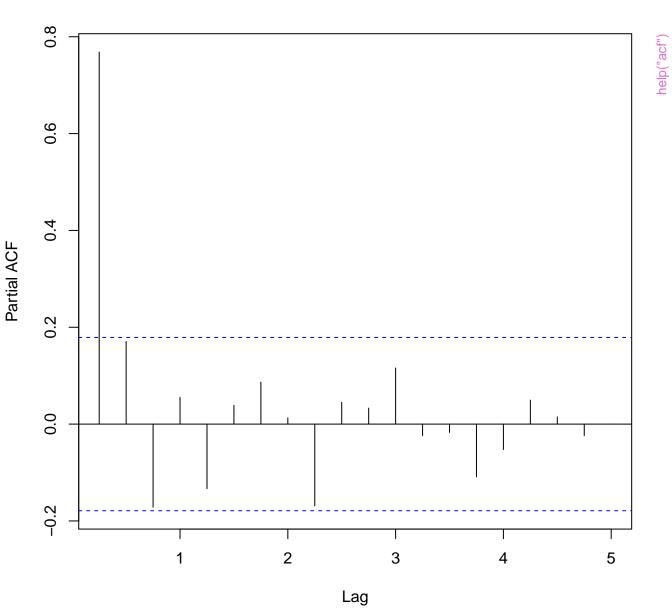
mdeaths & fdeaths

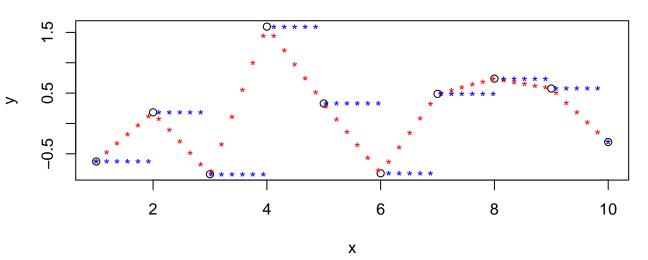


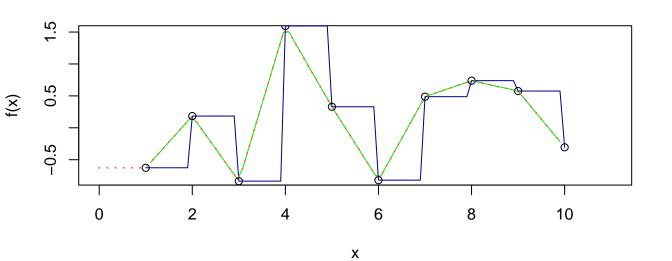
Series presidents



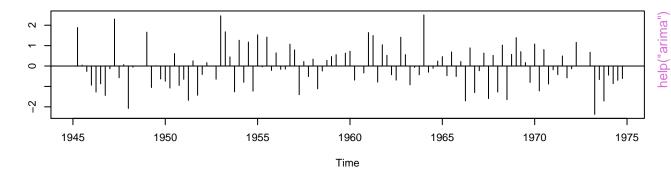
Series presidents



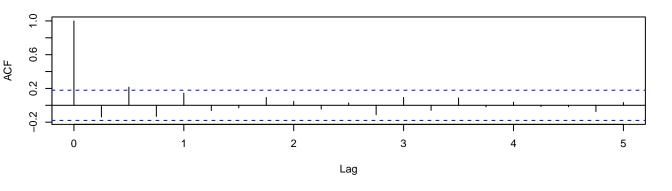




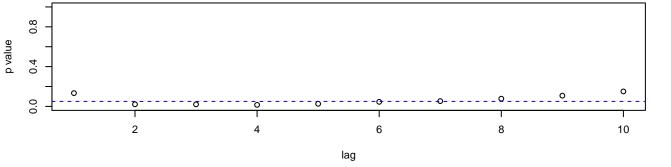
Standardized Residuals



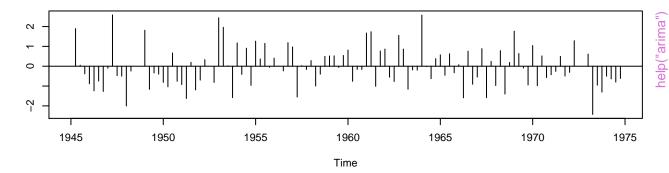
ACF of Residuals



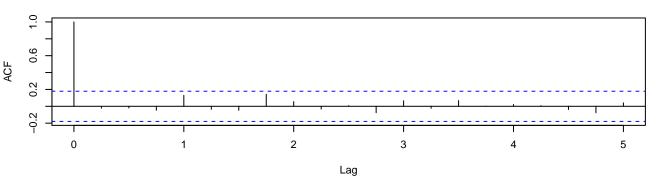
p values for Ljung-Box statistic



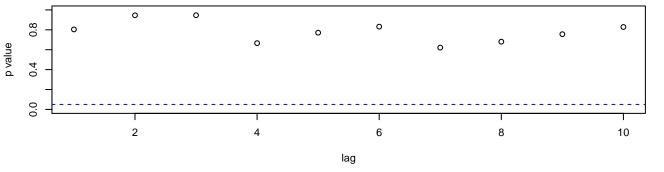
Standardized Residuals

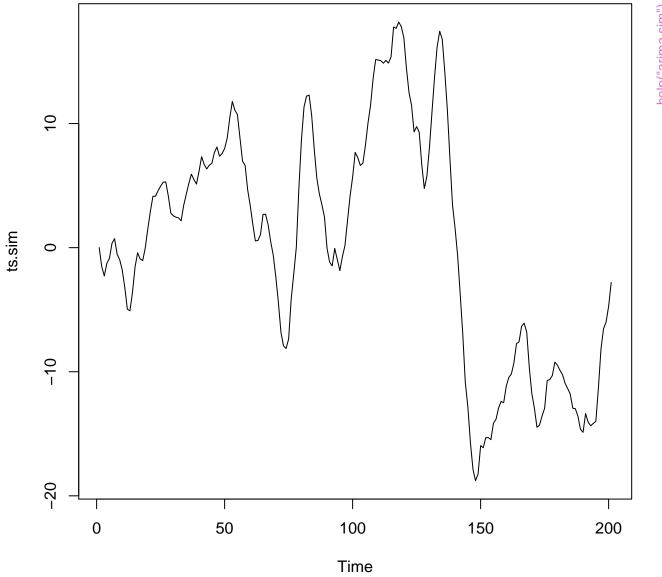


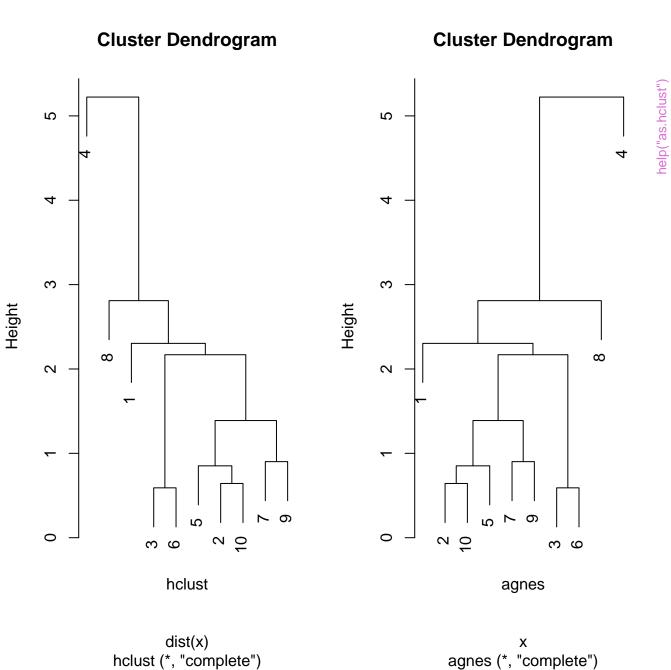
ACF of Residuals

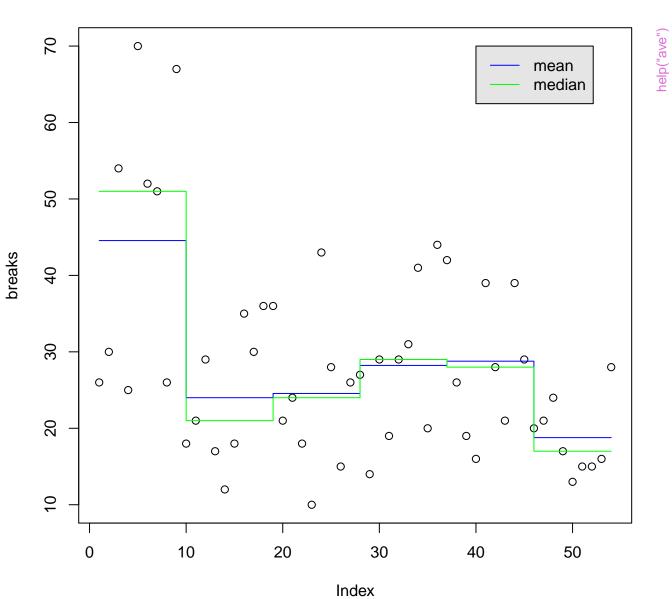


p values for Ljung-Box statistic

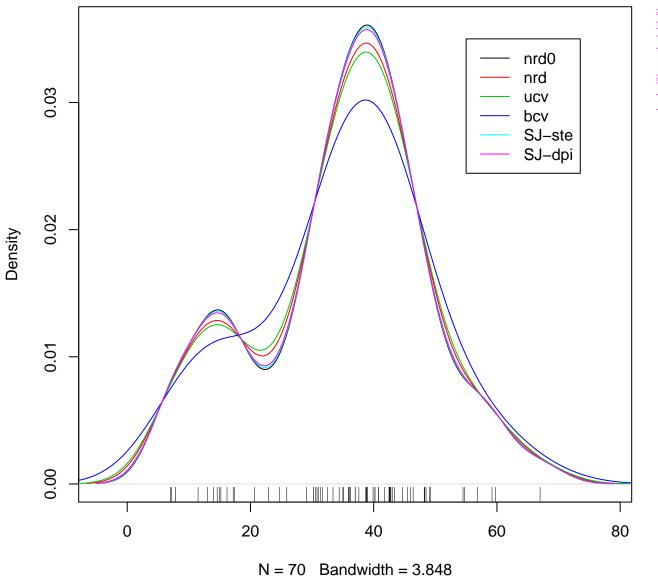


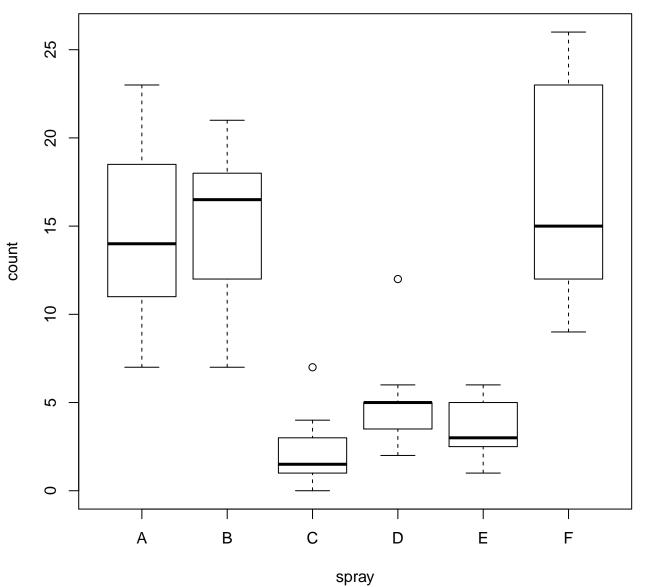


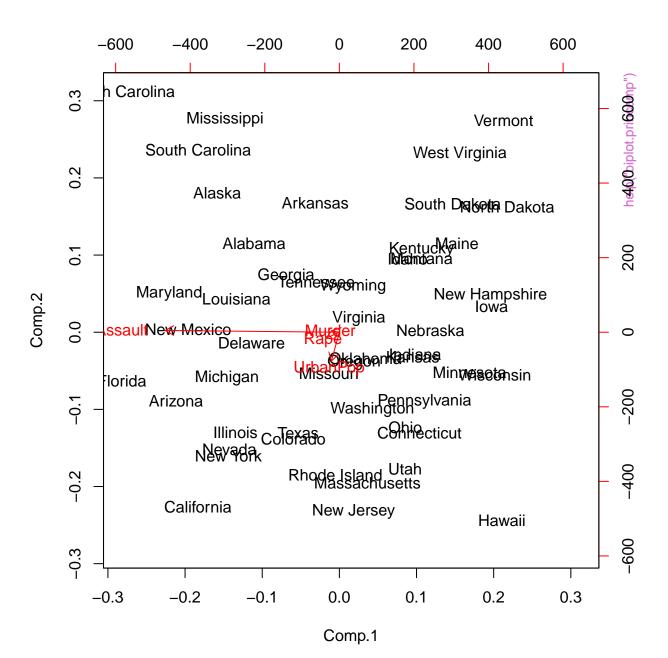




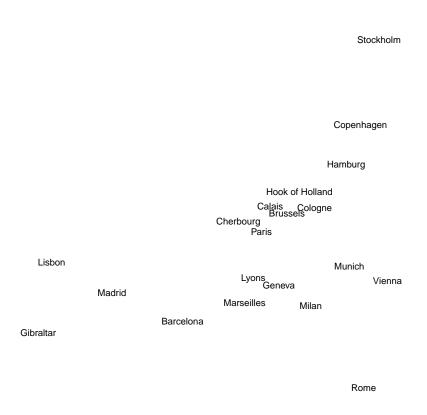
density.default(x = precip, n = 1000)



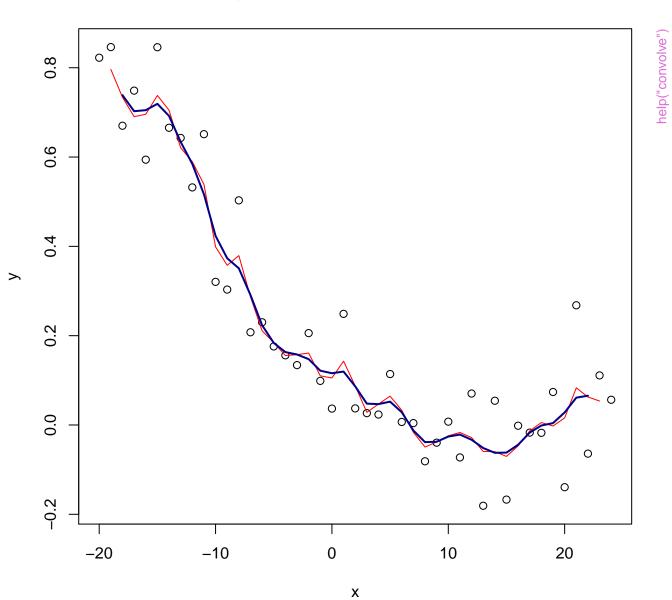




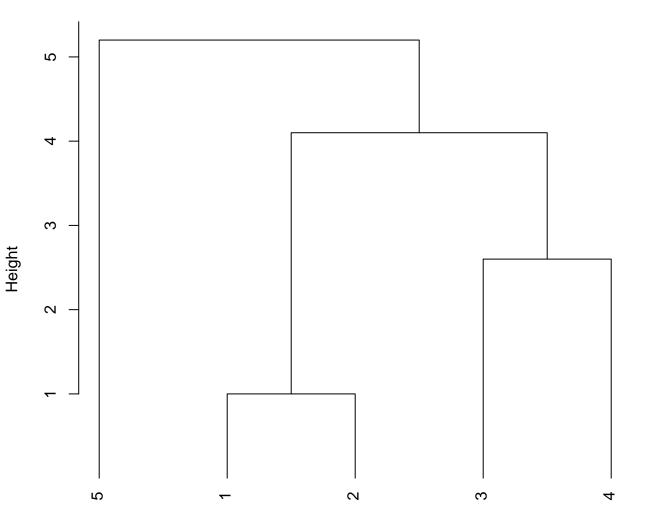
cmdscale(eurodist)



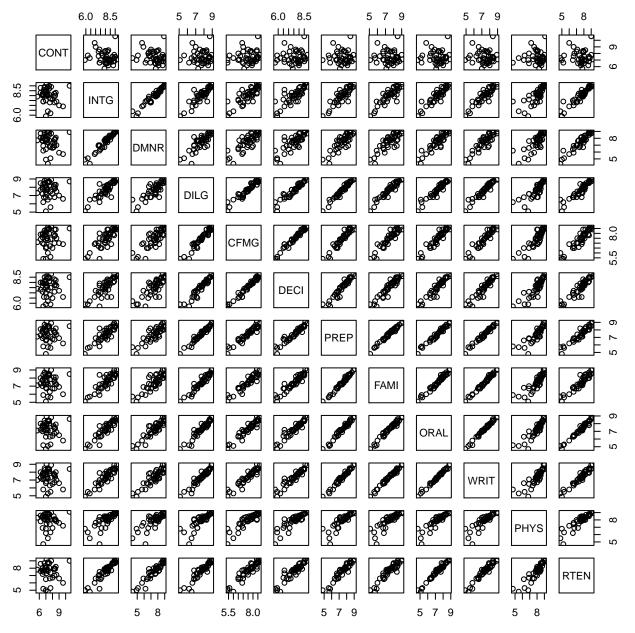
Using convolve(.) for Hanning filters

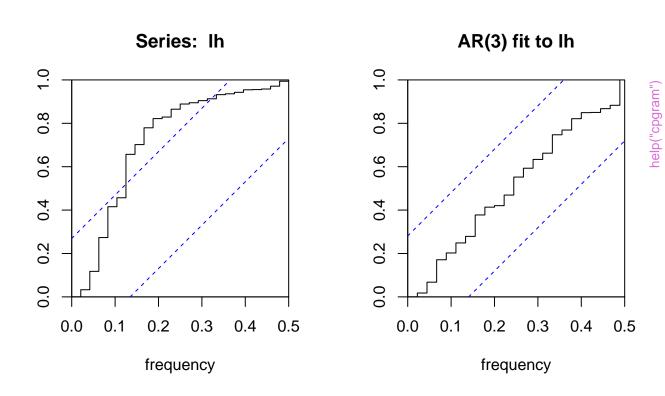


Cluster Dendrogram

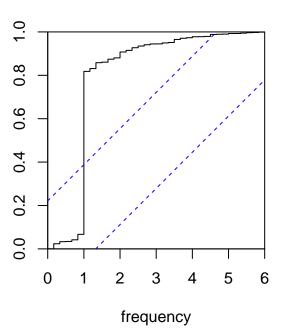


d0 hclust (*, "average")

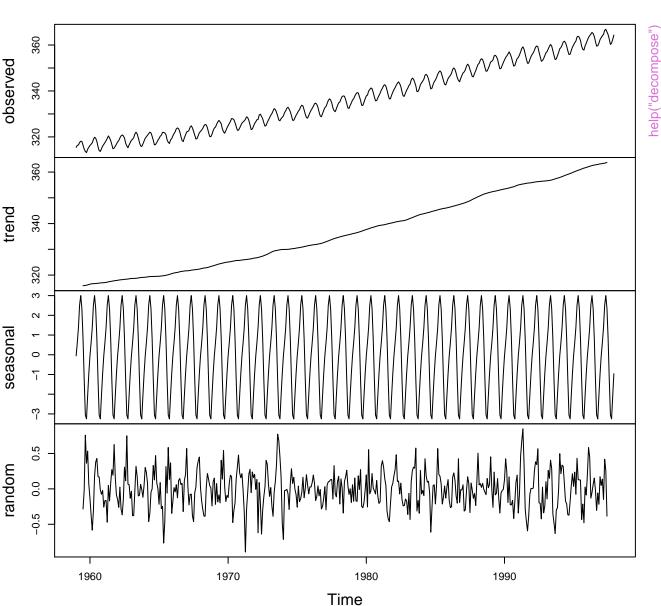


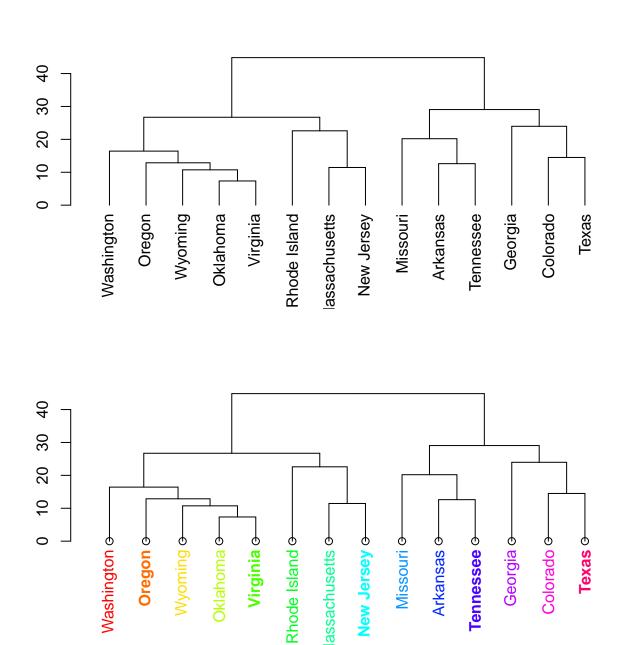


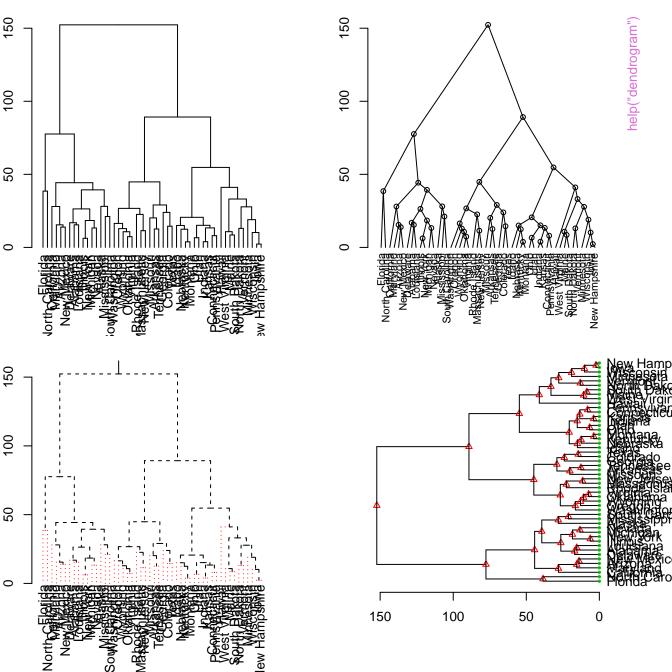
Series: Ideaths

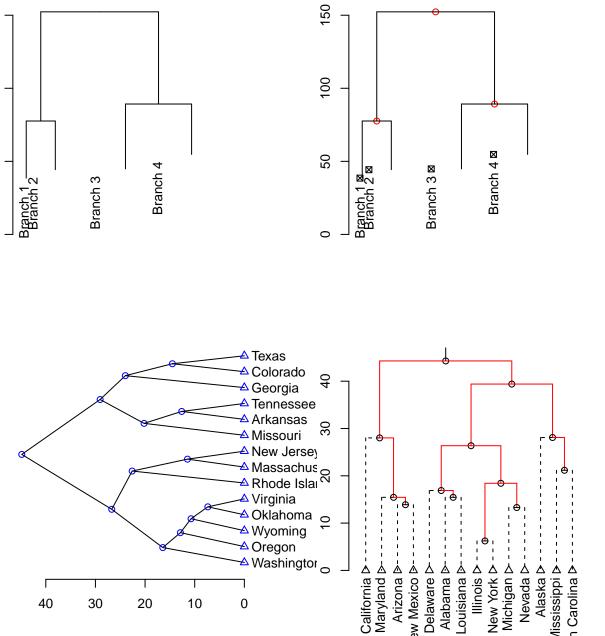


Decomposition of additive time series









Arizona 🕨

New Mexico ▶

_ouisiana ⊳ Illinois ⊳ New York ⊳ Michigan ⊳

Alabama ⊳ Delaware D

Alaska ⊳ Mississippi ⊳ outh Carolina ▶

150

100

20

0

40

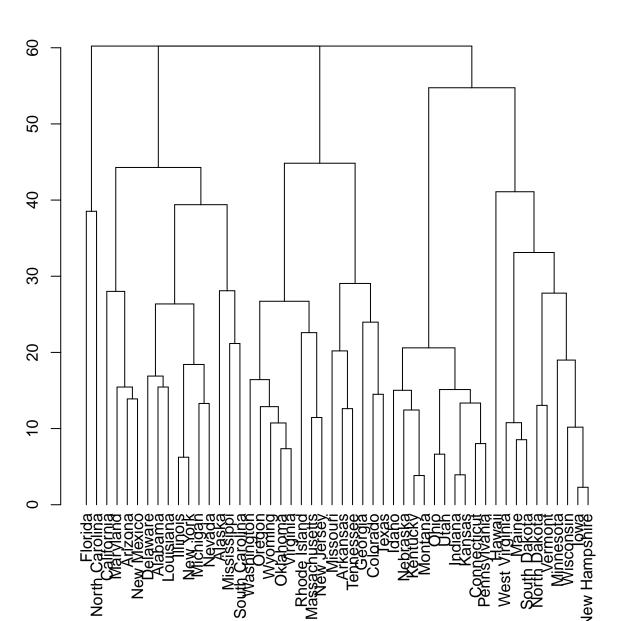
30

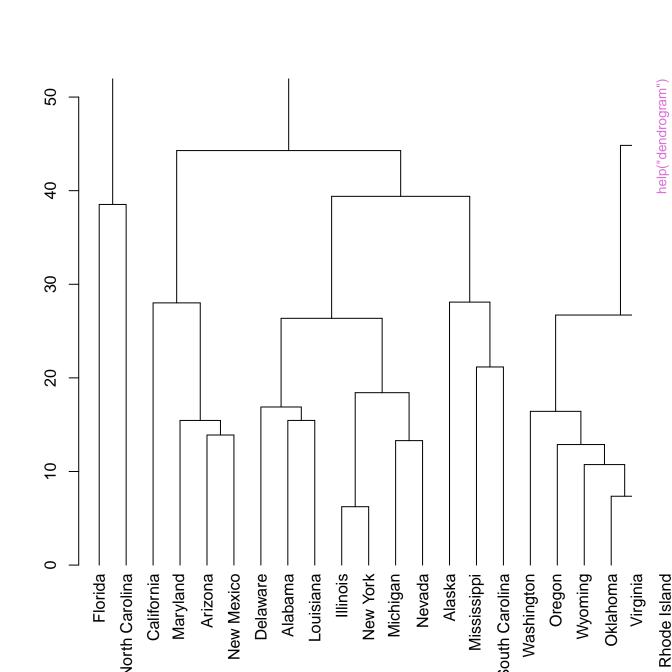
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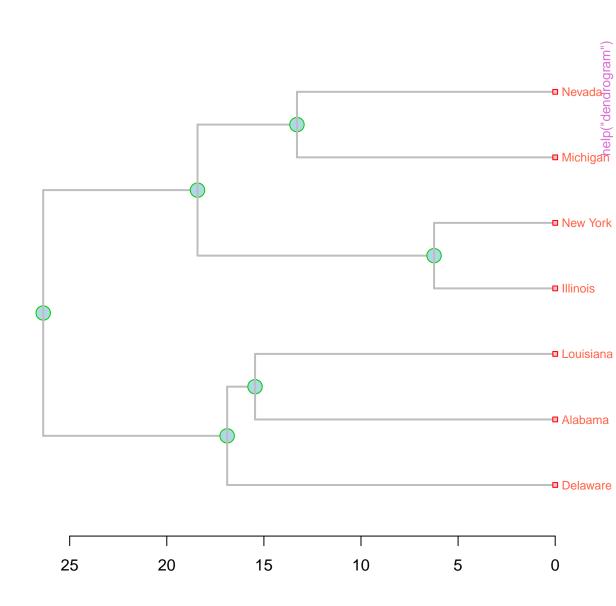
10

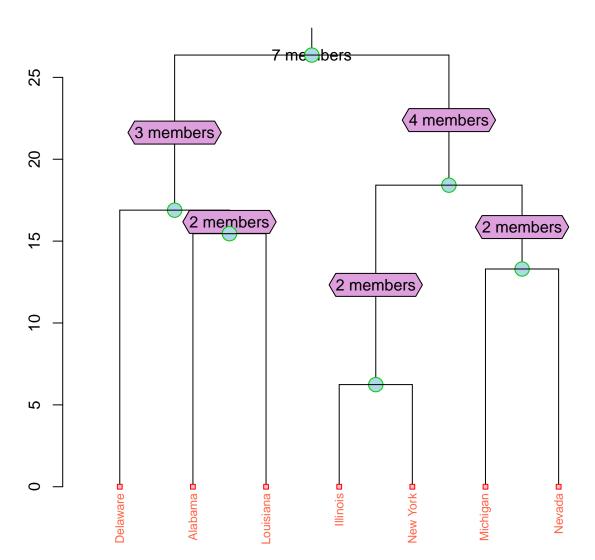
0

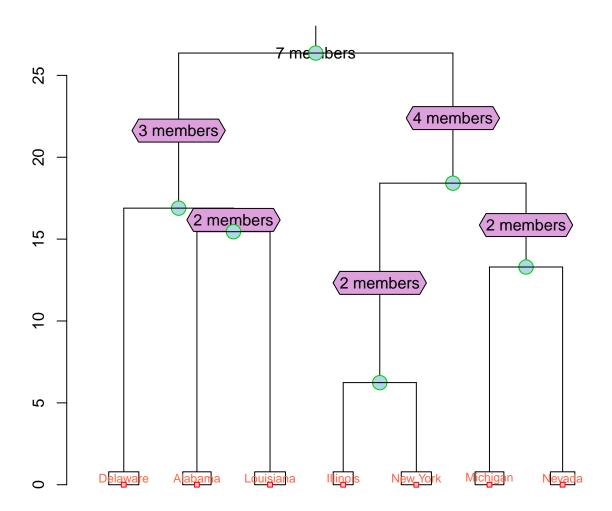
merge(d1, d2, d3, d4) |-> dendrogram with a 4-split

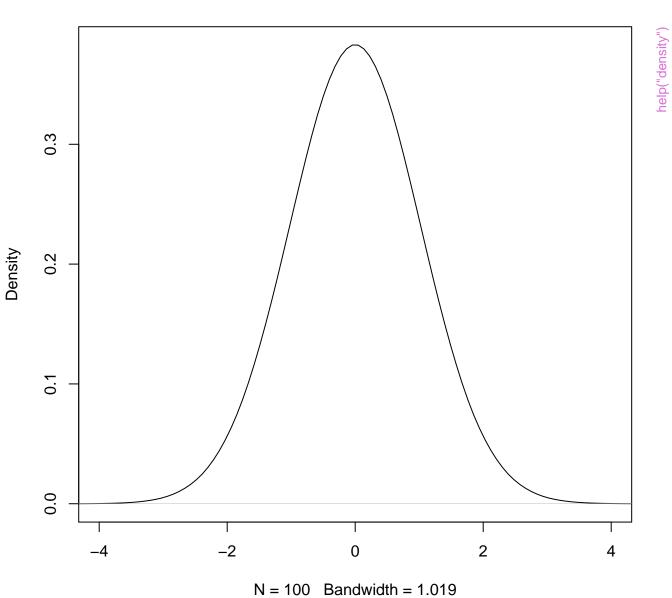


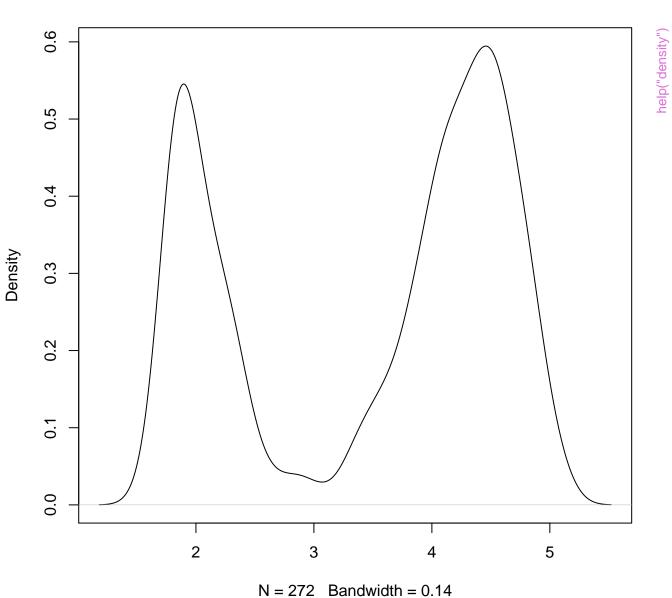




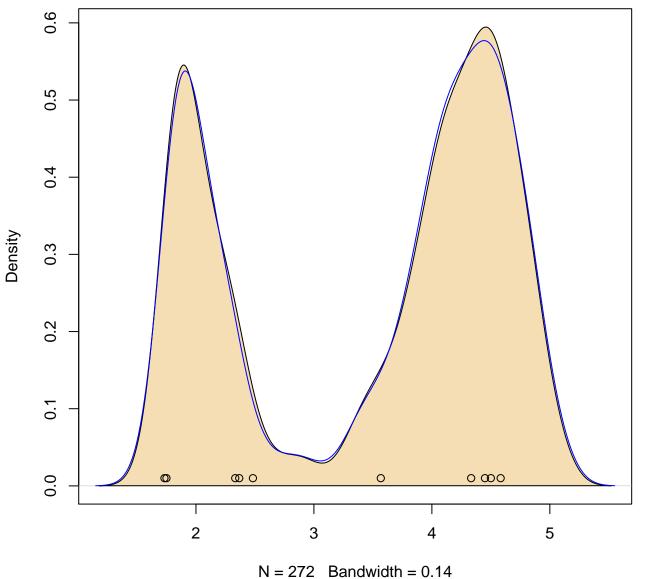




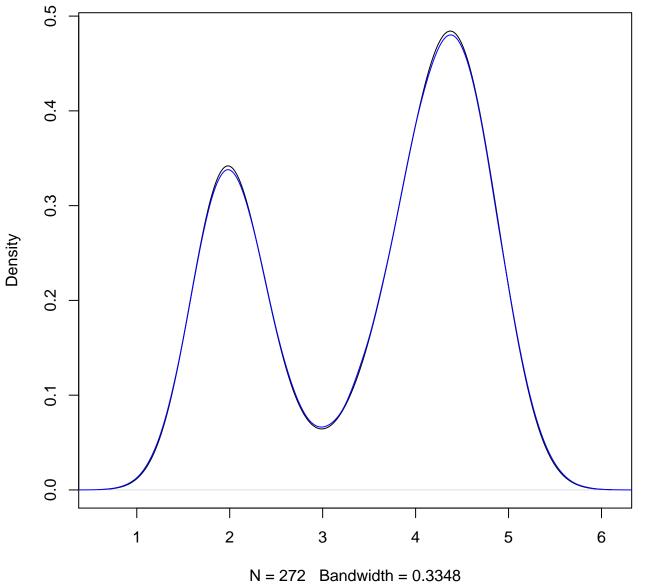




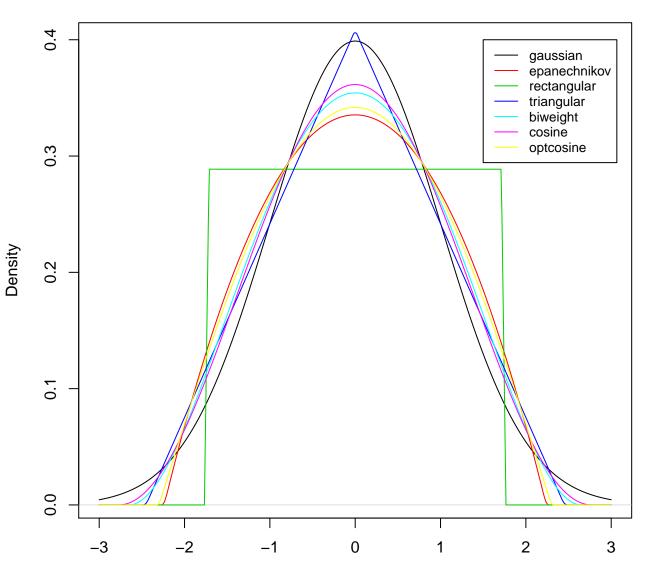




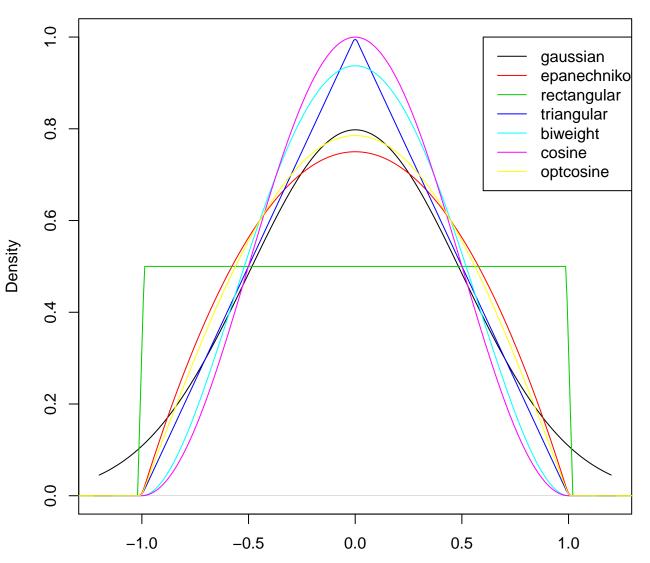




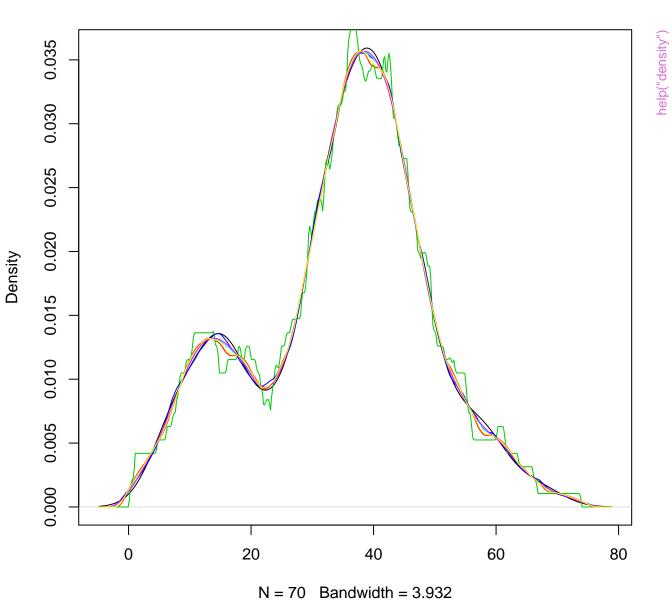
R's density() kernels with bw = 1



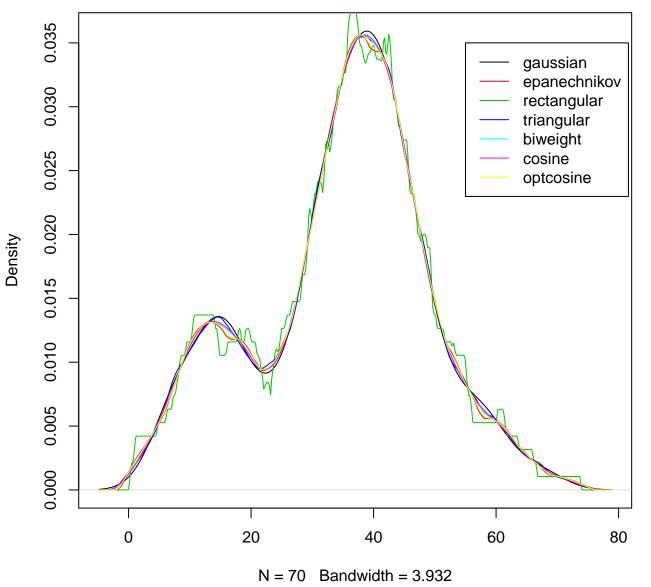
R's density() kernels with width = 1



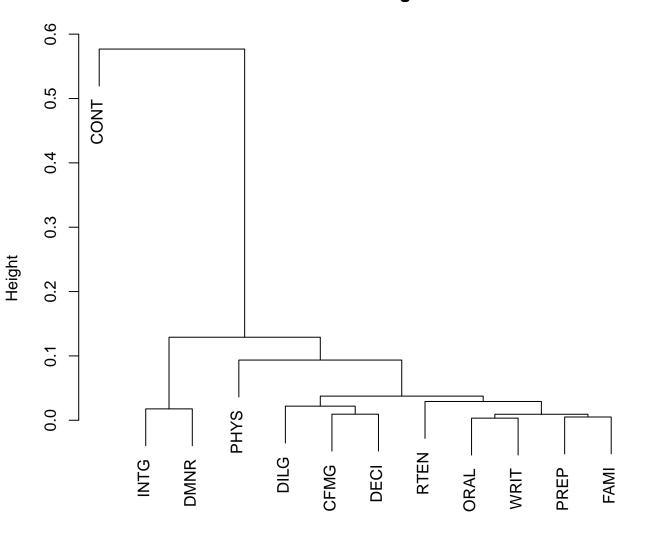
same sd bandwidths, 7 different kernels



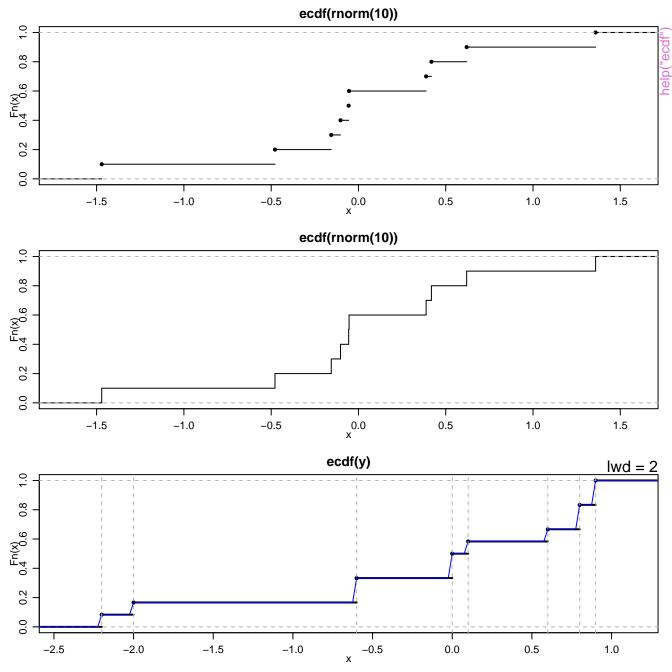
equivalent bandwidths, 7 different kernels

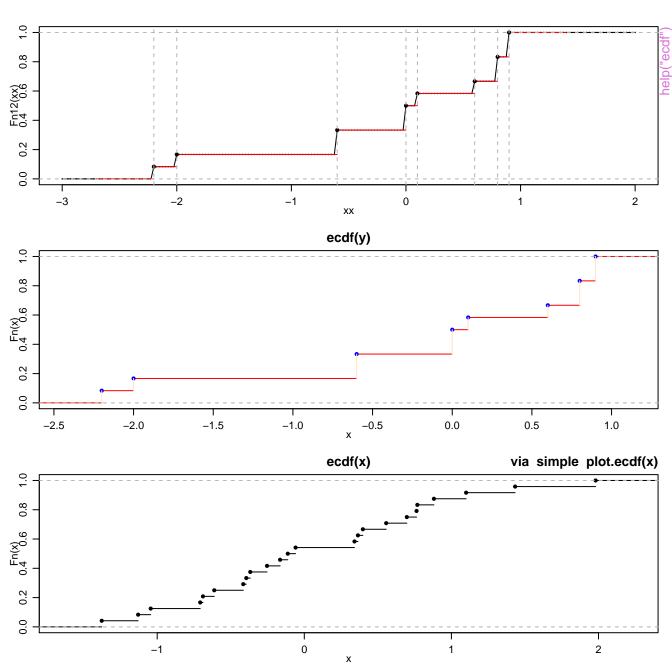


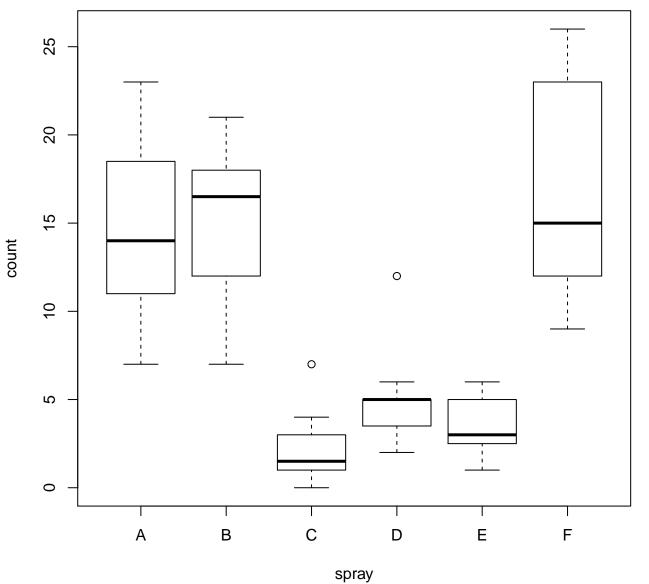
Cluster Dendrogram



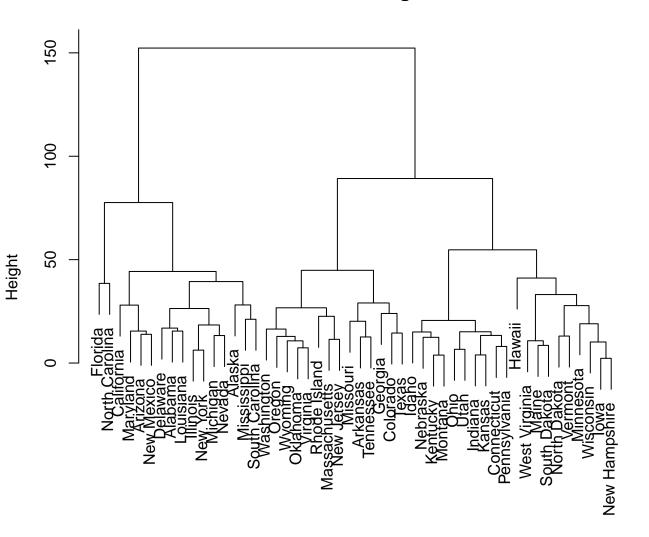
dd hclust (*, "complete")





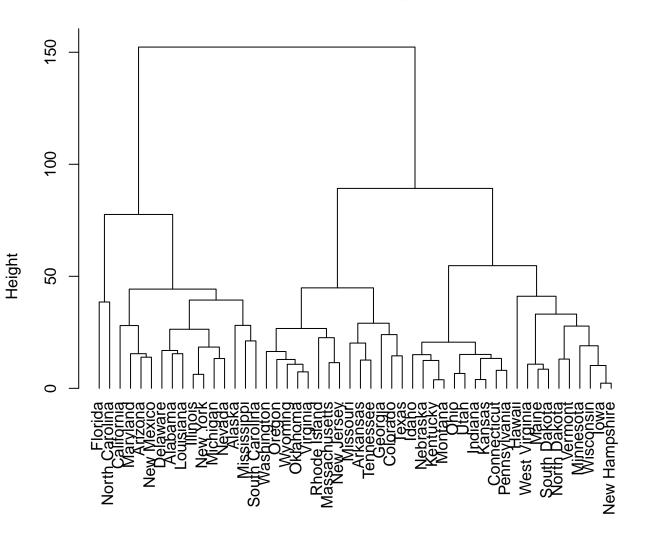


Cluster Dendrogram

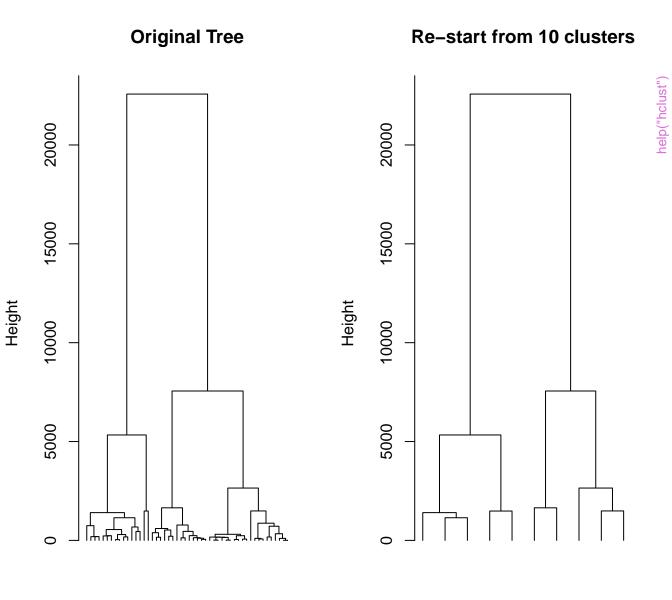


dist(USArrests)
hclust (*, "average")

Cluster Dendrogram

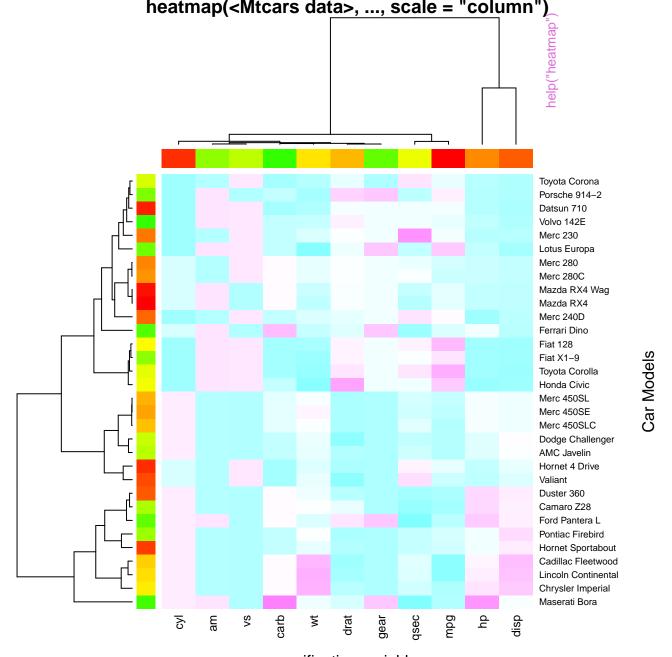


dist(USArrests)
hclust (*, "average")

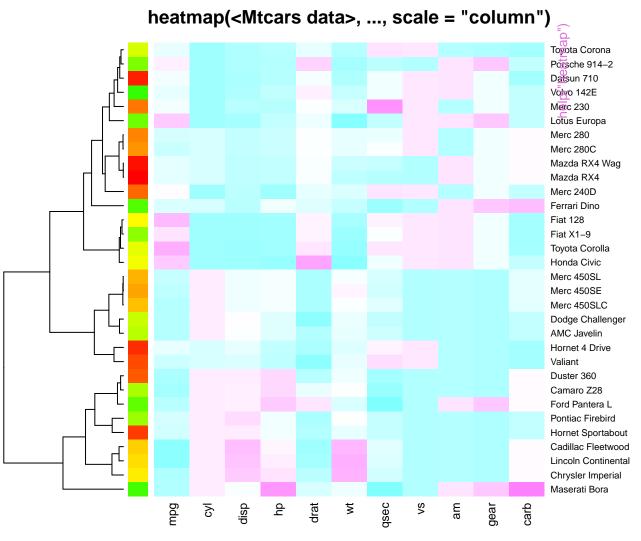


dist(USArrests)^2 hclust (*, "centroid")

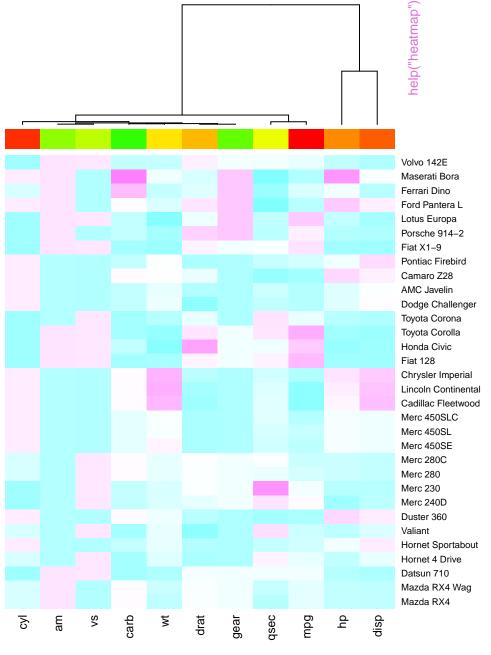
dist(cent)^2 hclust (*, "centroid")



specification variables

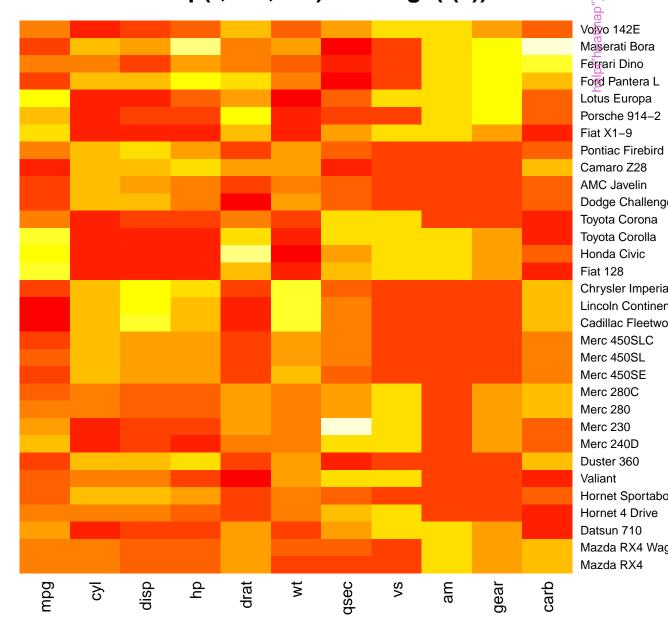


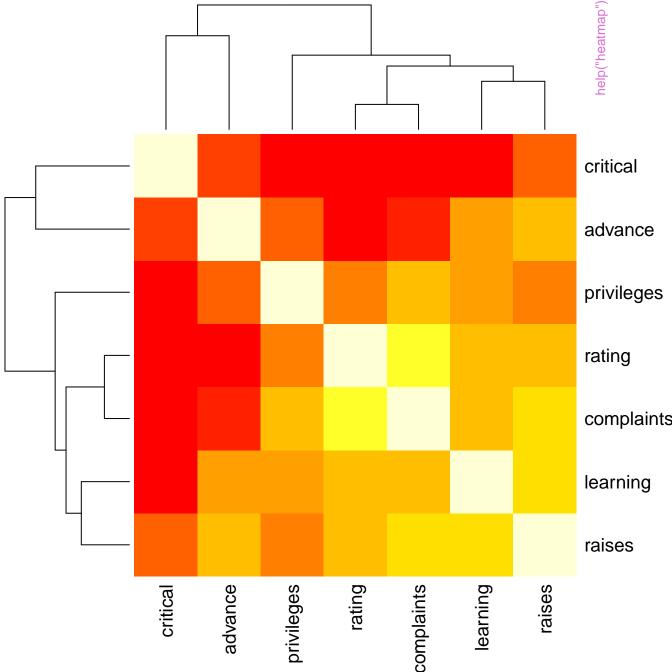
specification variables

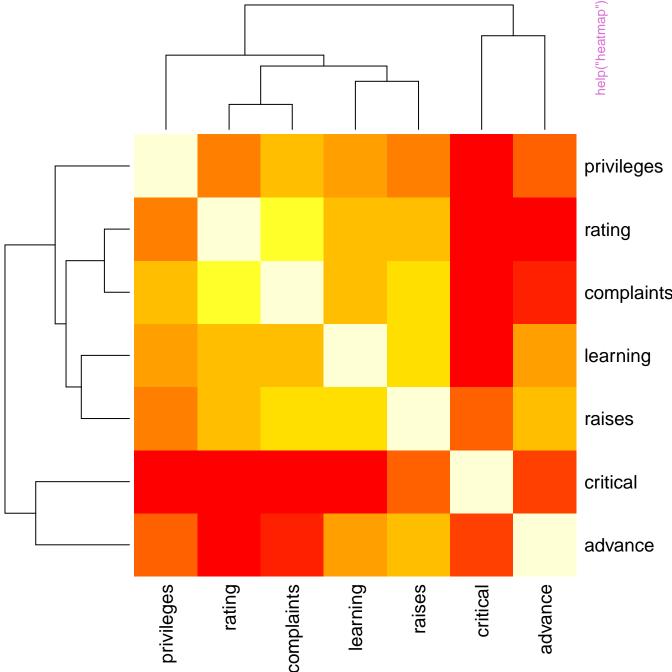


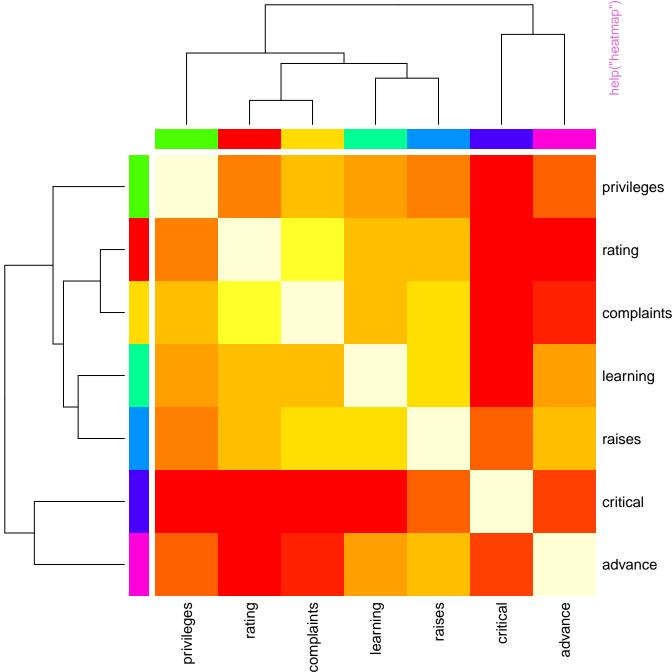
xlab

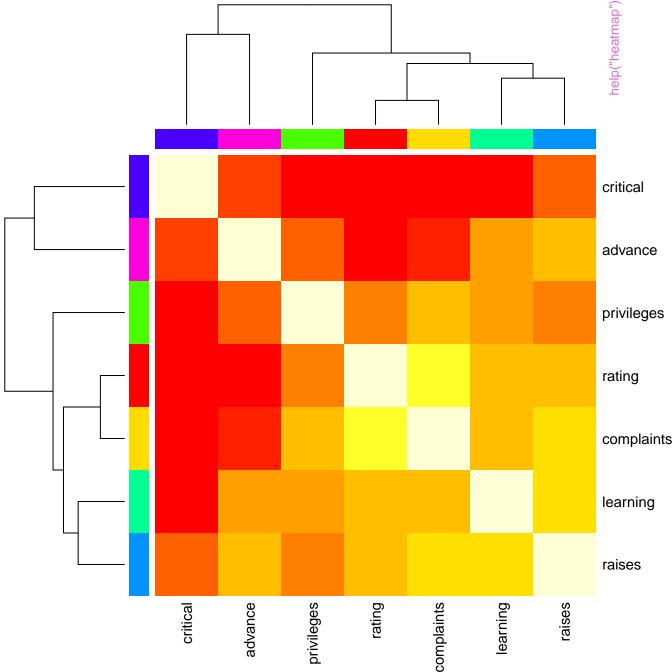
heatmap(*, NA, NA) \sim image(t(x))

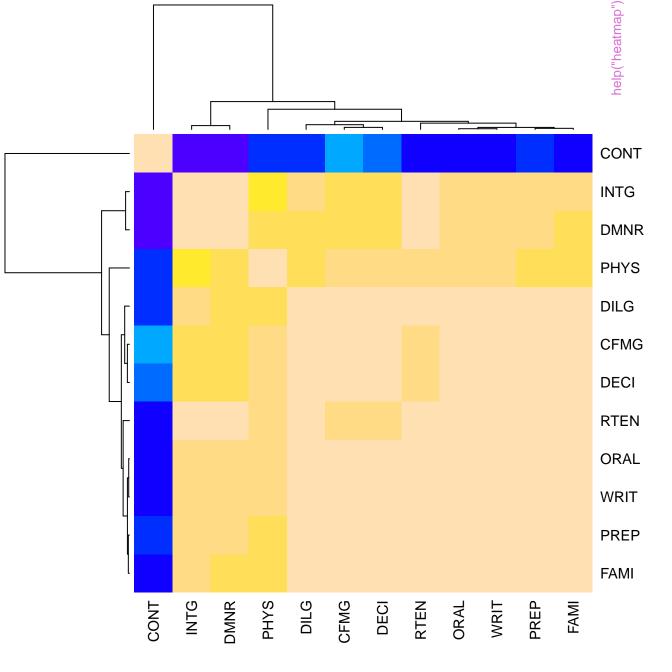


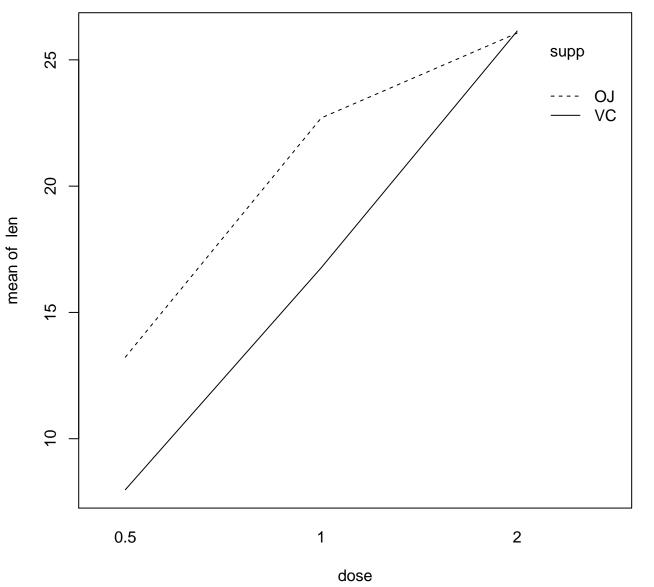


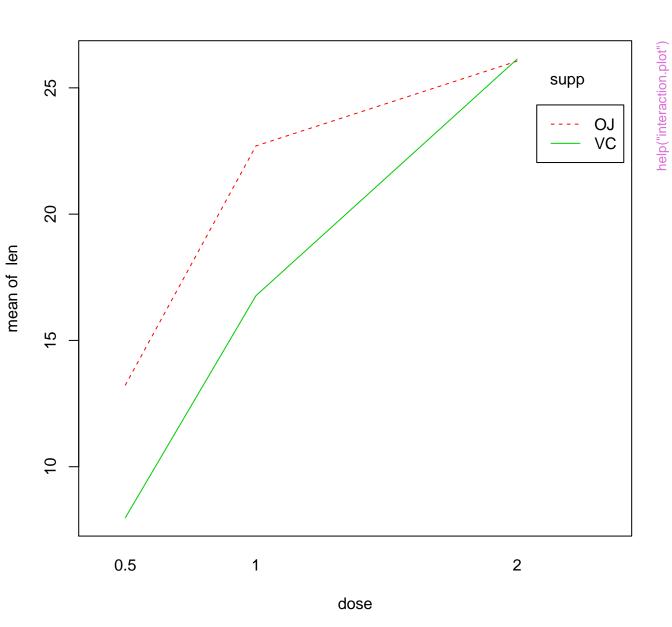


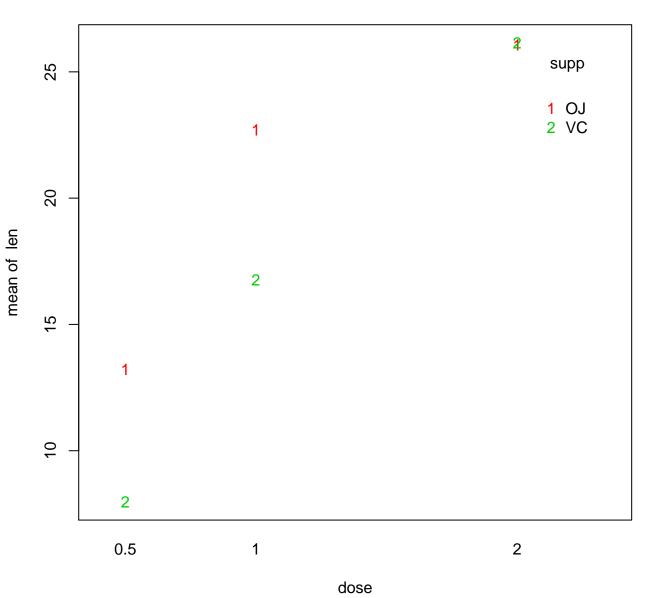


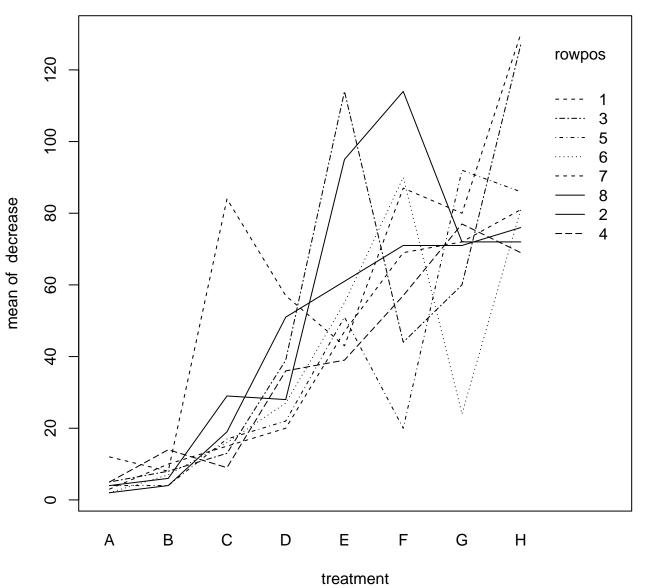


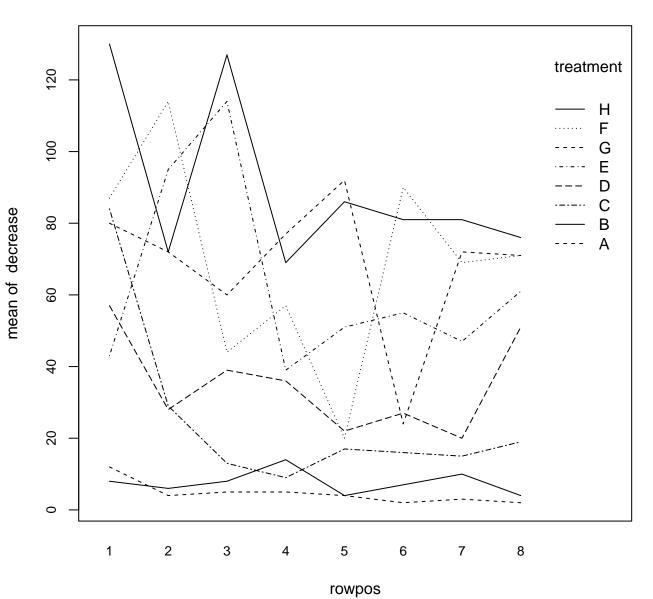


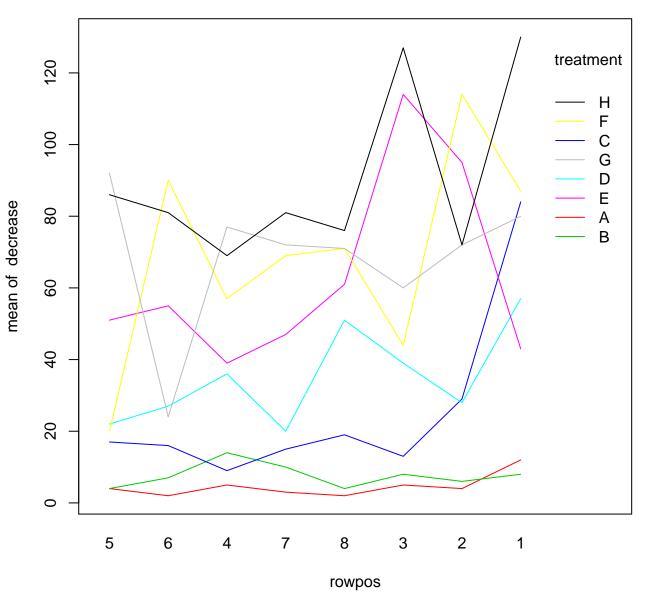




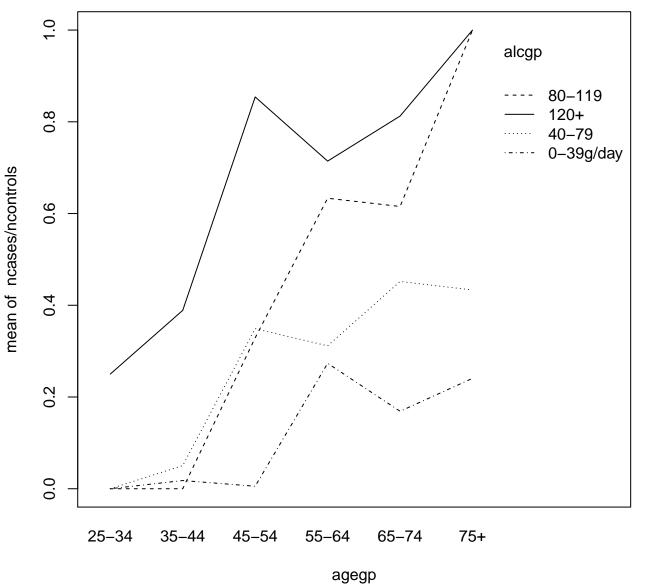


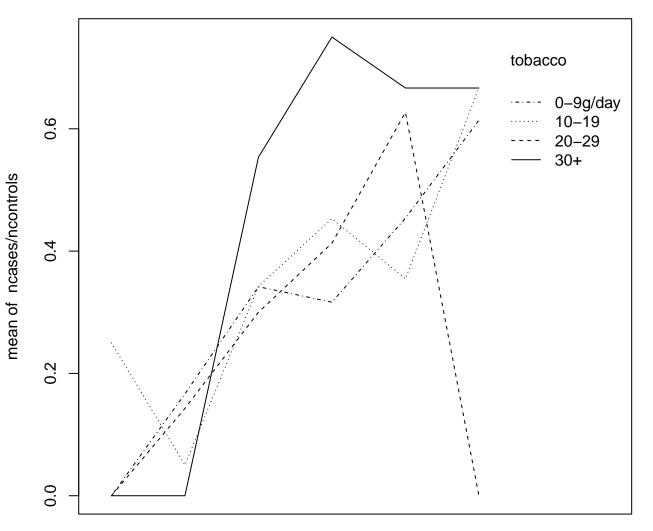


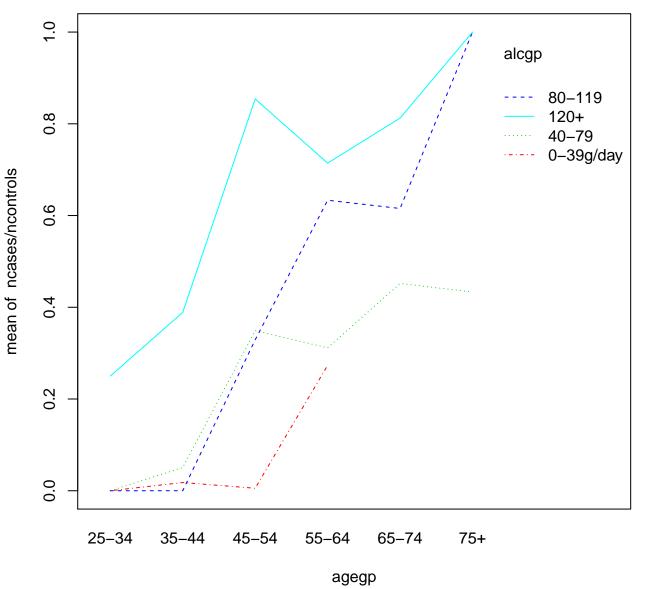


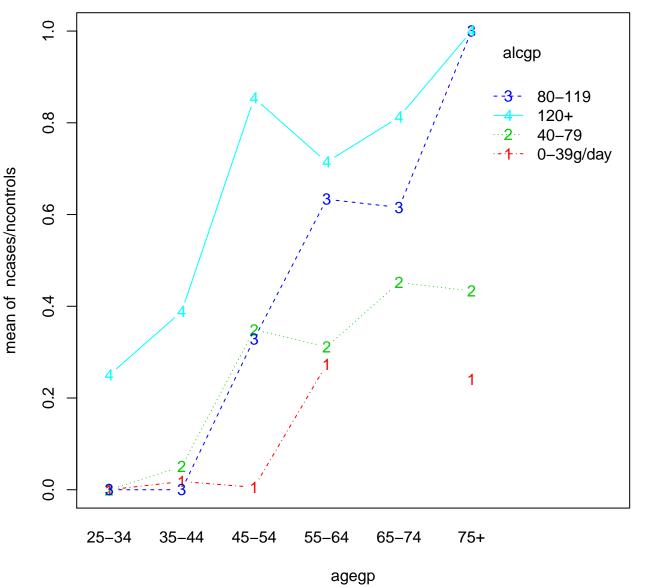


'esoph' Data



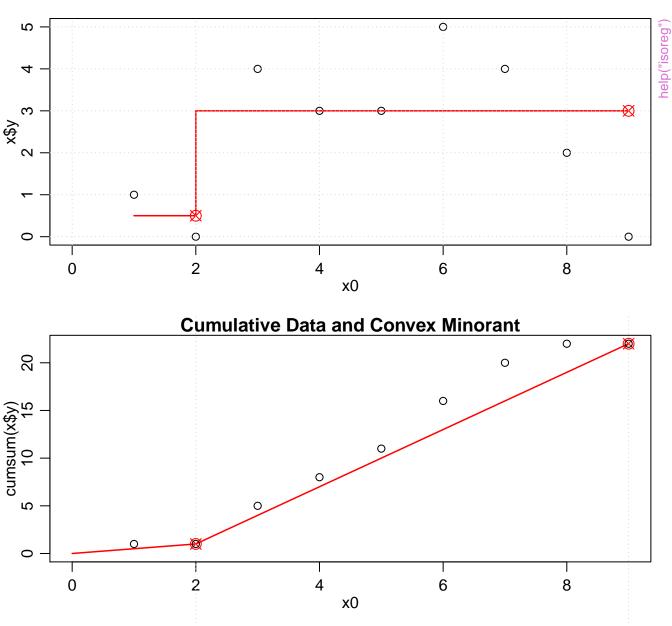


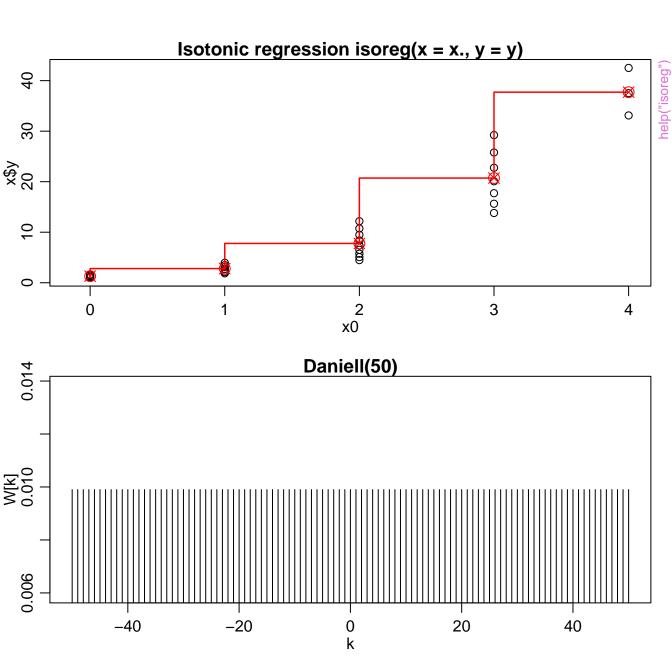


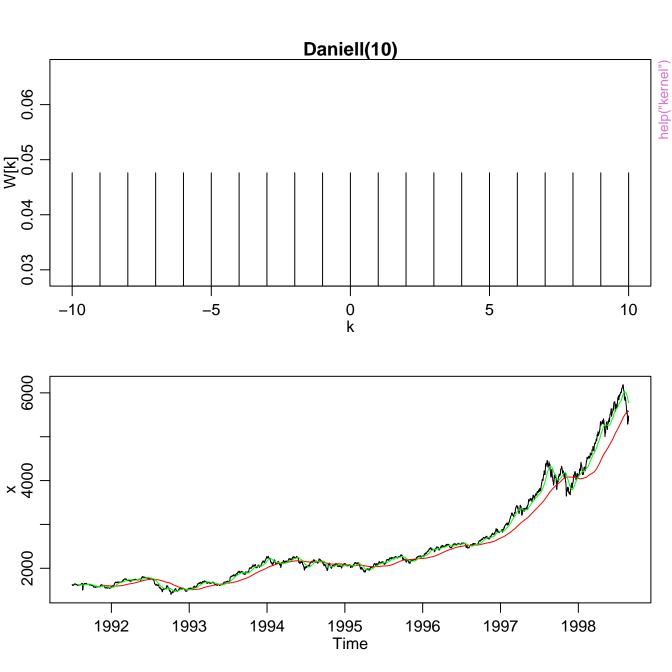


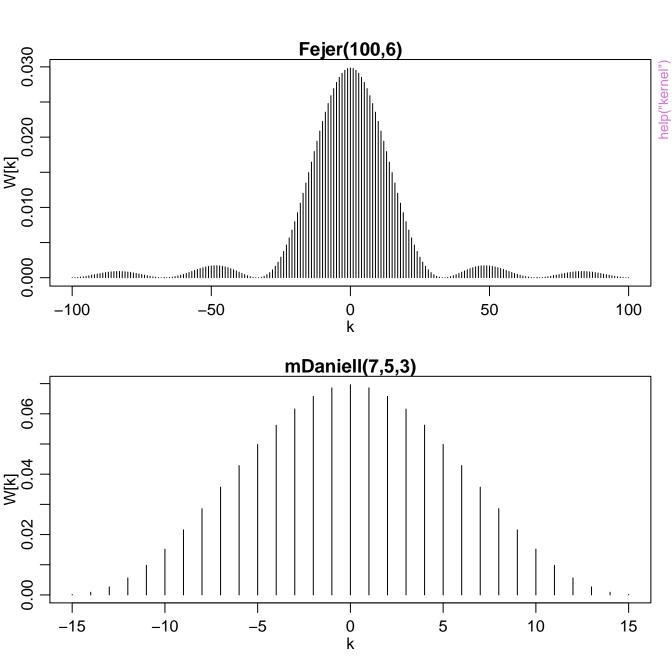
agegp
function(x) mean(x, na.rm=TRUE)

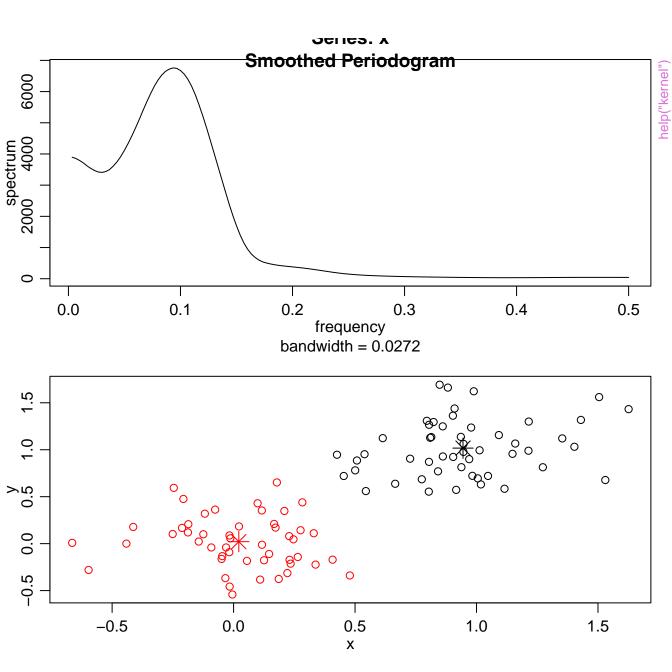
Isotonic regression isoreg(x = c(1, 0, 4, 3, 3, 5, 4, 2, 0))

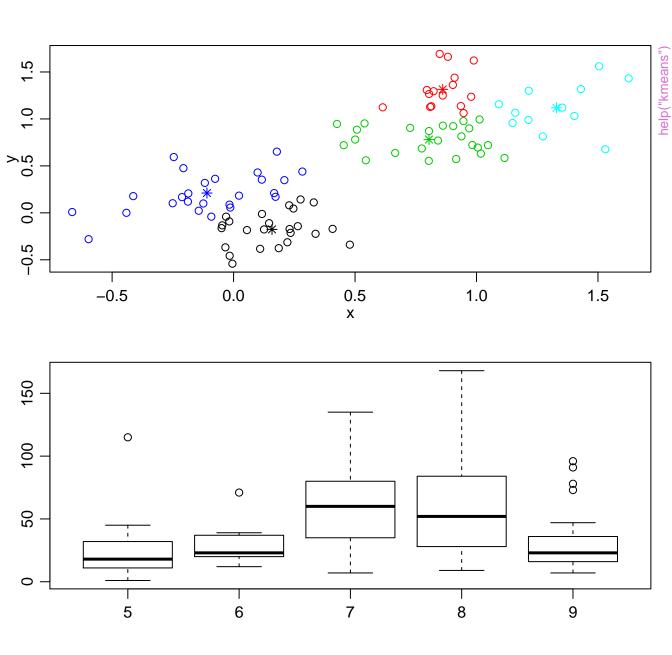


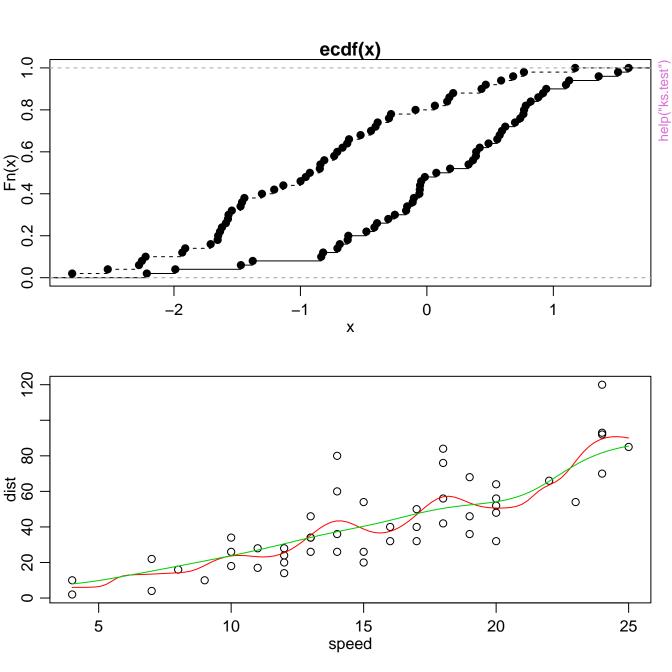


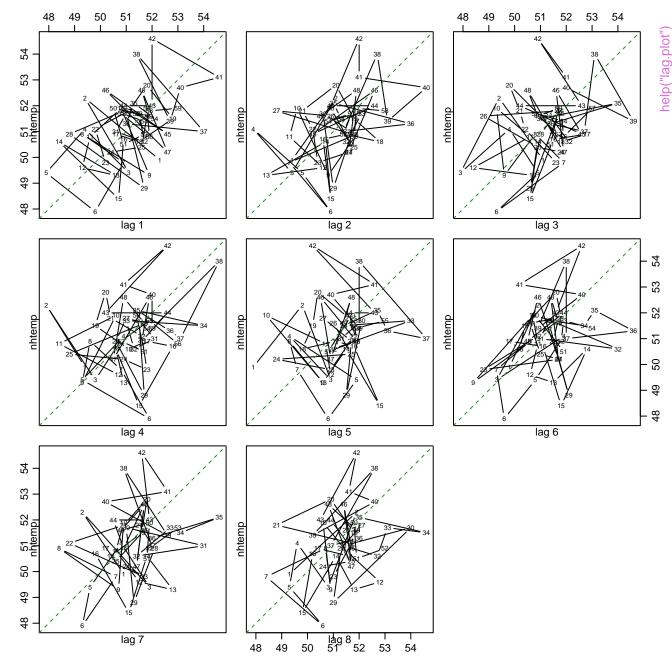




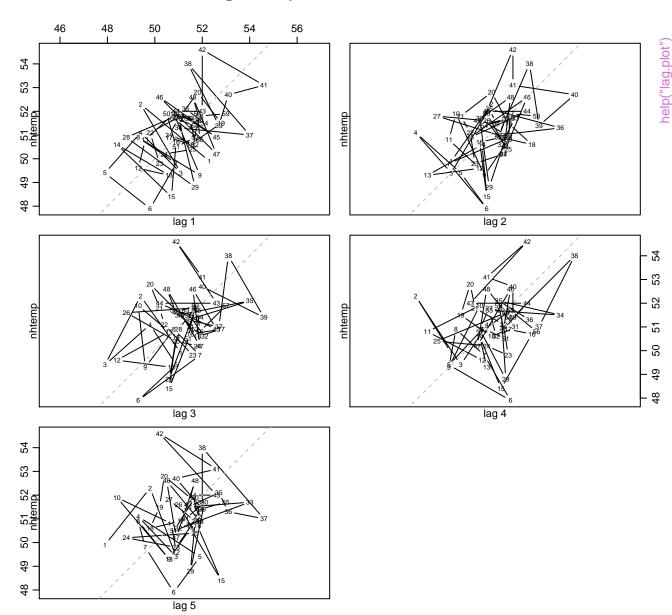




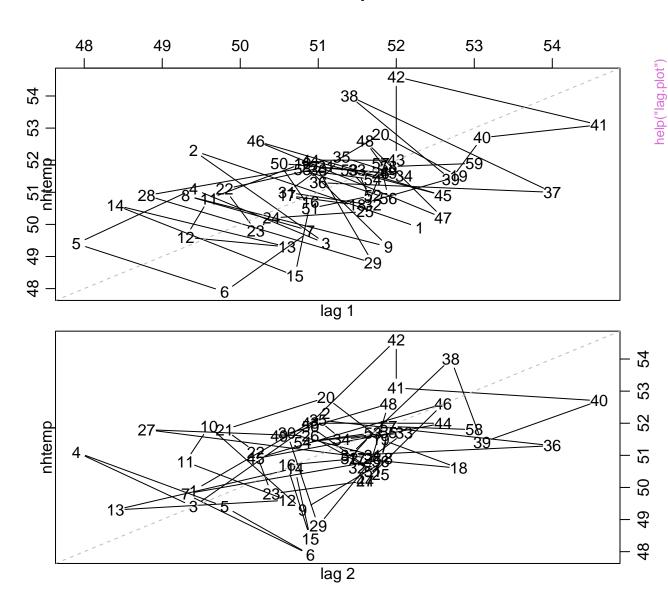




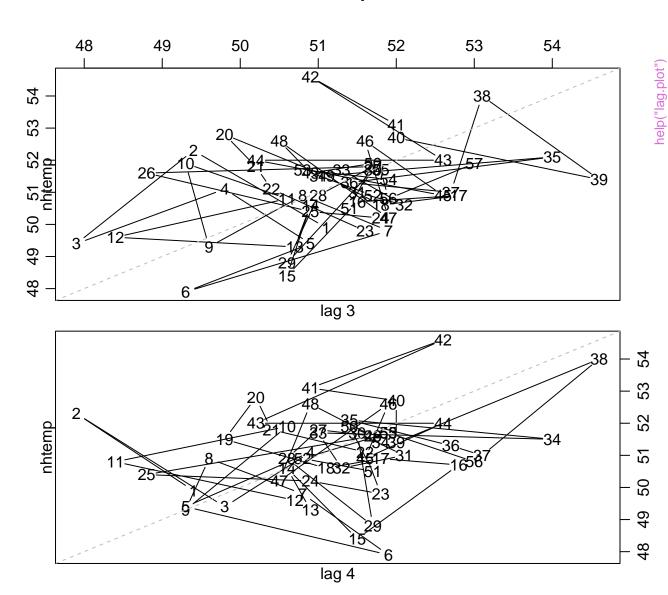
Average Temperatures in New Haven



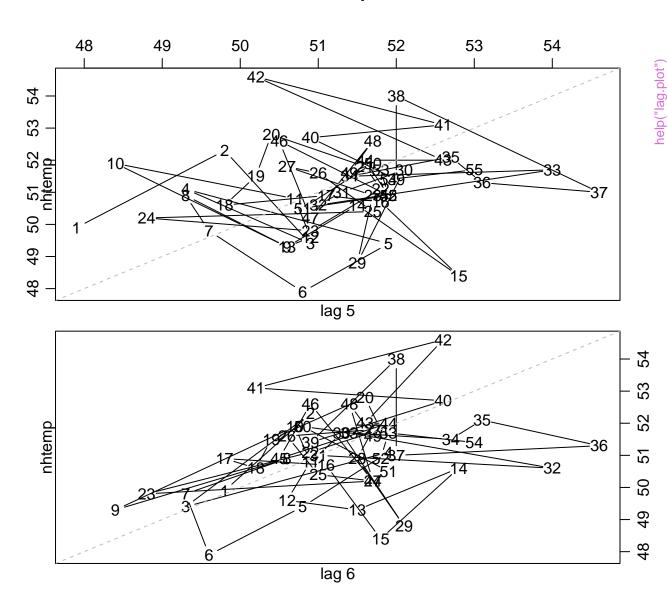
New Haven Temperatures

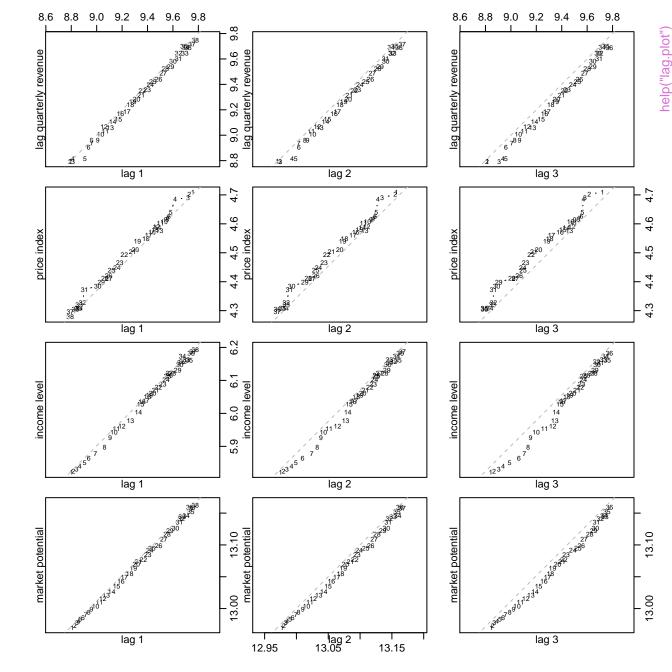


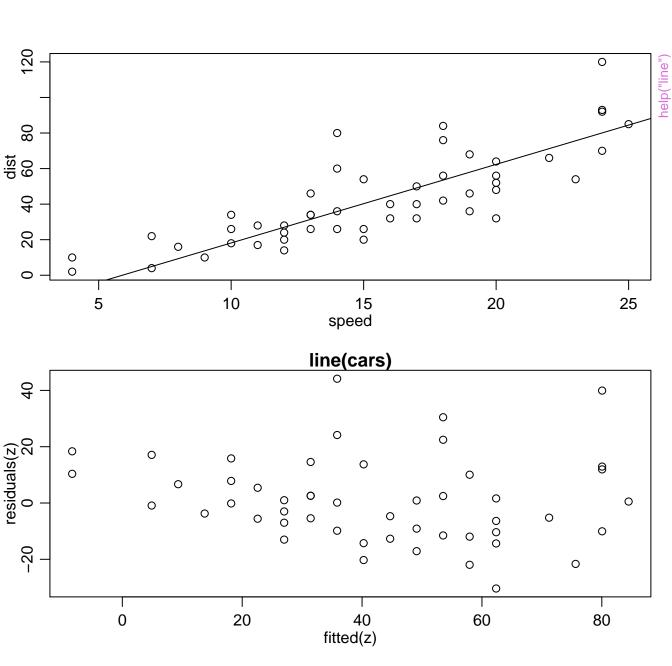
New Haven Temperatures

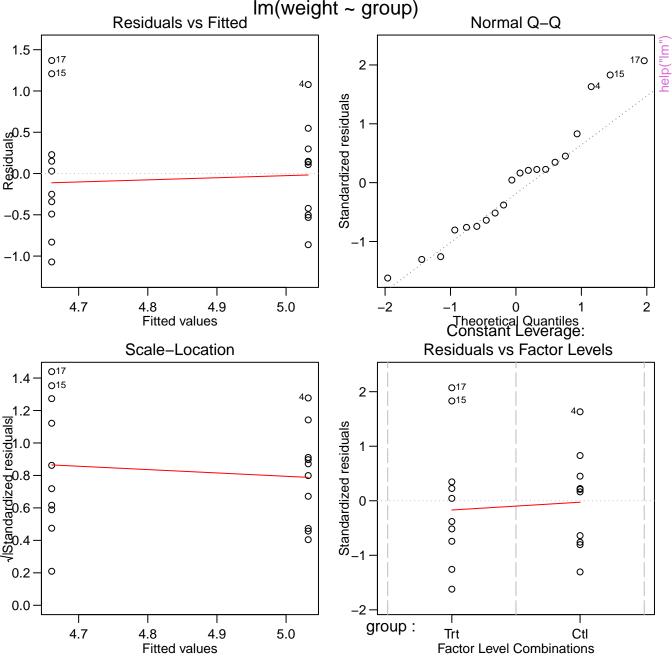


New Haven Temperatures

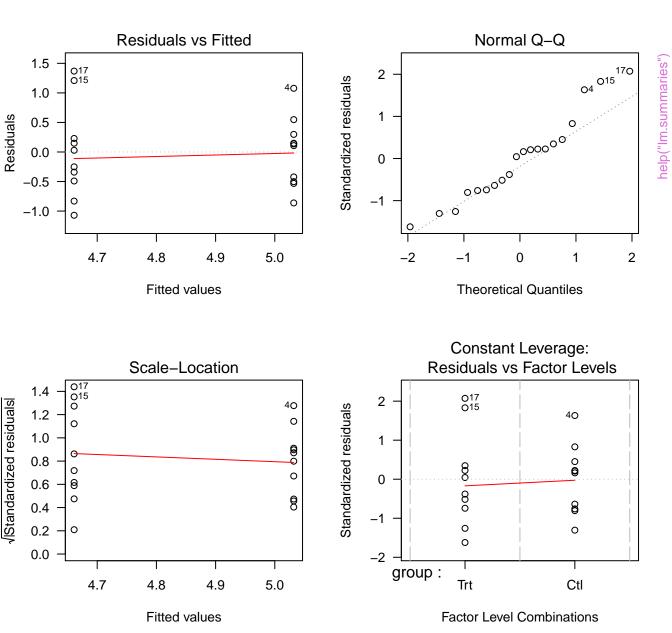


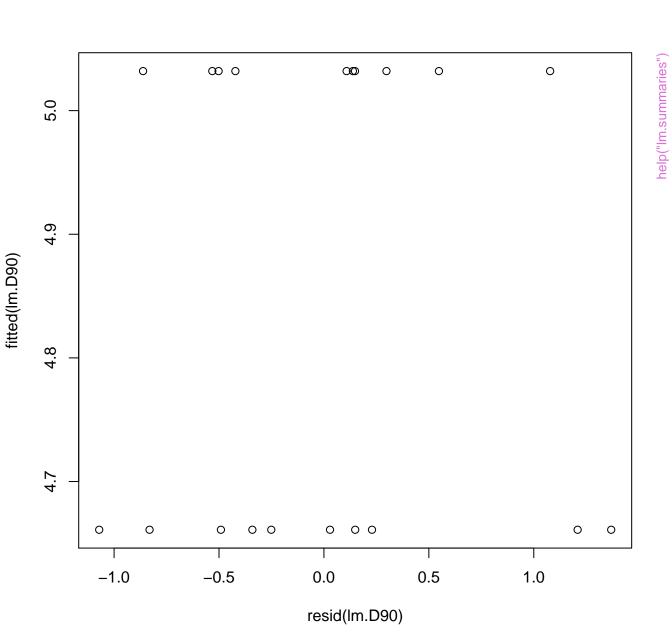




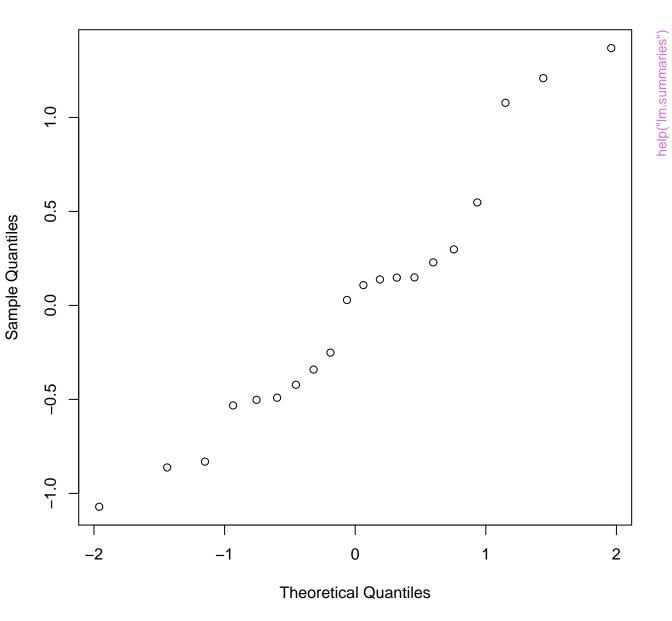


lm(weight ~ group)

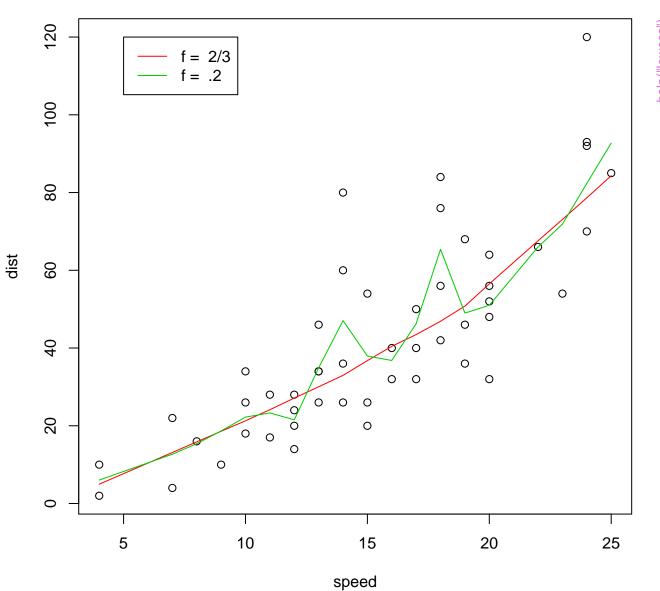




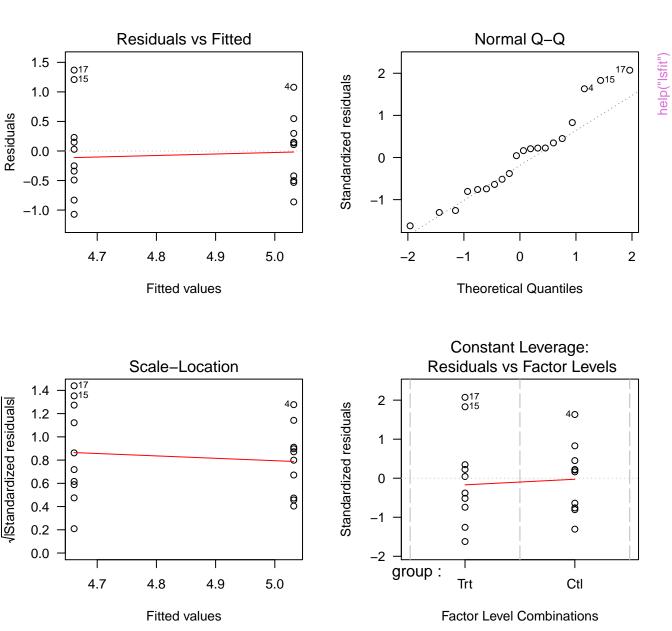
Normal Q-Q Plot



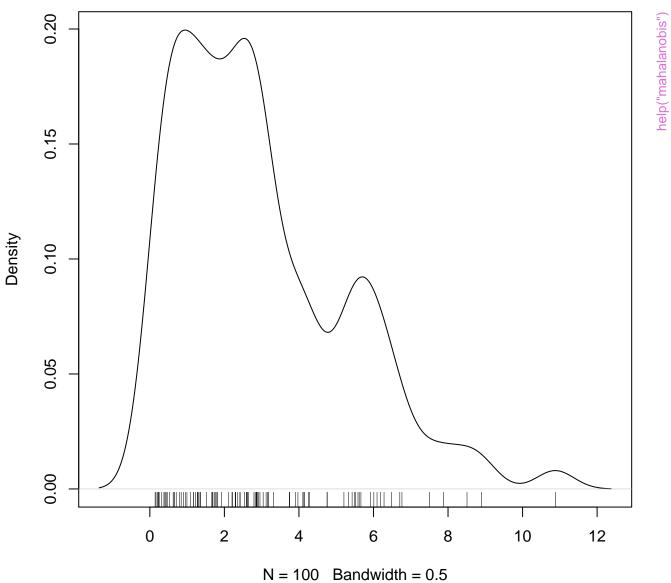




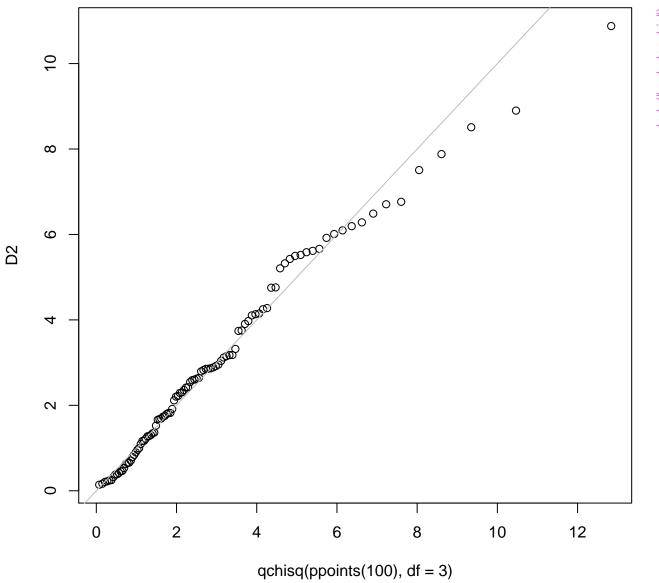
Im(weight ~ group)

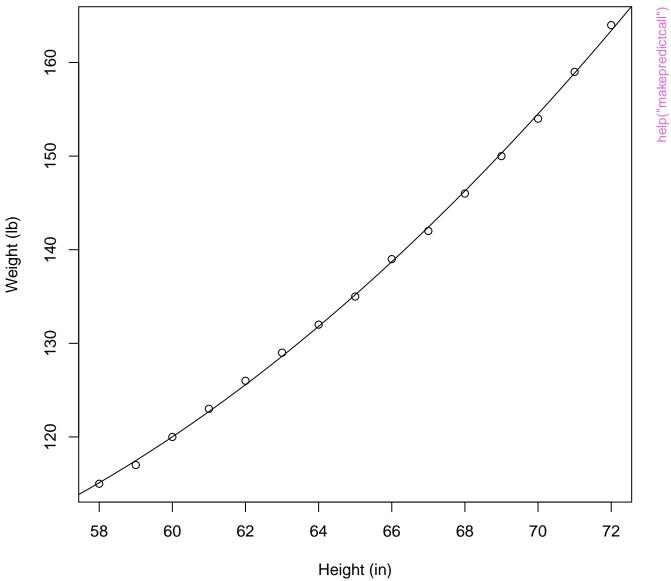


Squared Mahalanobis distances, n=100, p=3

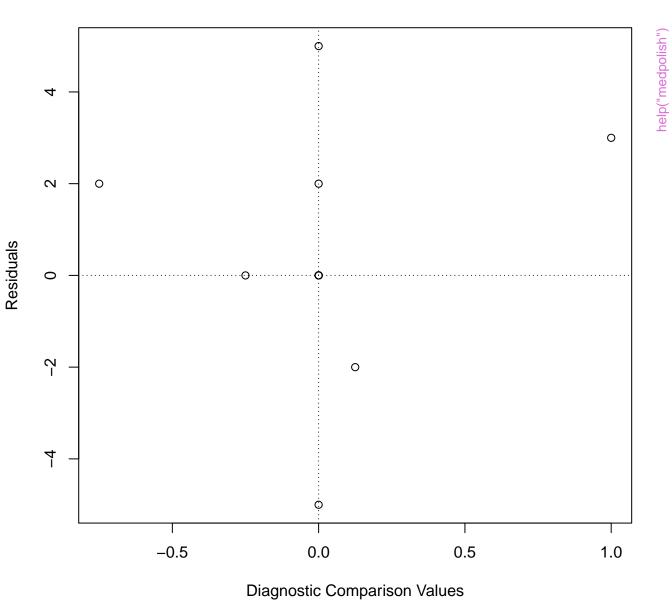


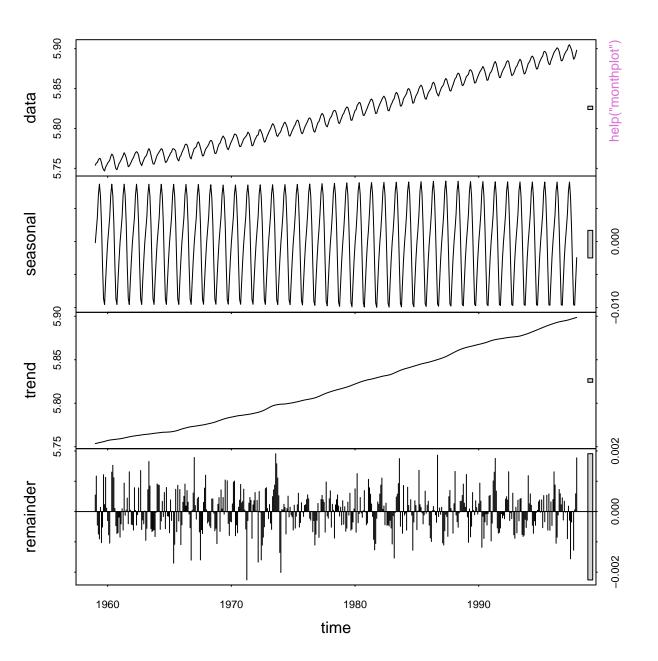


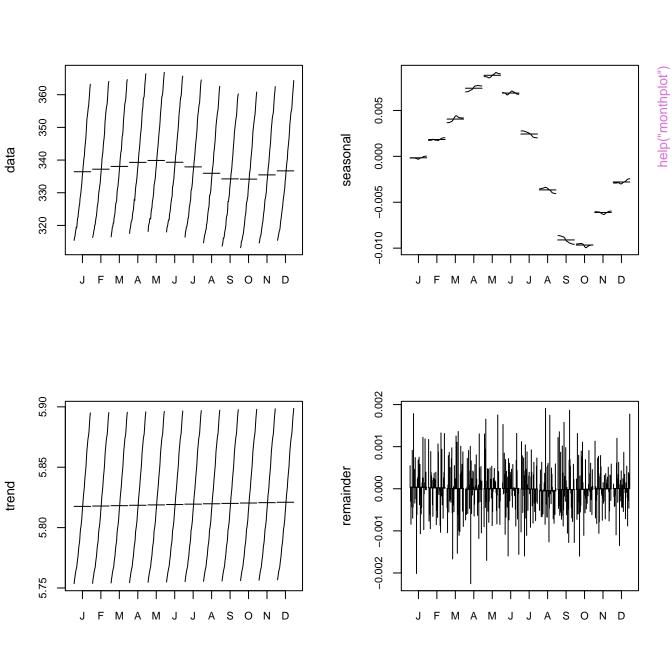


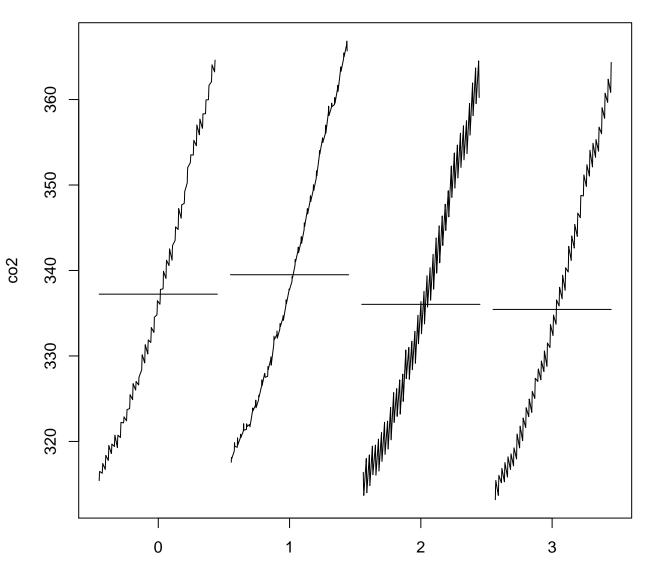


Tukey Additivity Plot

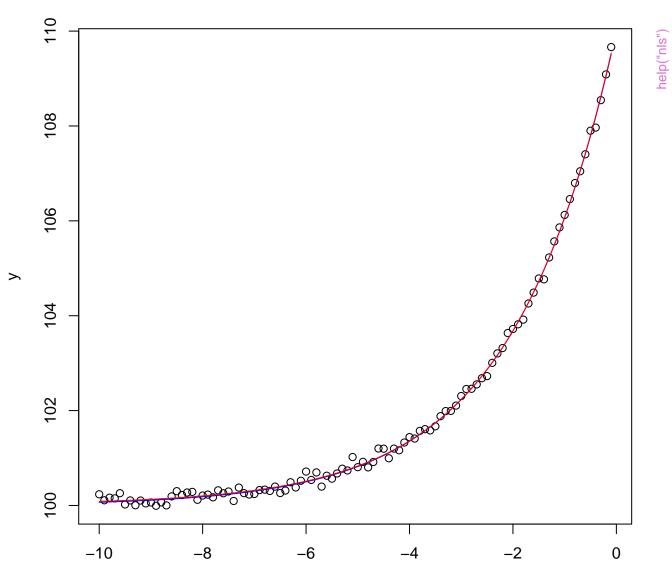




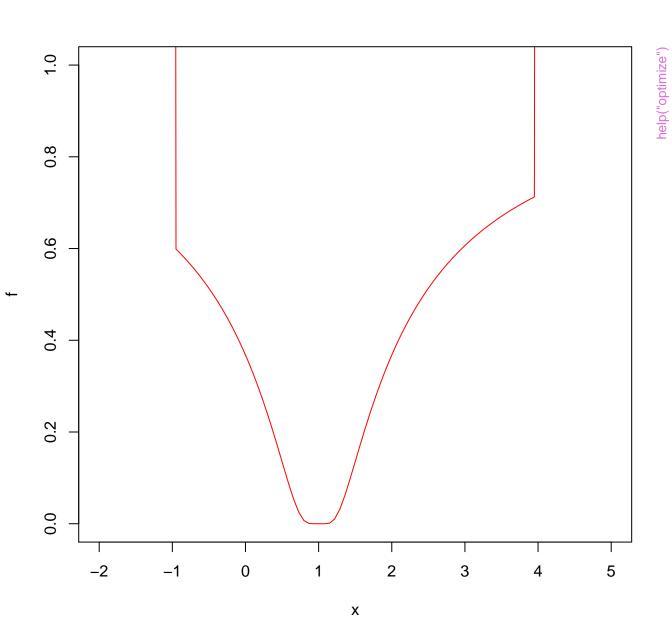




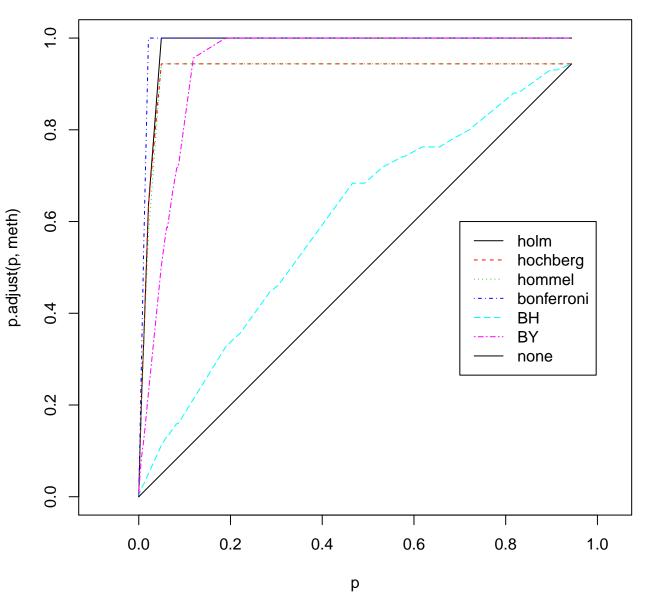
nls(*), data, true function and fit, n=100

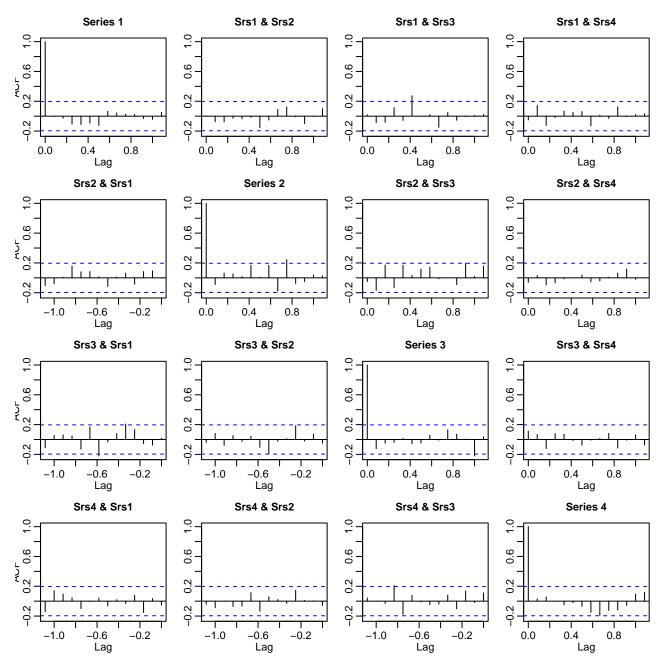


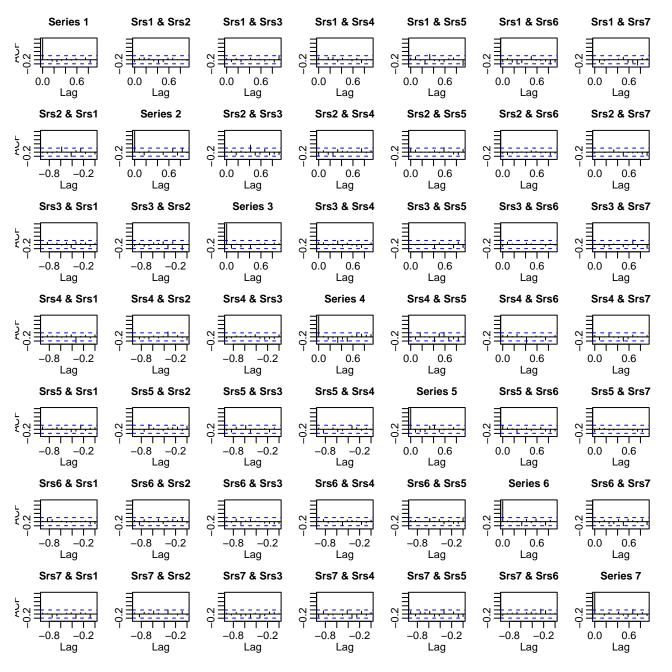
Χ

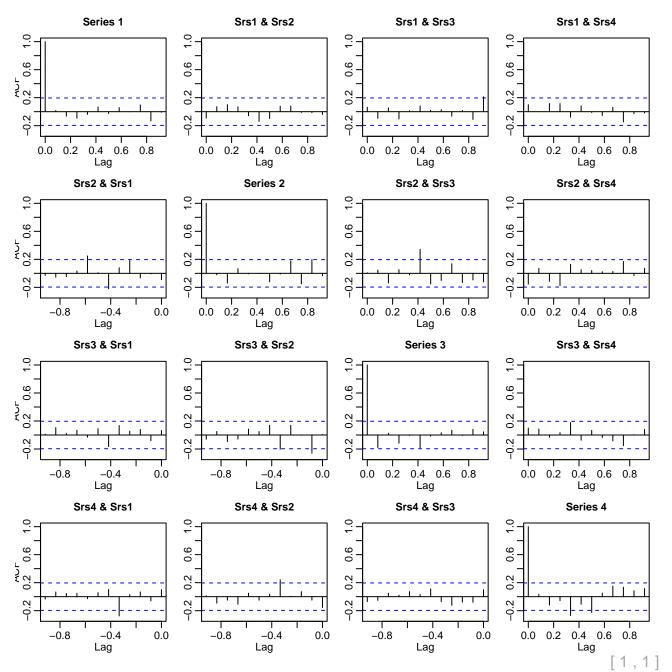


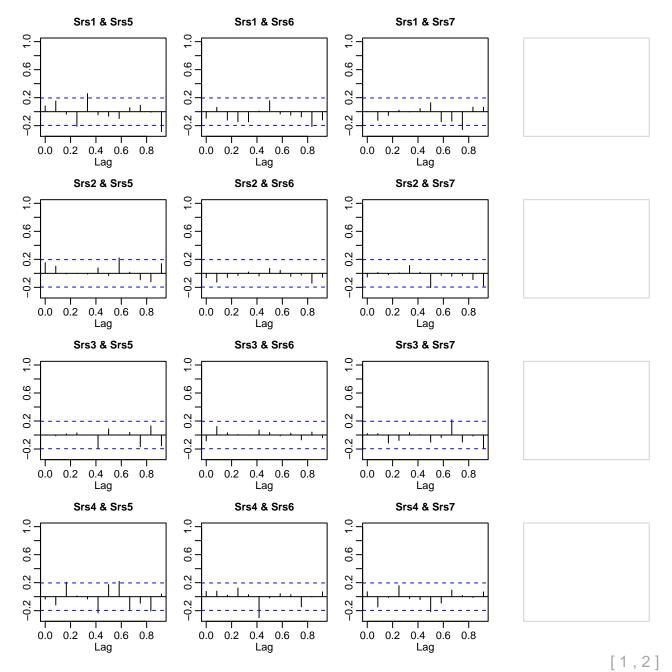
P-value adjustments

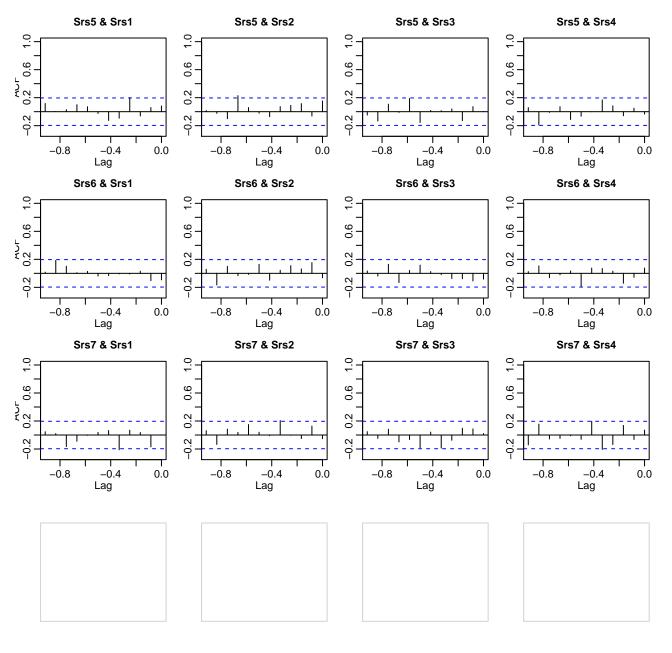


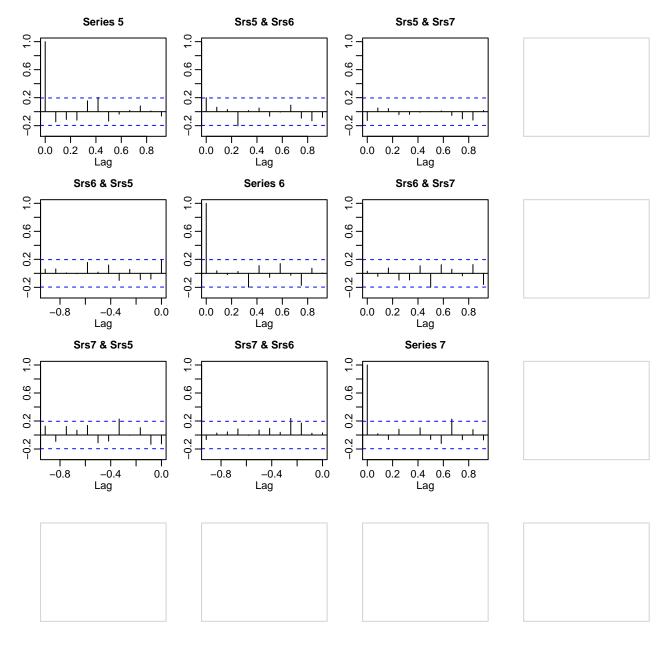




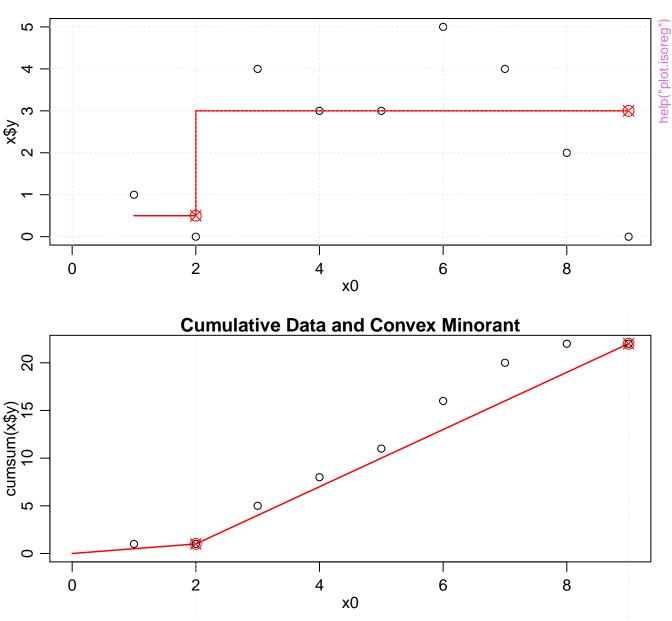


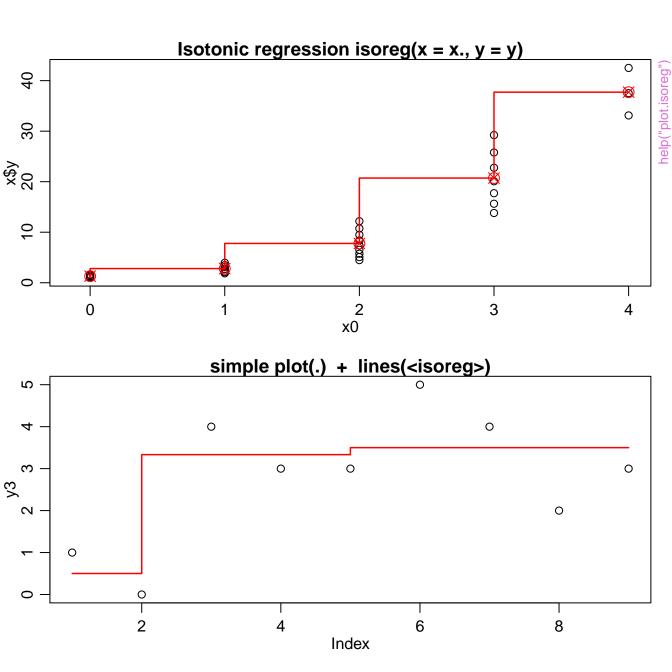




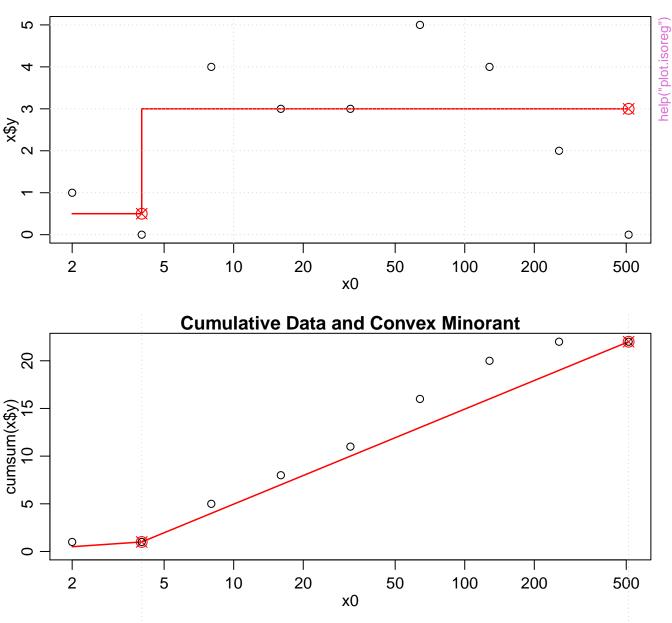


Isotonic regression isoreg(x = c(1, 0, 4, 3, 3, 5, 4, 2, 0))

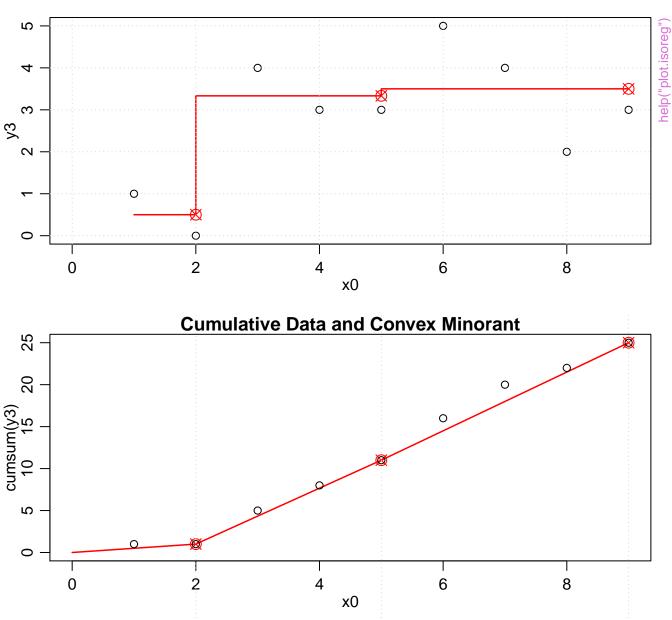




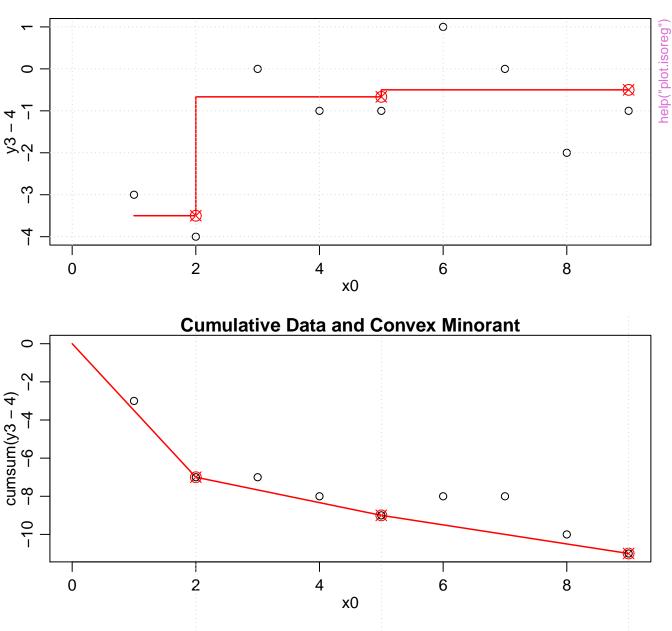
Isotonic regression isoreg(x = $2^{(1:9)}$, y = c(1, 0, 4, 3, 3, 5, 4, 2, 0))



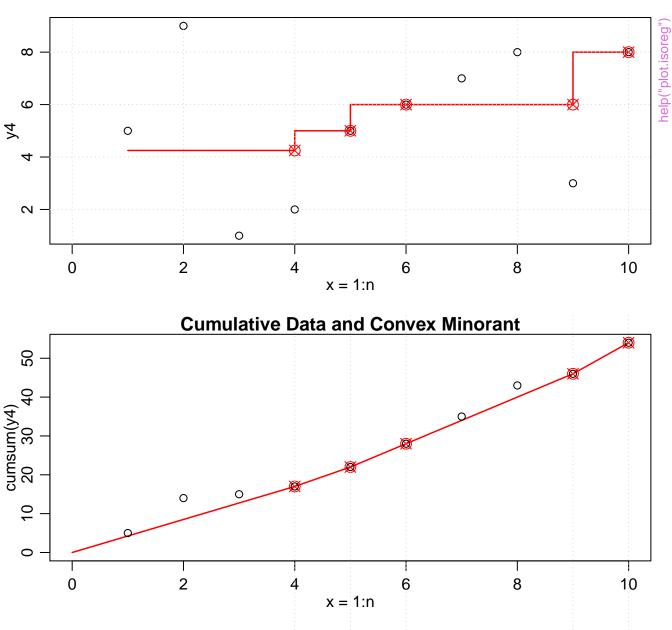
Isotonic regression isoreg(x = y3 < -c(1, 0, 4, 3, 3, 5, 4, 2, 3))



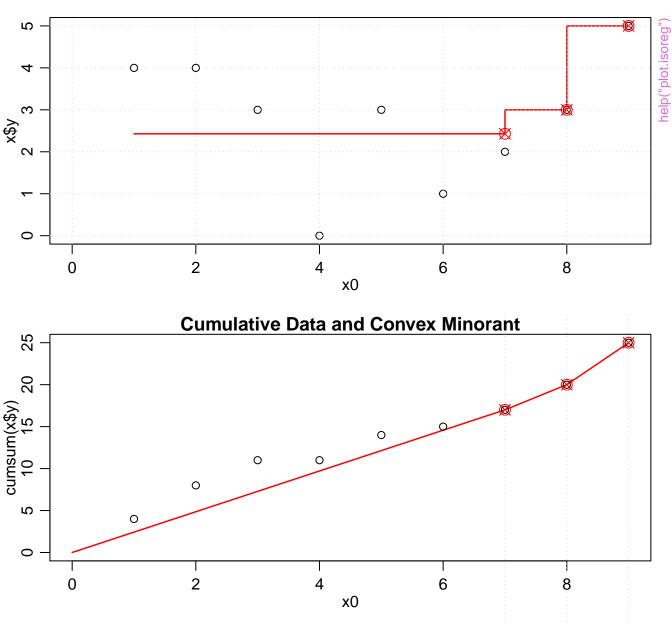
Isotonic regression isoreg(x = y3 - 4)



Isotonic regression isoreg(x = 1:10, y = y4 < -c(5, 9, 1:2, 5:8, 3, 8))



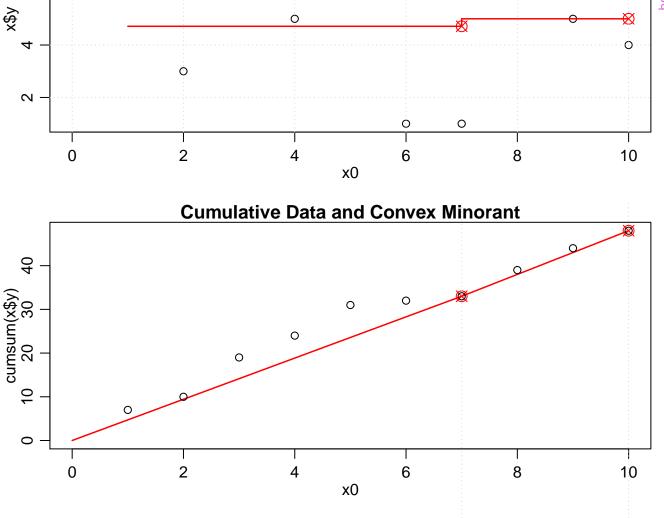
Isotonic regression isoreg(x = sample(9), y = y3)

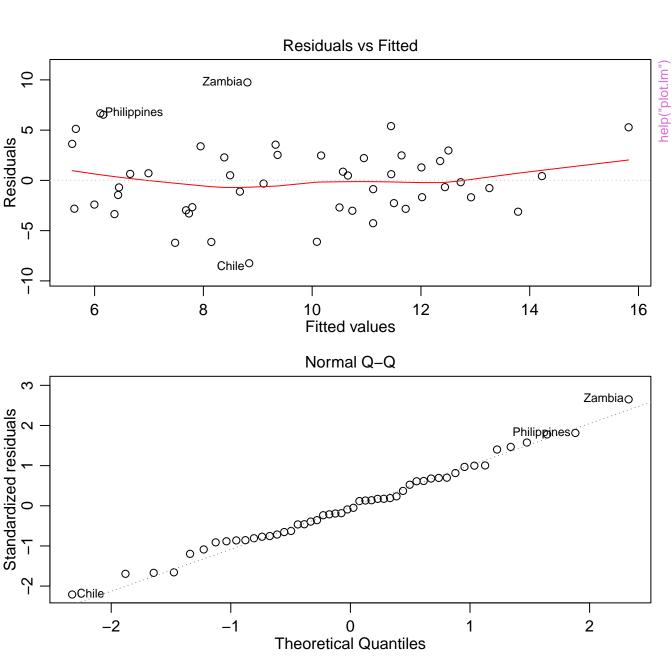


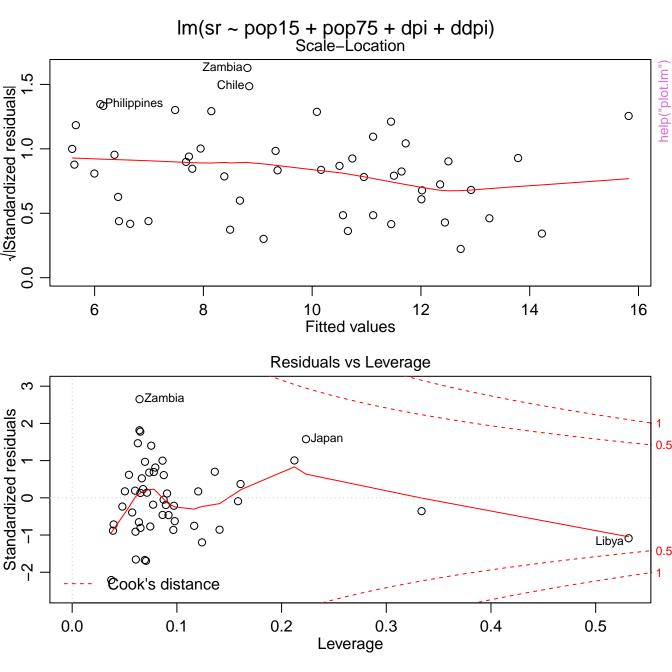
Isotonic regression isoreg(x = sample(9), y = y3)

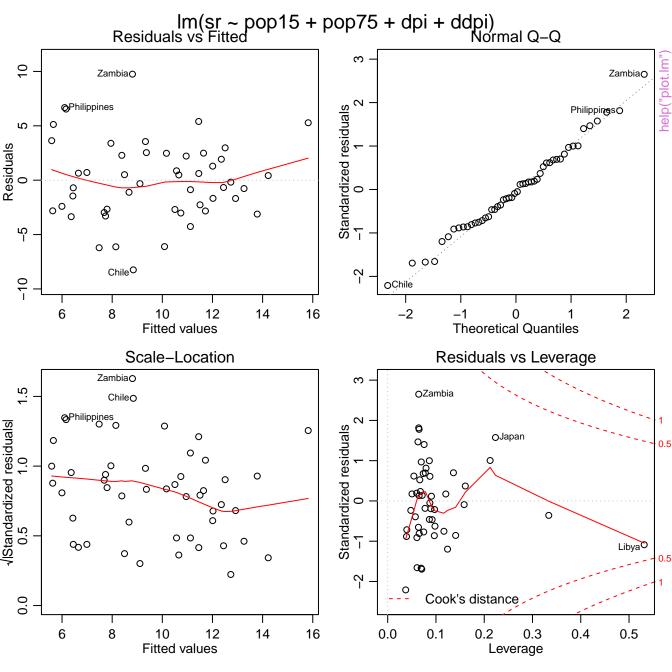
Cumulative Data and Convex Minora help("plot.isoreg cumsum(x\$y) x\$y \sim x0 x0

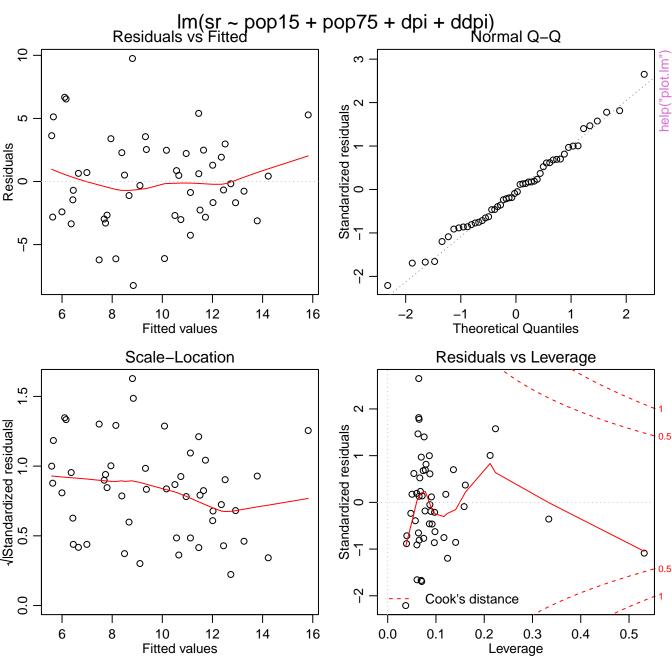
Isotonic regression isoreg(x = sample(10), y = sample(10, replace = TRUE)) help("plot.isoreg' ∞ 0 0 Ó 2 10 0 8 x0 **Cumulative Data and Convex Minorant**

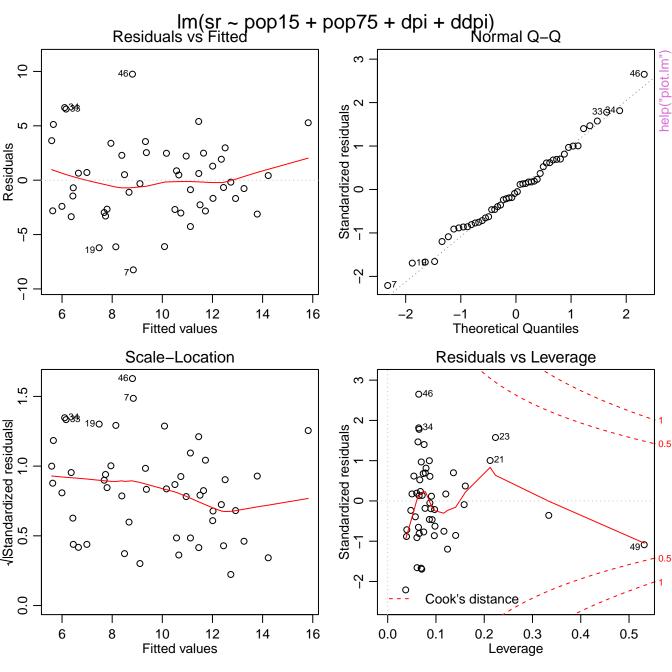


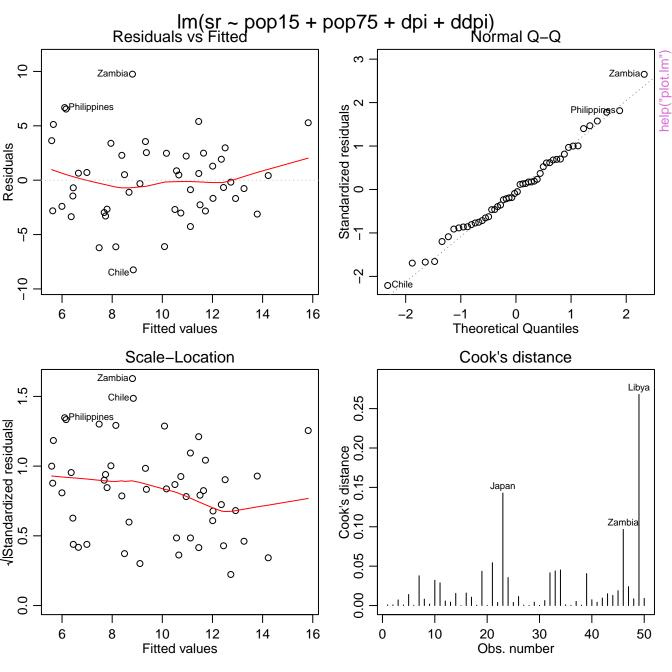


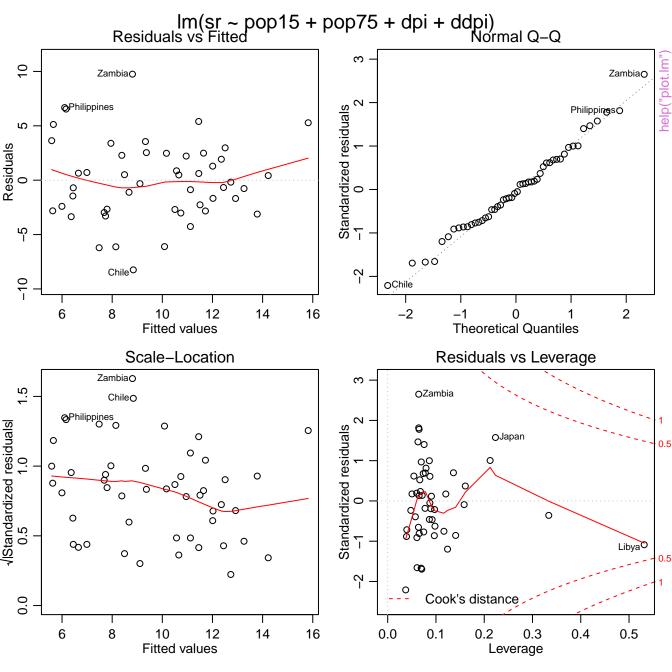


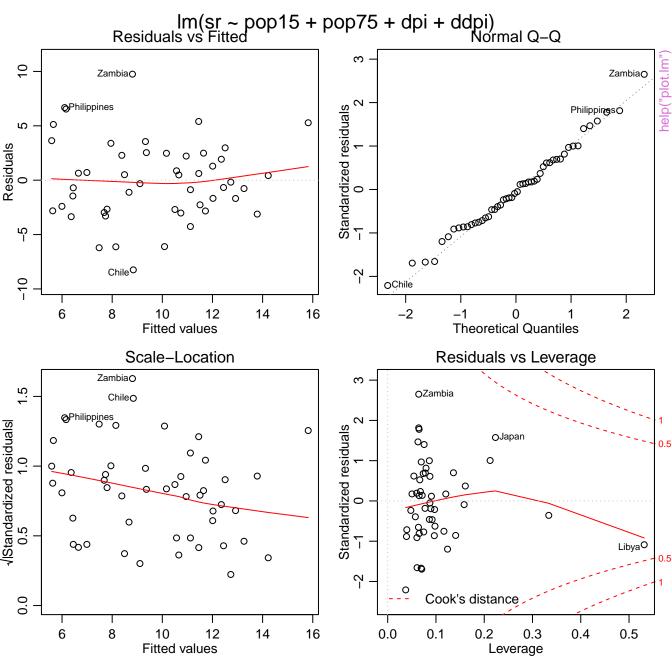


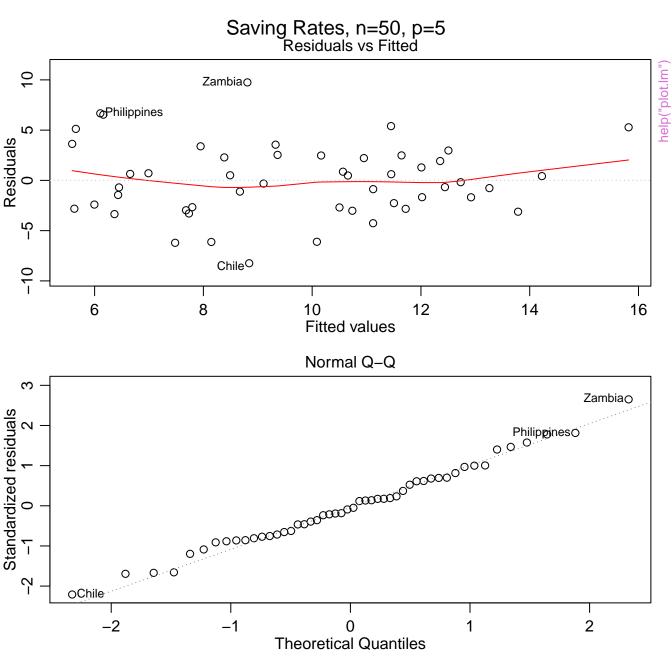


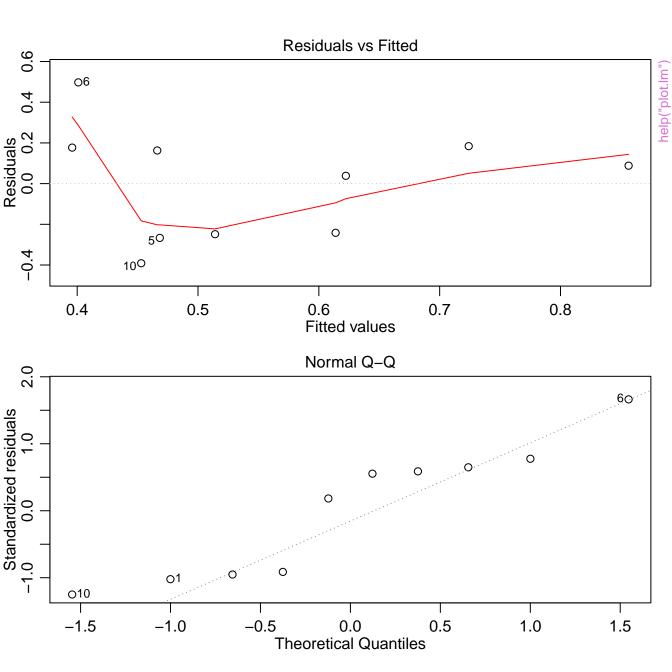


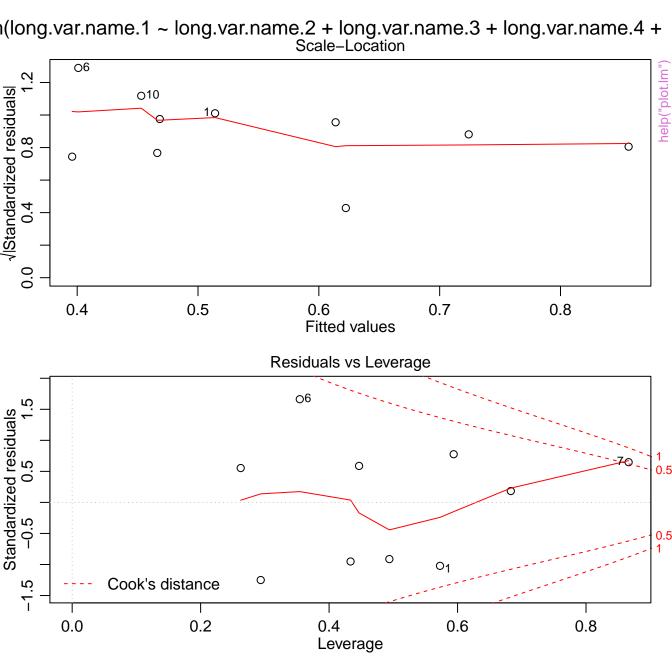


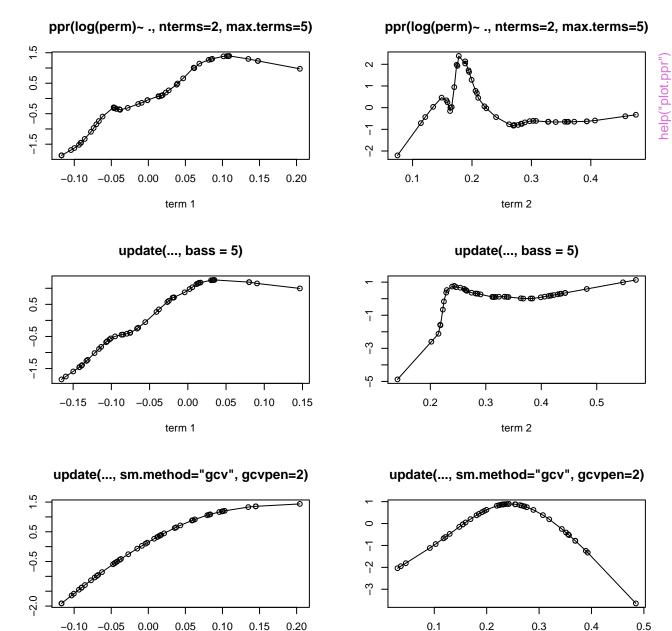






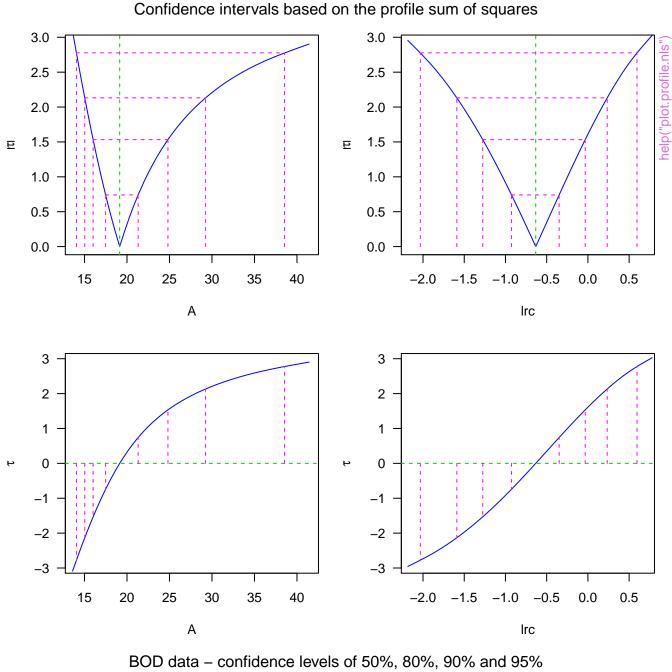


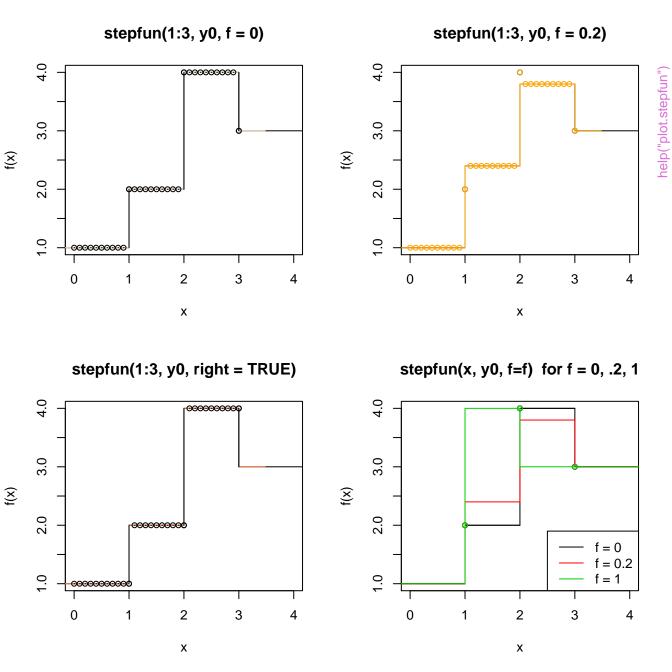




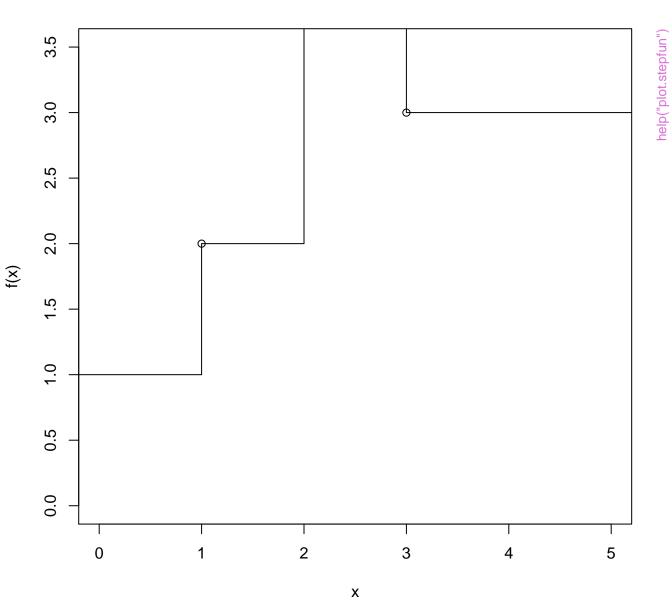
term 2

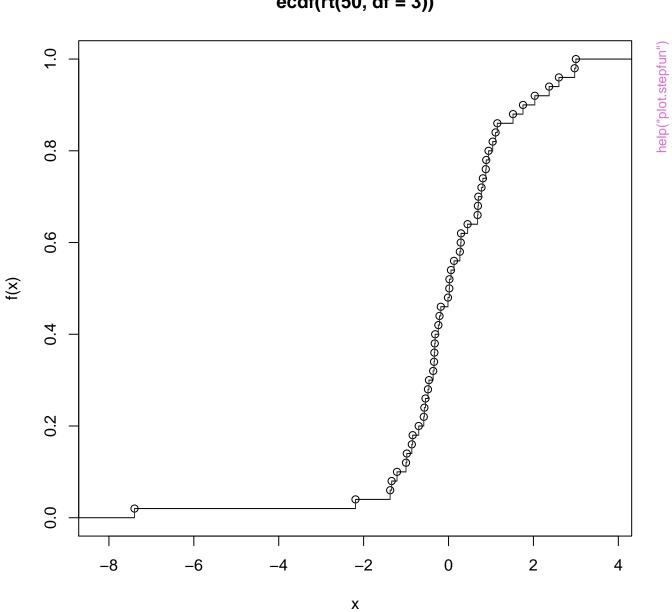
term 1

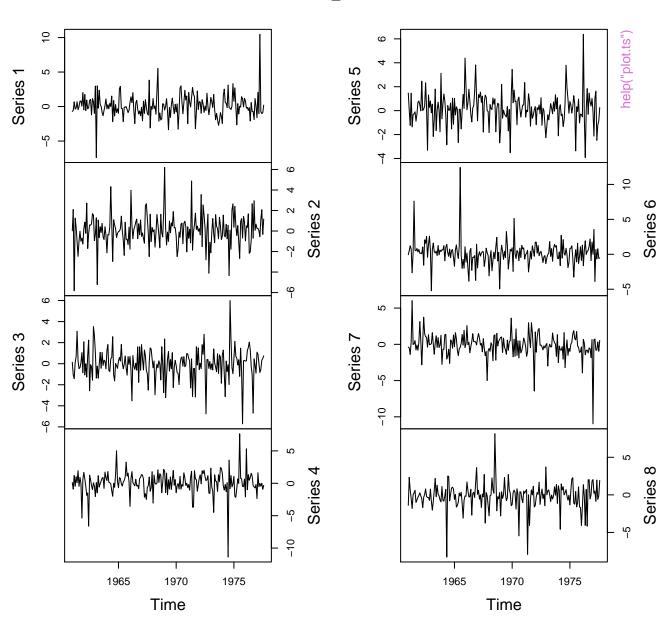




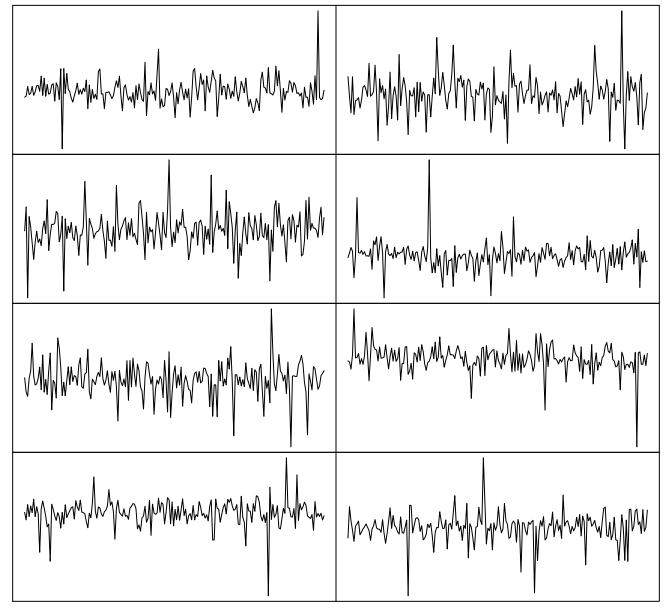
plot(stepfun(*), xlim= . , ylim = .)

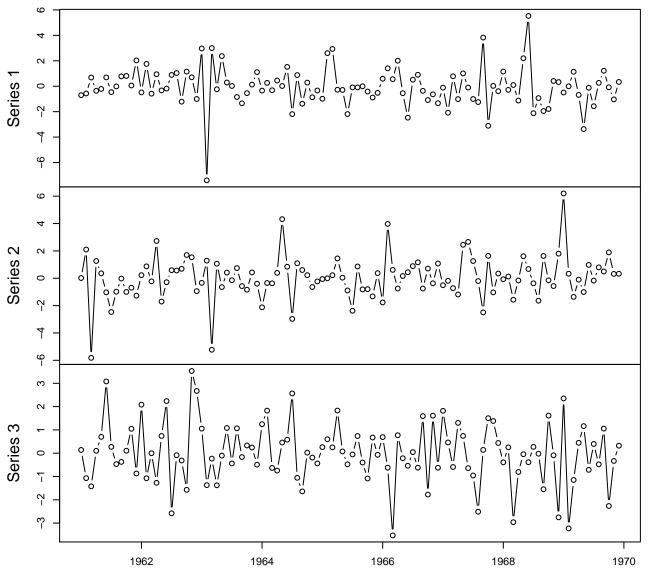




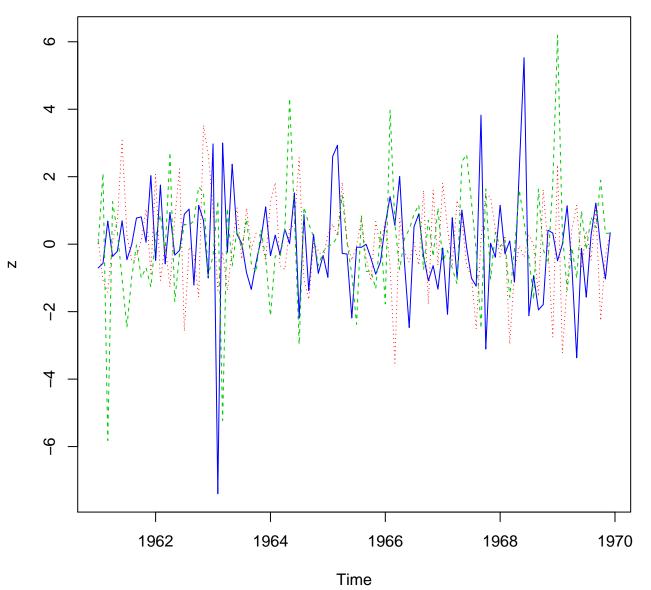


plot(ts(..), axes=FALSE, ann=FALSE, frame.plot=TRUE, mar..., oma...)

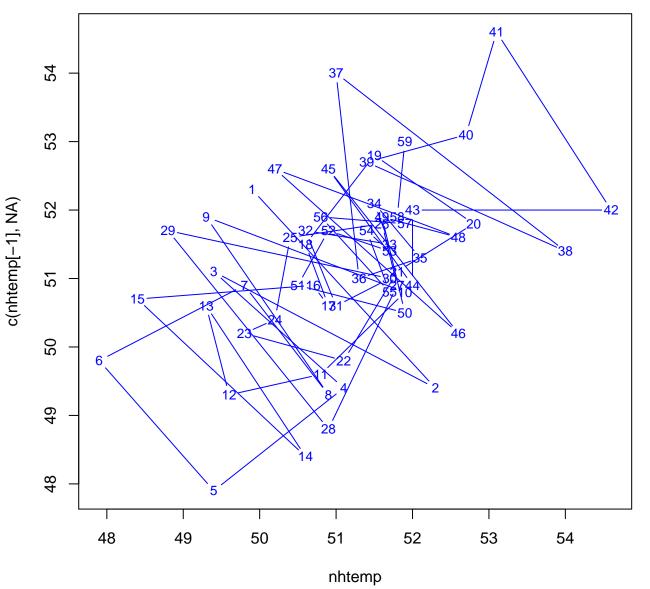


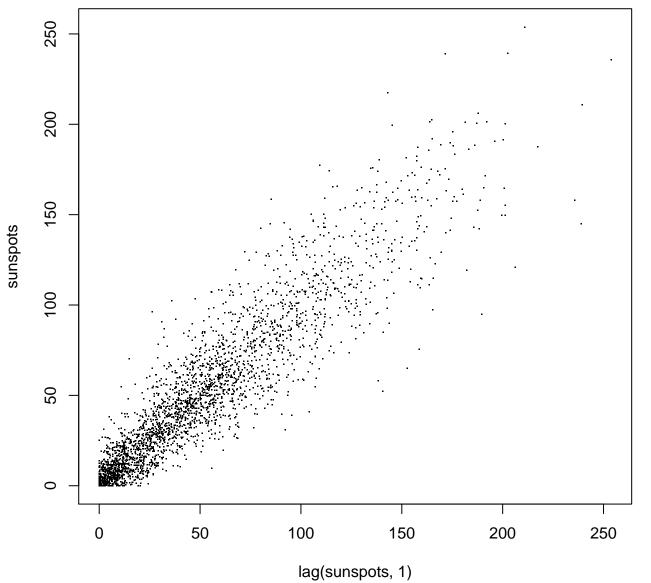


Time

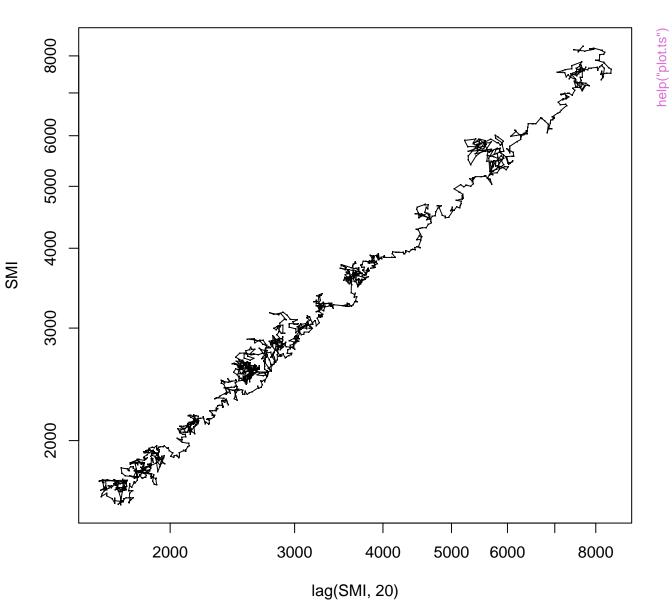


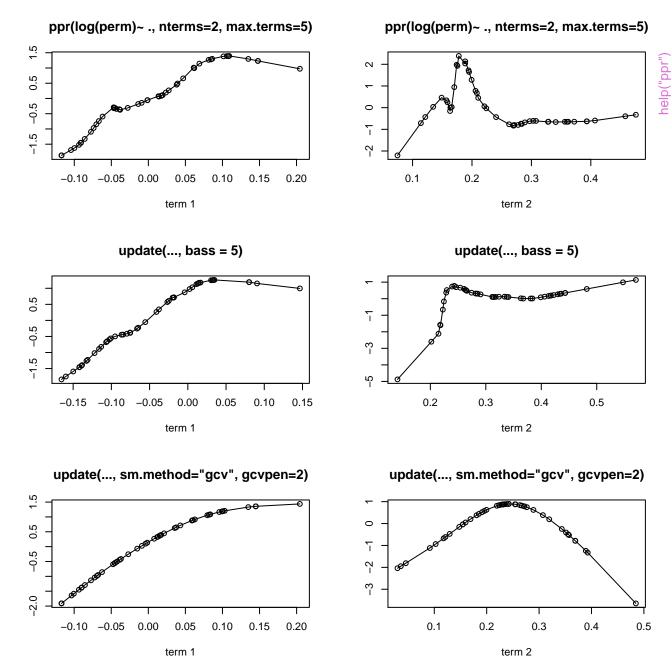
Lag plot of New Haven temperatures



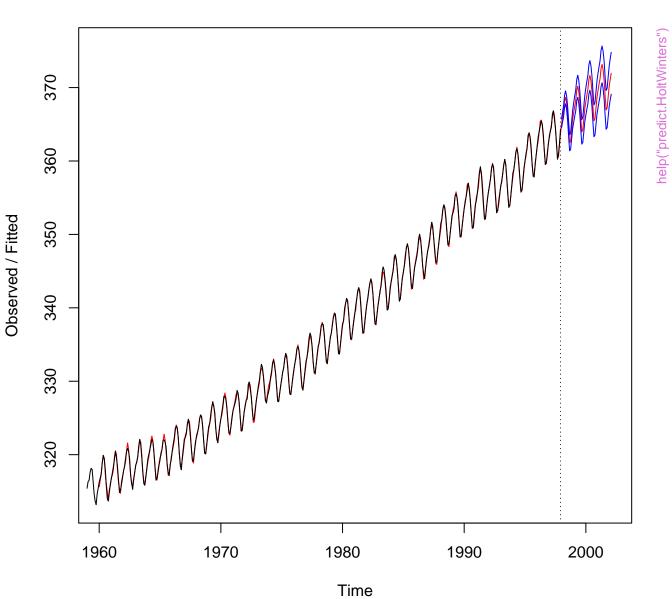


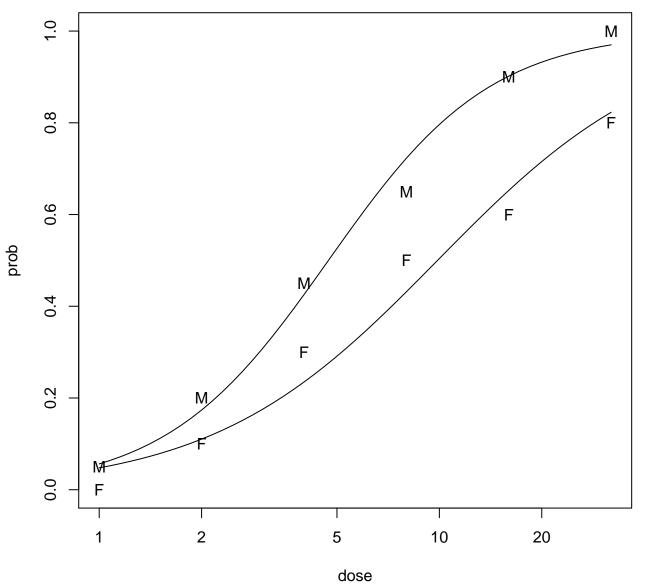
lag(SMI, 1)



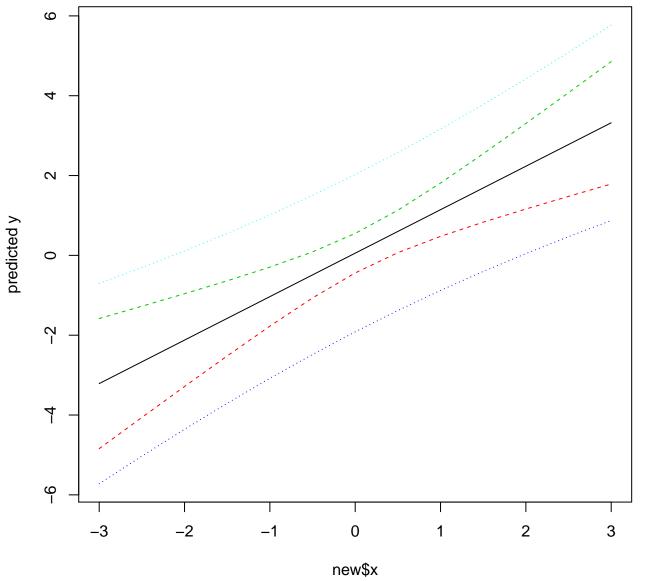


Holt-Winters filtering

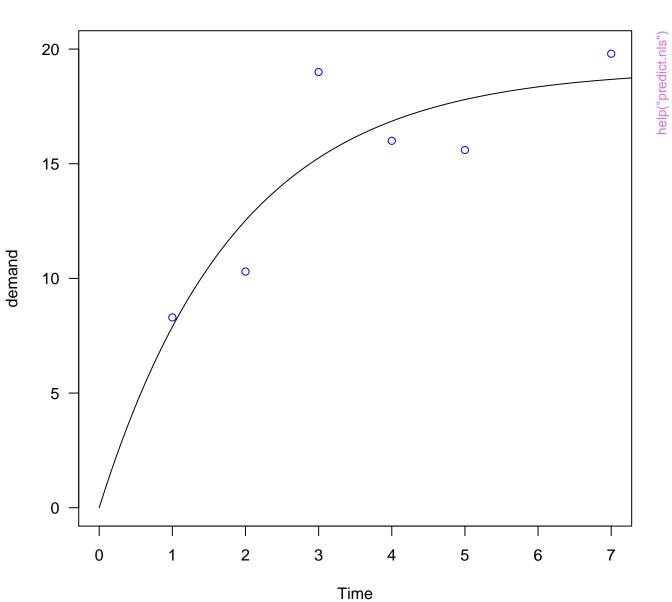


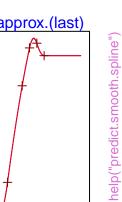


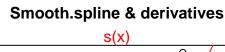


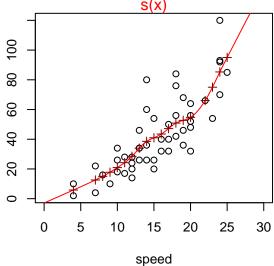


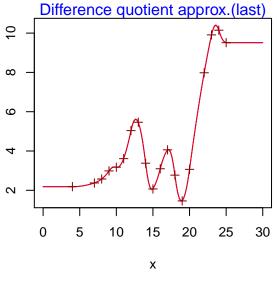
BOD data and fitted first-order curve



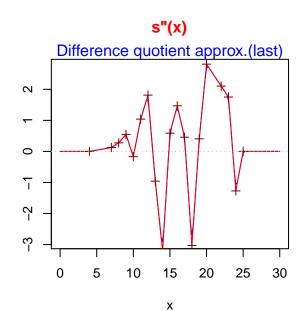


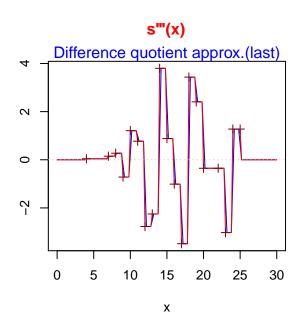




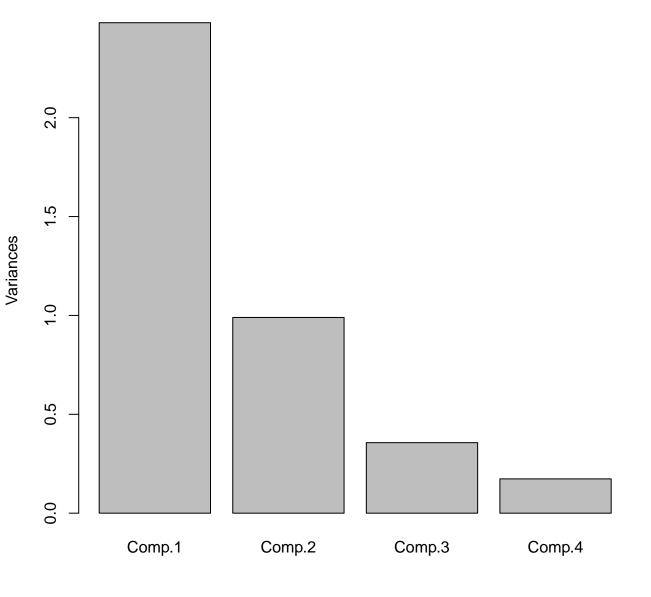


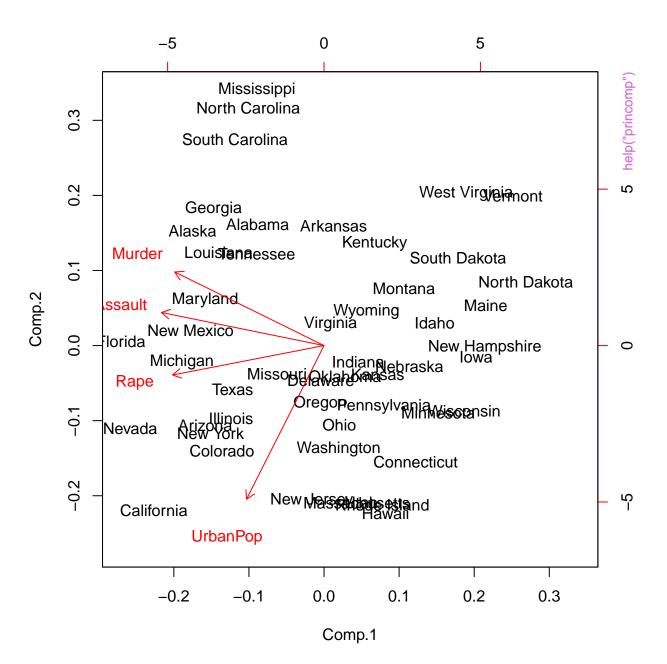
s'(x)



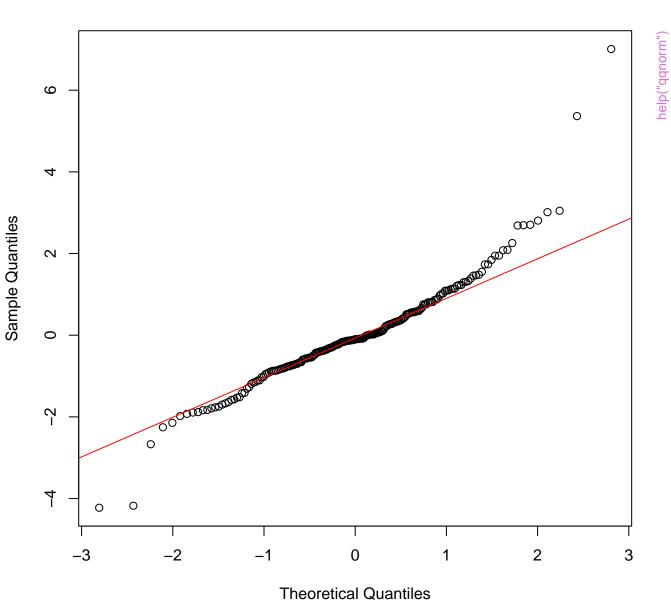


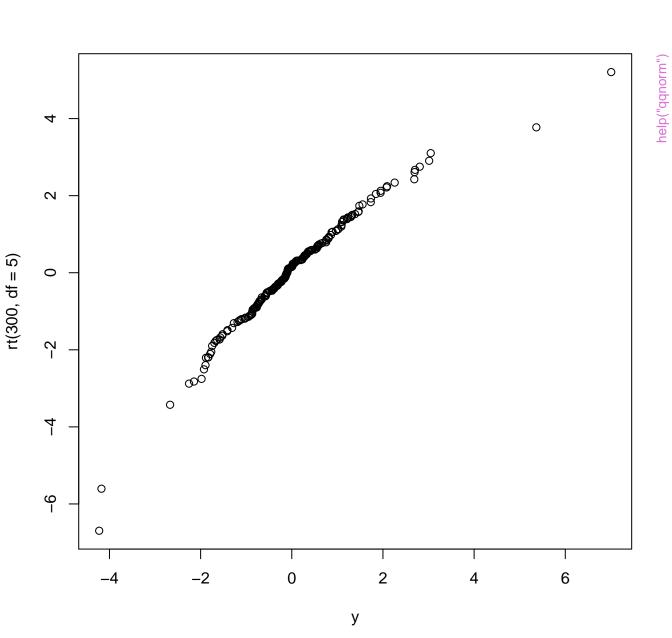


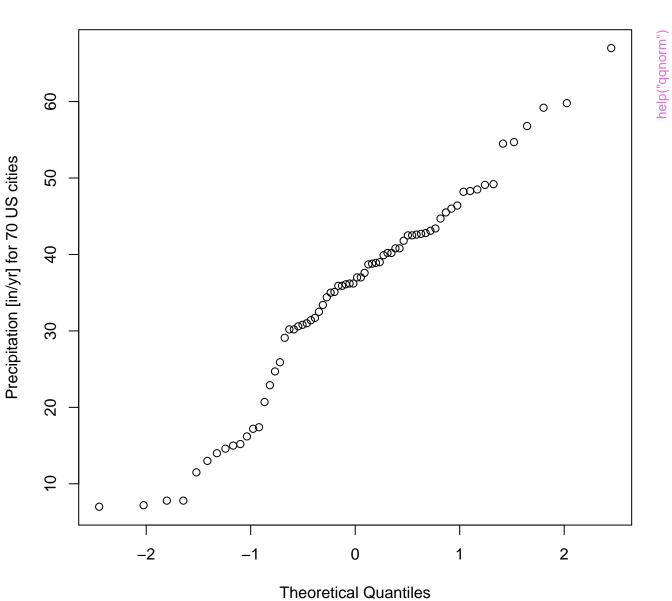


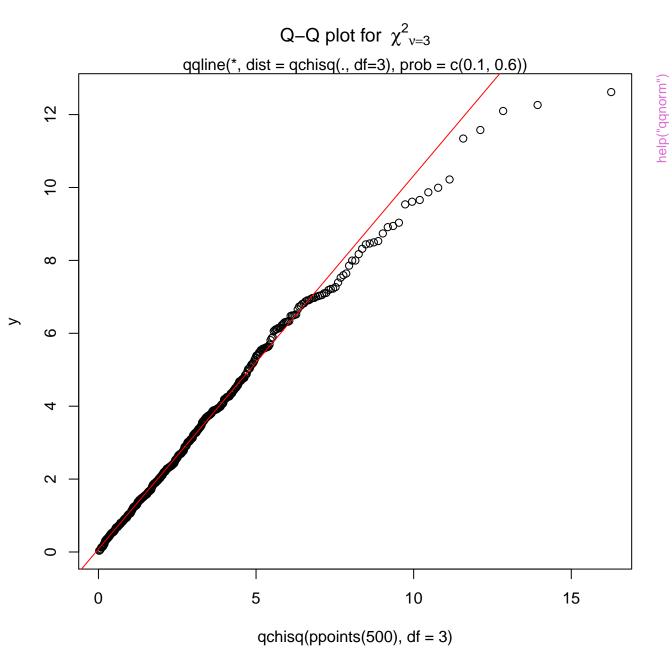


Normal Q-Q Plot

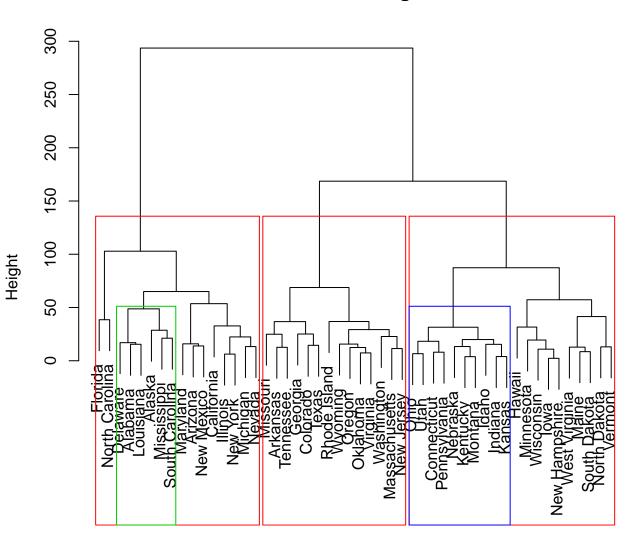






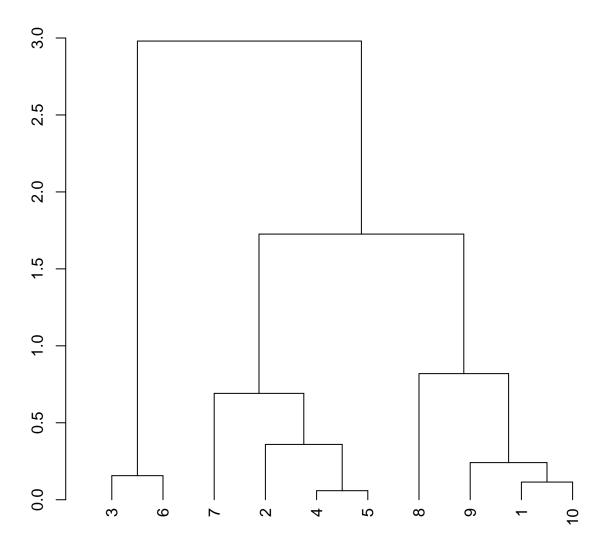


Cluster Dendrogram



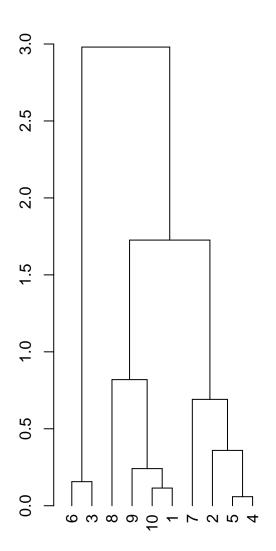
dist(USArrests) hclust (*, "complete")

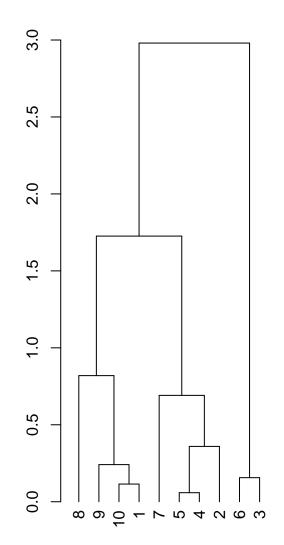
random dendrogram 'dd'



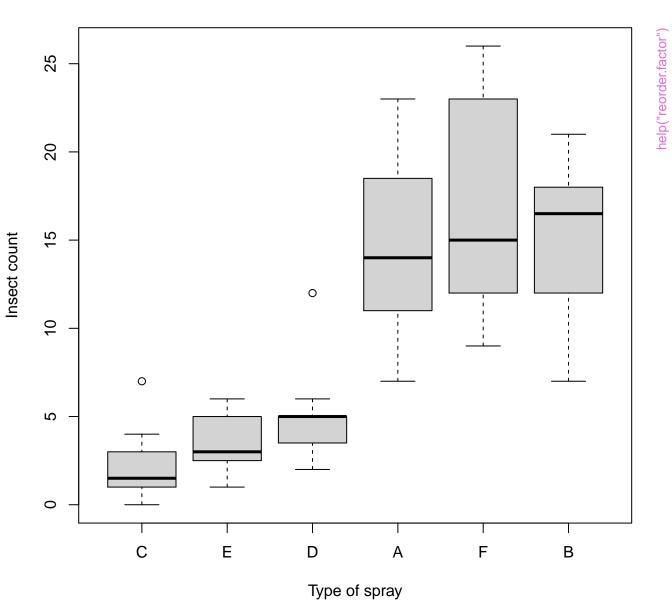
reorder(dd, 10:1)

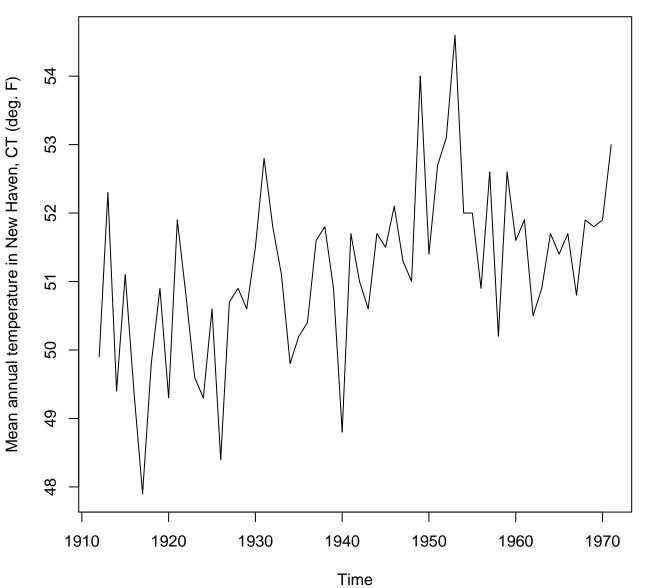
reorder(dd, 10:1, mean)





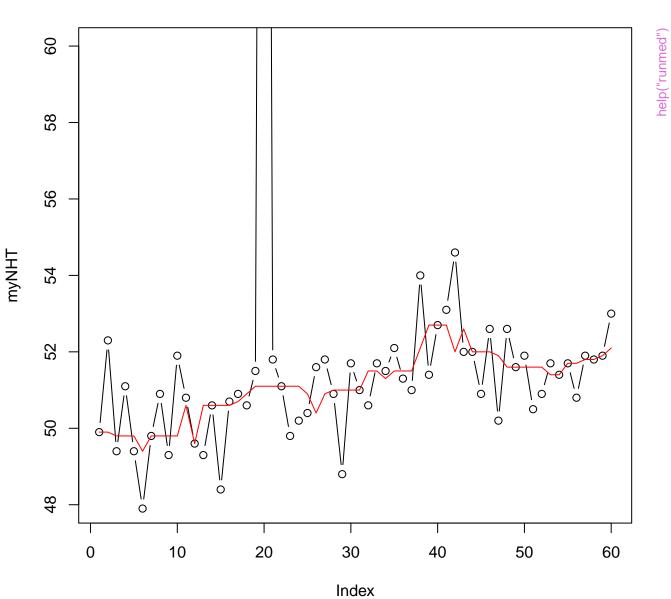
InsectSprays data



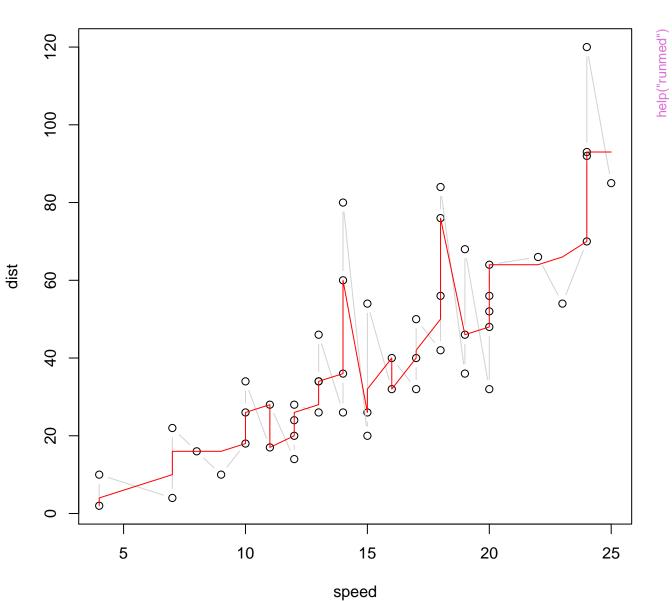


help("runmed")

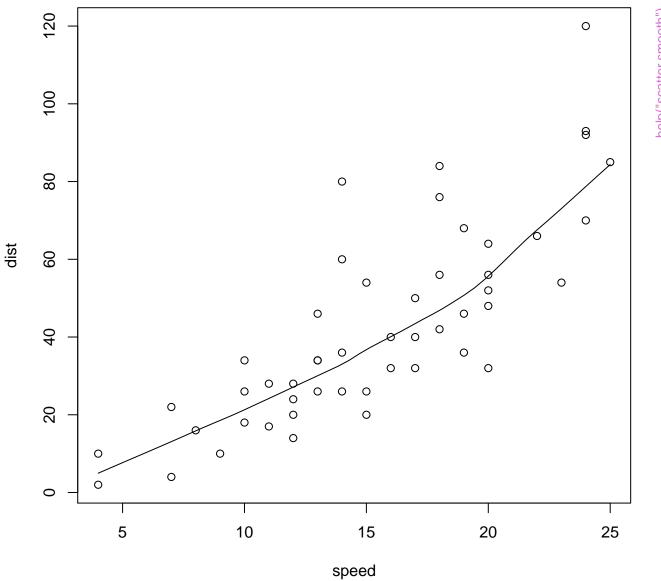
Running Medians Example



'cars' data and runmed(dist, 3)

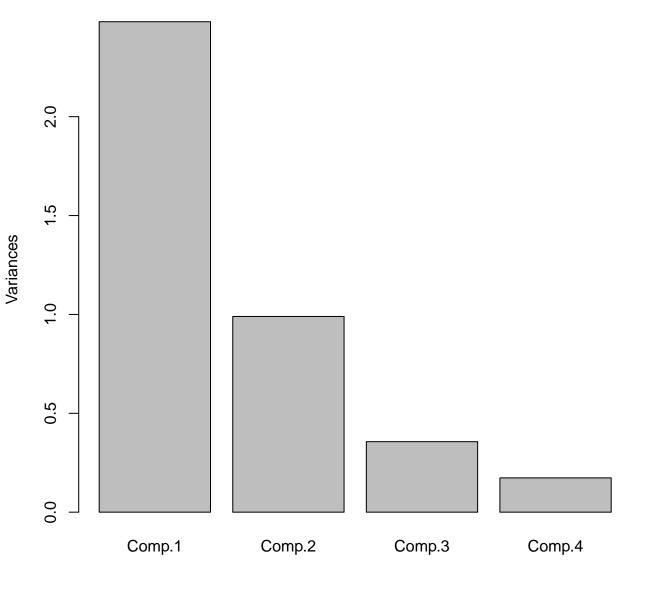


Index

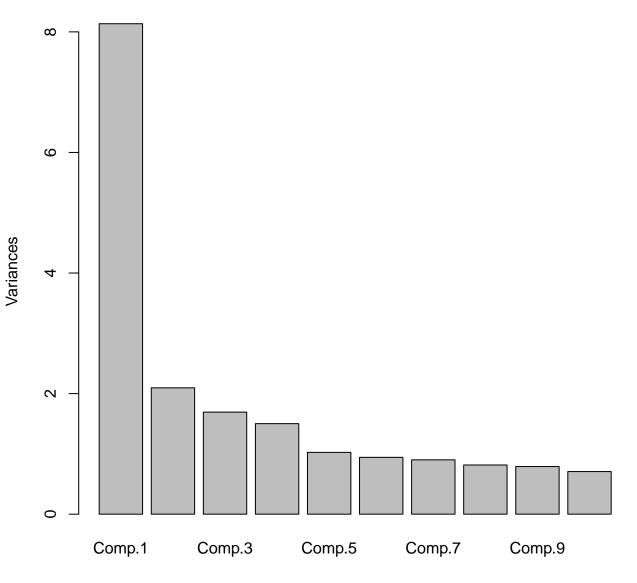


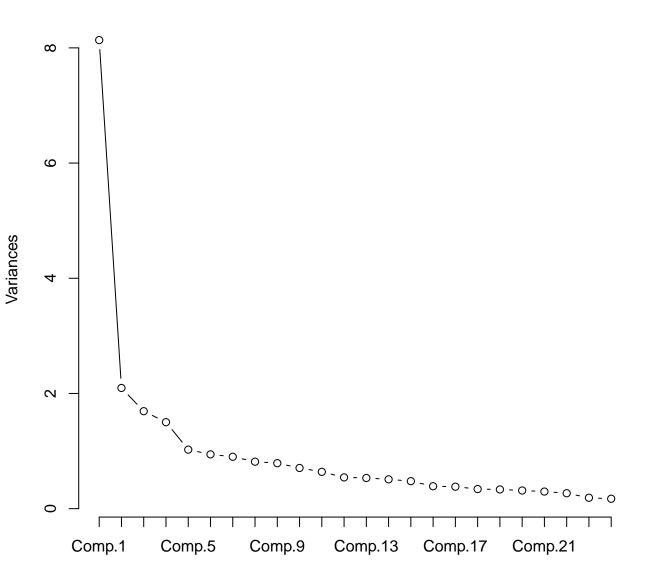
speed

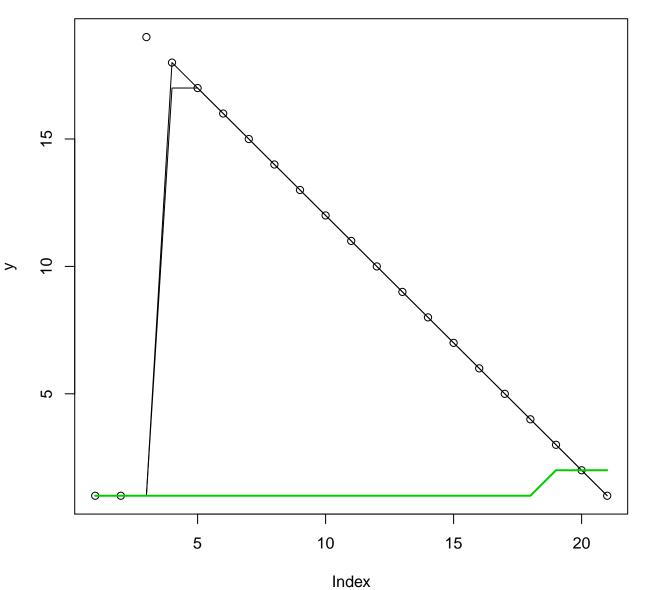


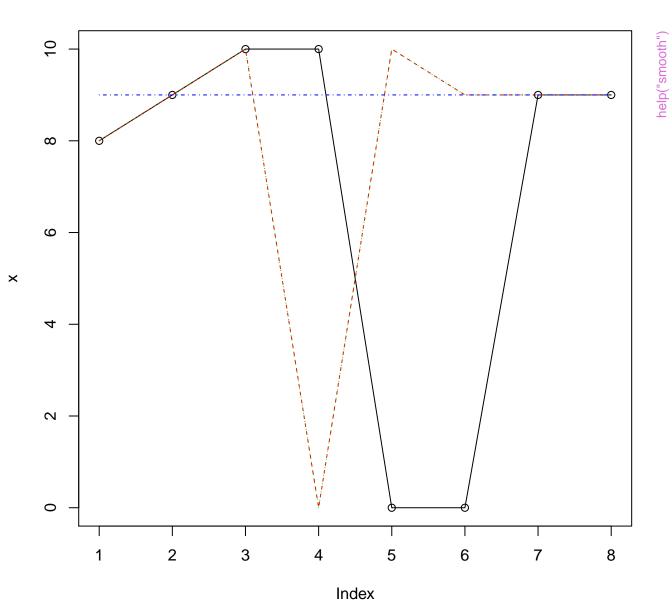




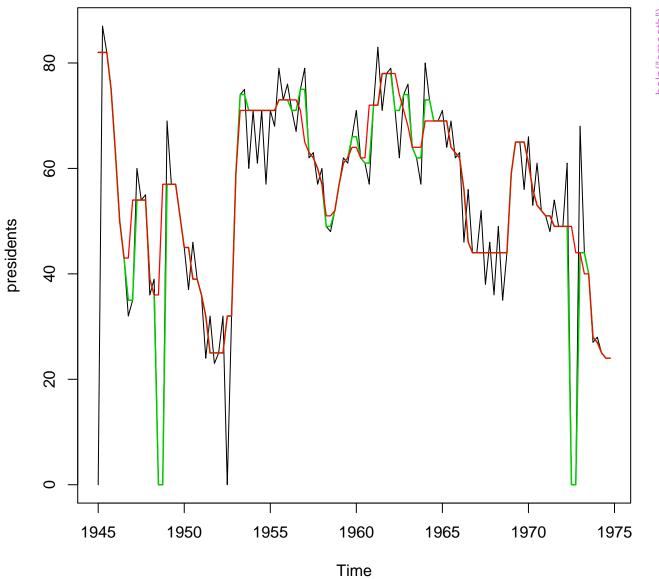


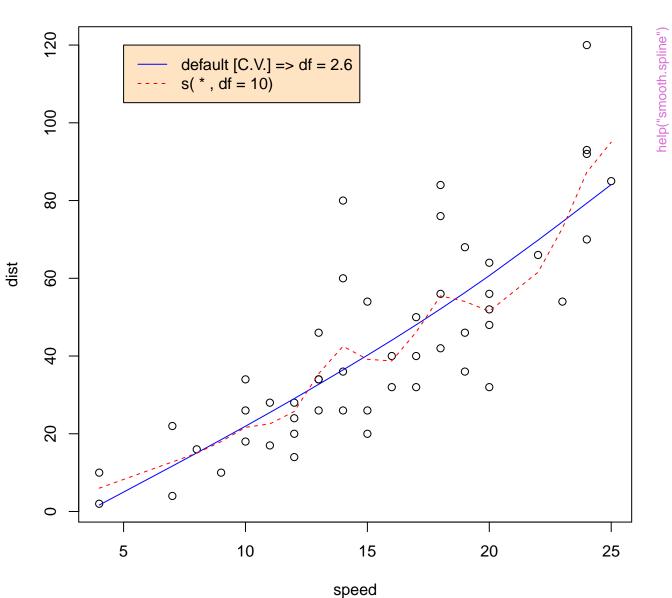


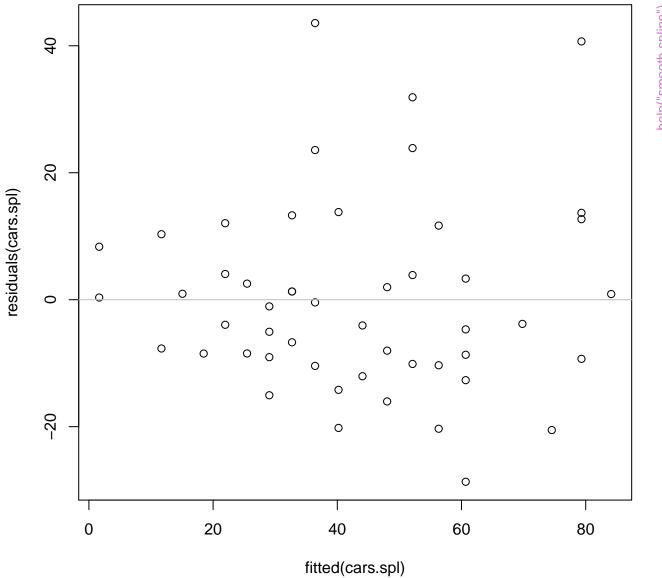


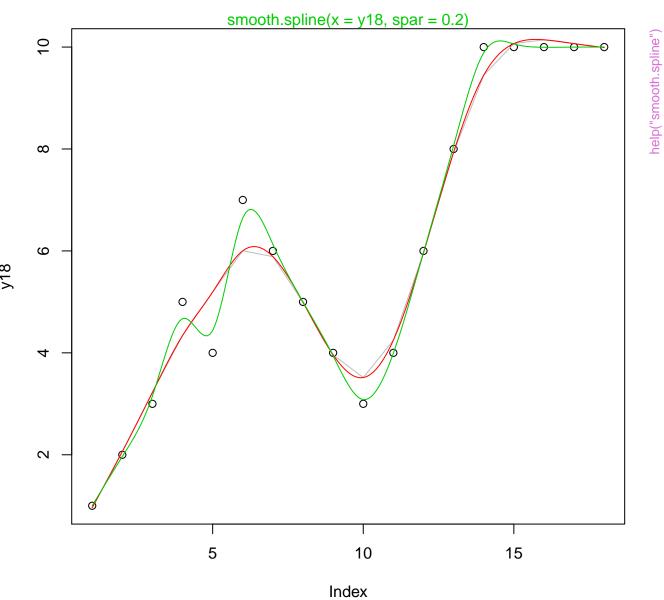


smooth(presidents0, *): 3R and default 3RS3R

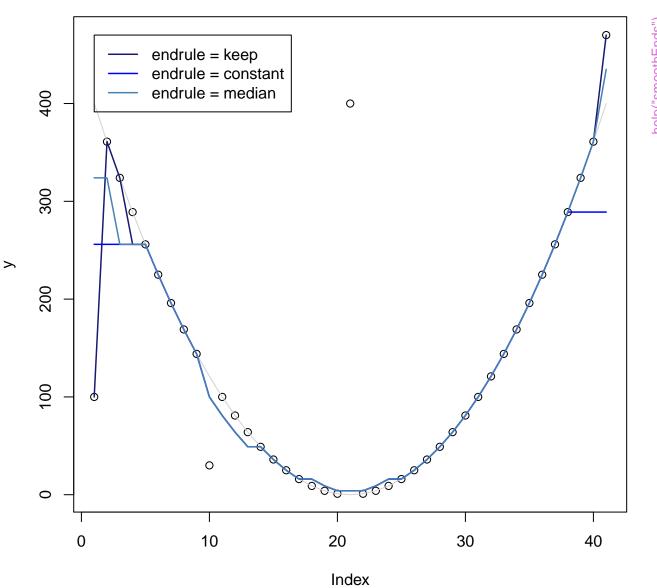




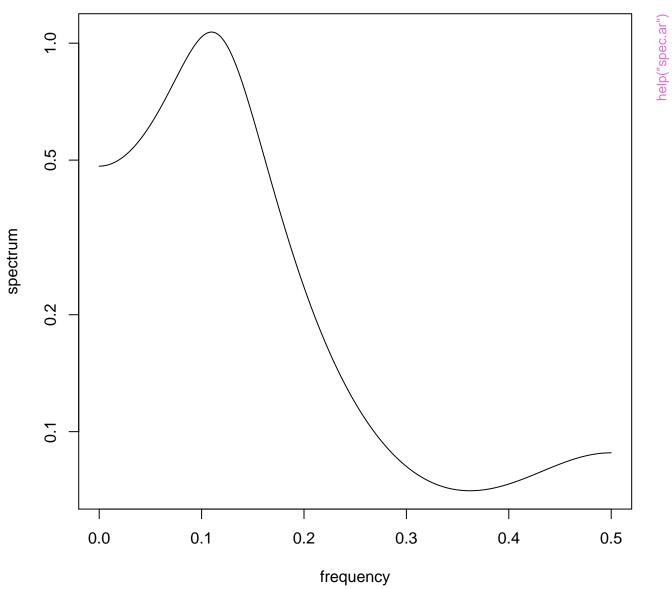




Running Medians -- runmed(*, k=7, end.rule = X)



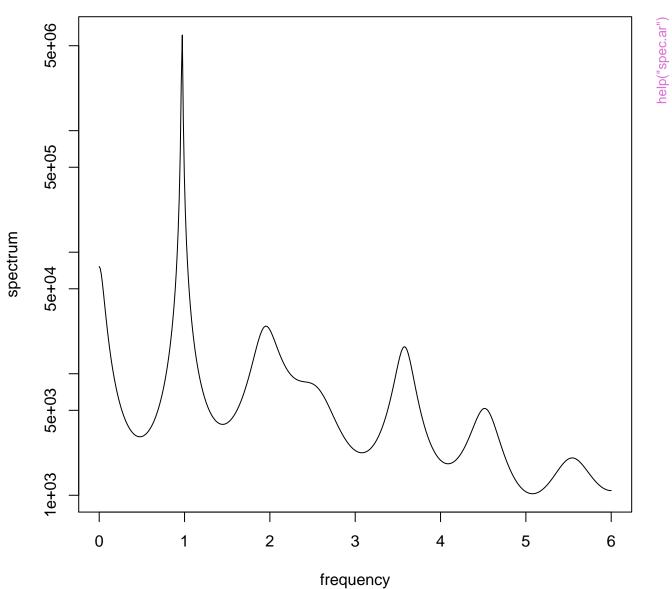
Series: Ih AR (3) spectrum



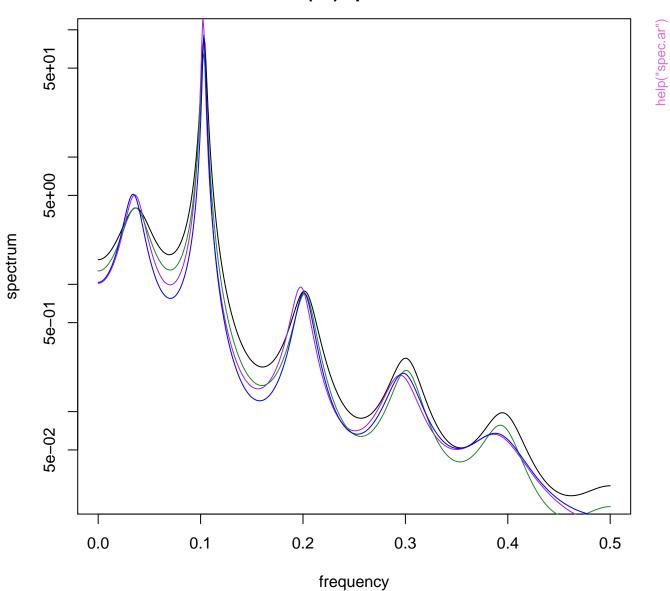
Series: Ideaths AR (10) spectrum 2e+06 help("spec.ar") 2e+05 5e+05 spectrum 2e+04 5e+04 5e+03 2e+03 5 0 2 3 4 6

frequency

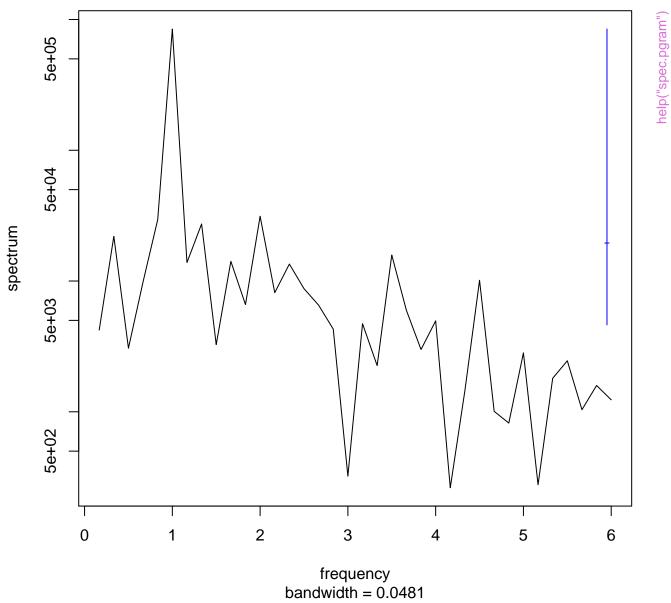
Series: Ideaths AR (13) spectrum



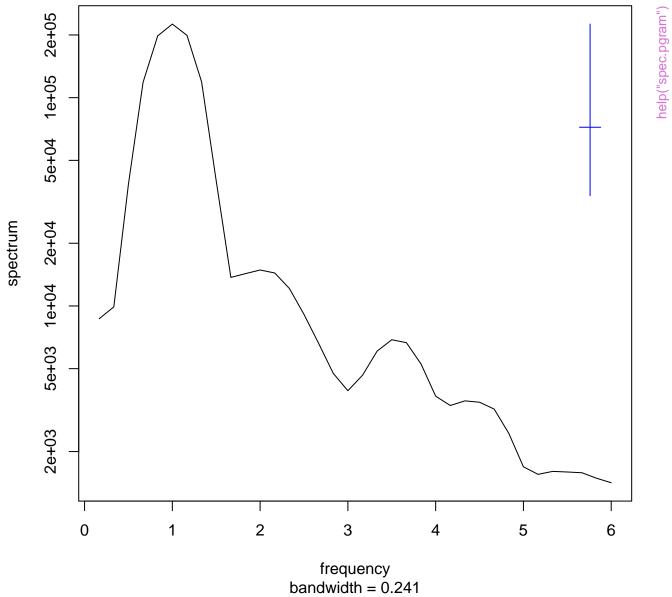
Series: log(lynx) AR (11) spectrum



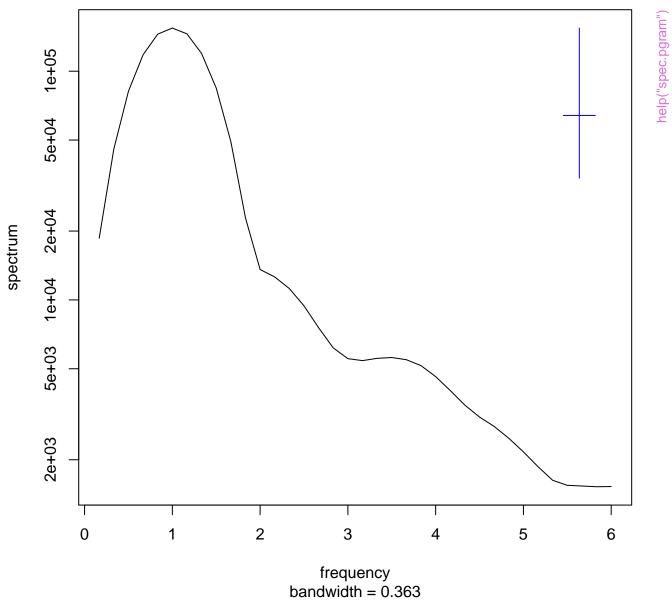
Series: x Raw Periodogram

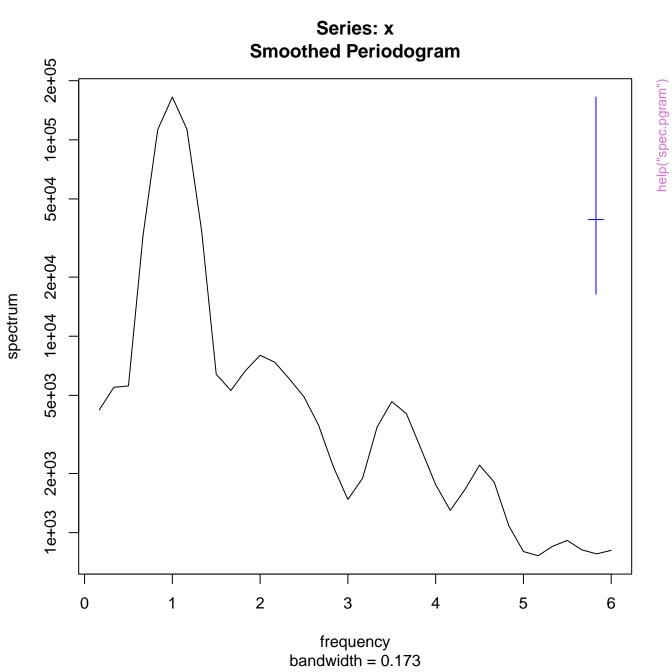


Series: x Smoothed Periodogram

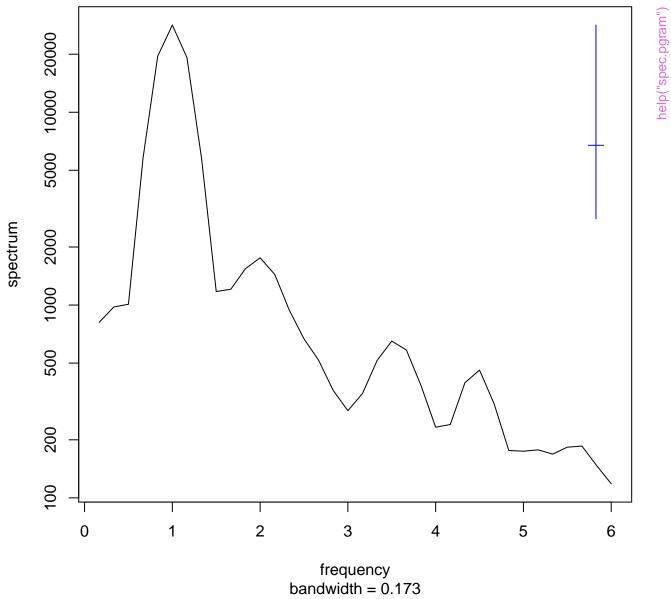


Series: x Smoothed Periodogram

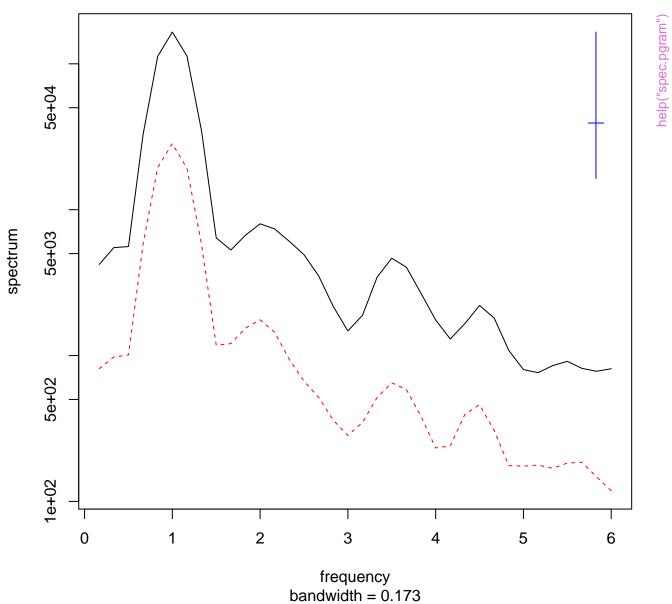




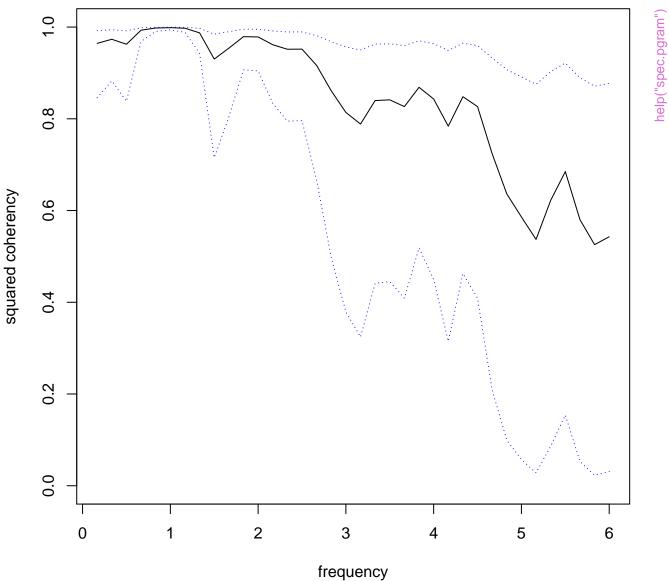
Series: x Smoothed Periodogram

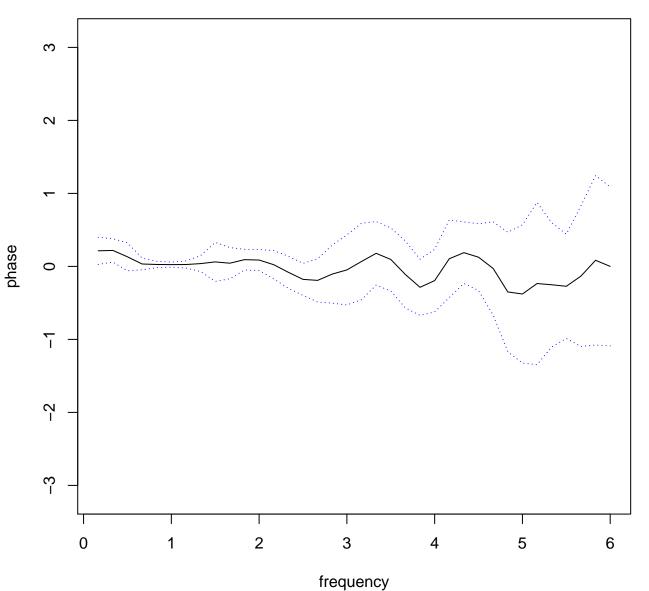


Series: ts.union(mdeaths, fdeaths)
Smoothed Periodogram

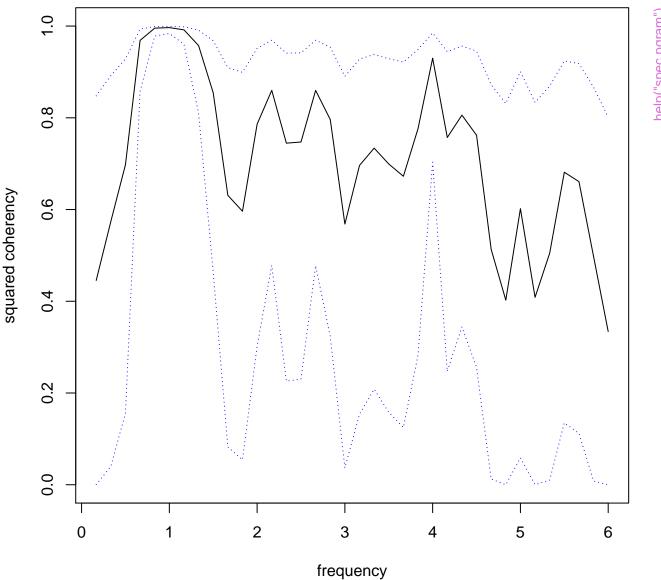


Series: ts.union(mdeaths, fdeaths) -- Squared Coherency

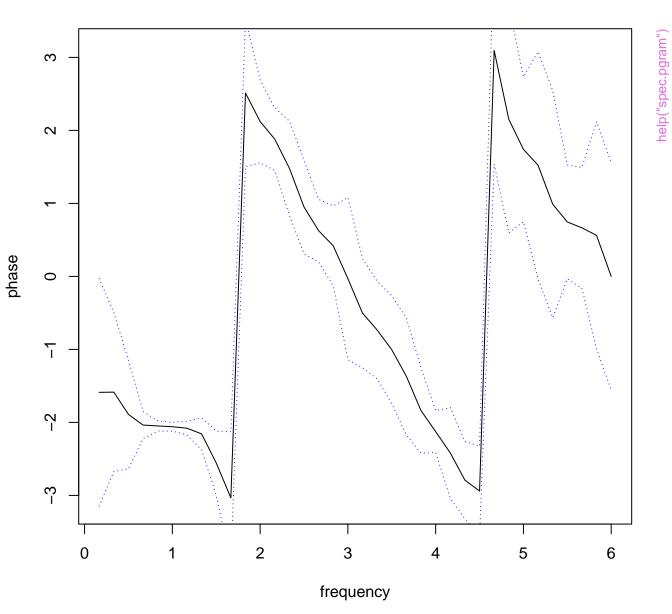




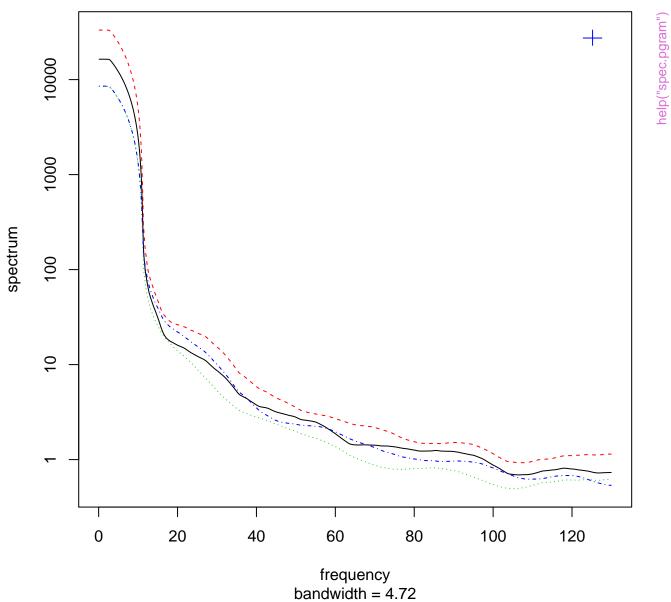




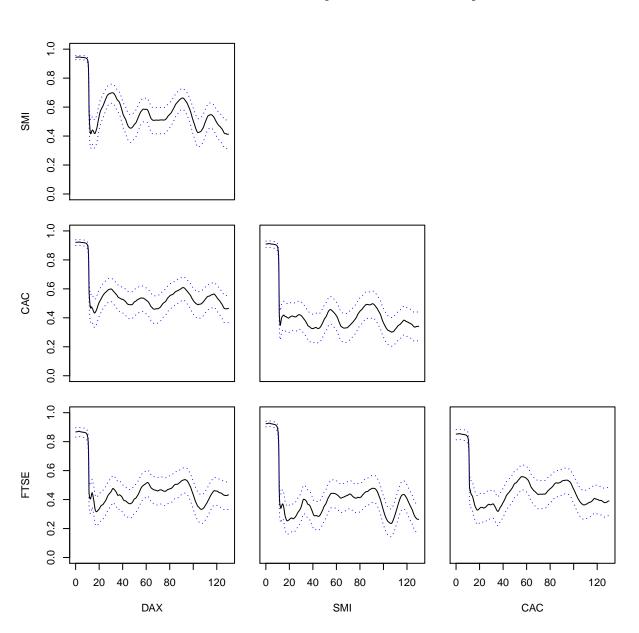
Series: ts.intersect(mdeaths, lag(fdeaths, 4)) -- Phase spectrum



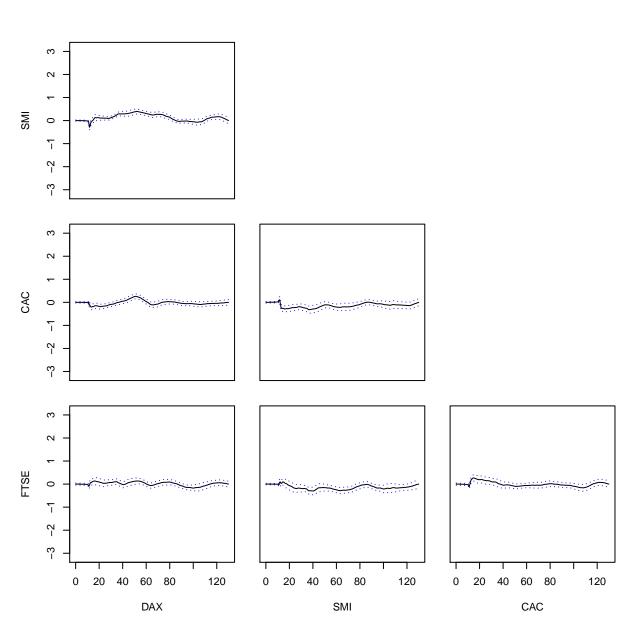
Series: x Smoothed Periodogram



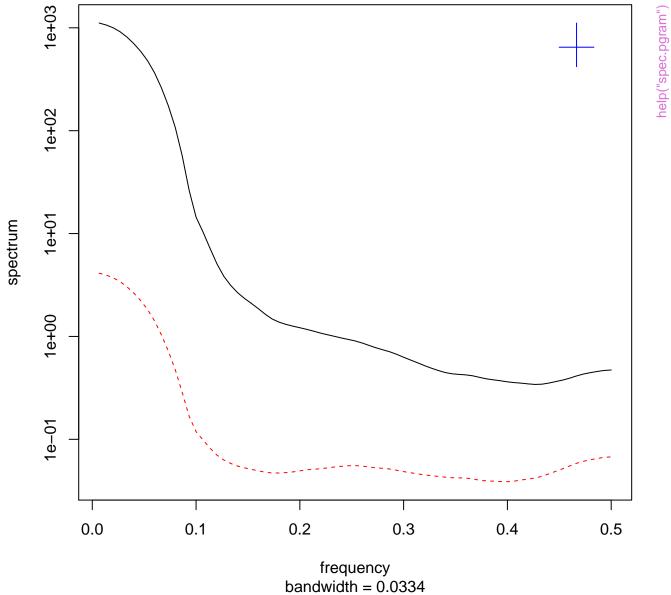
Series: x -- Squared Coherency

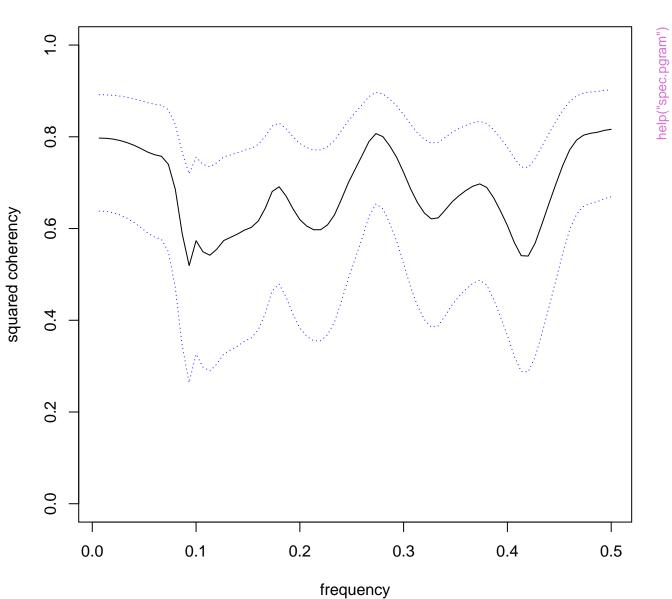


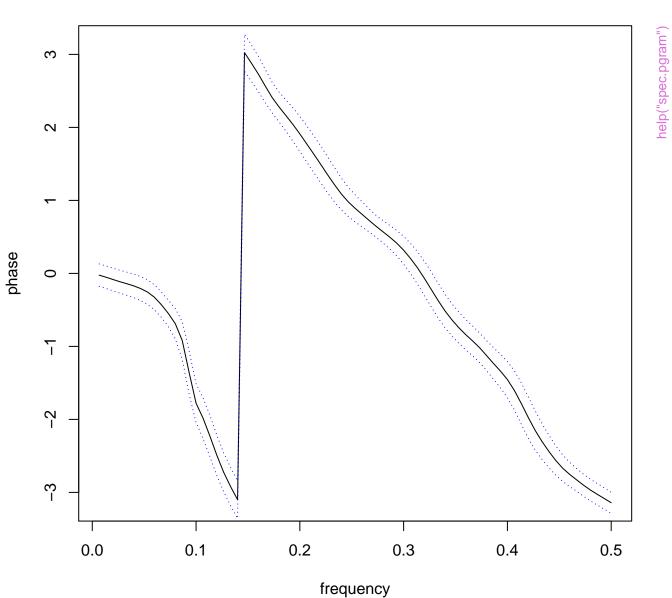
Series: x -- Phase spectrum

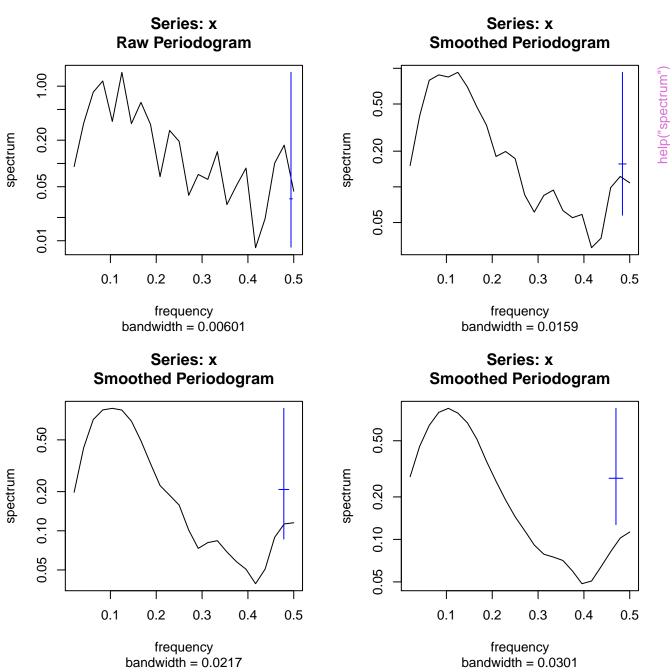


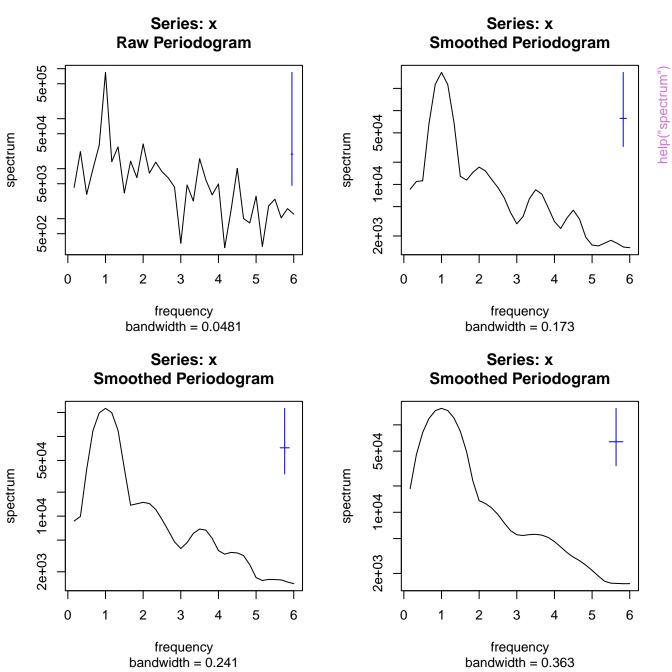
Series: x Smoothed Periodogram

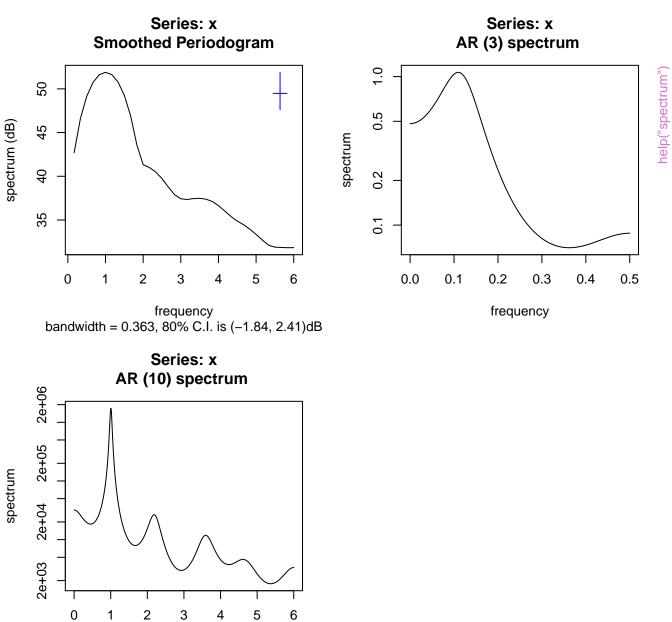






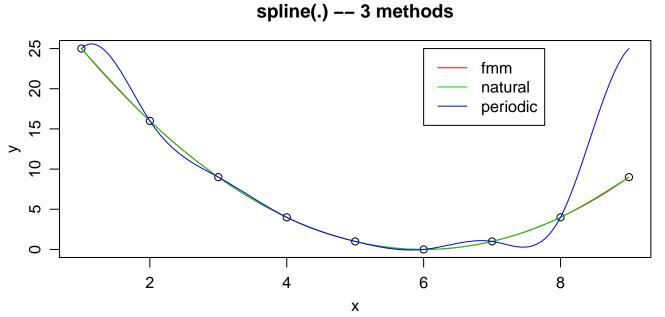


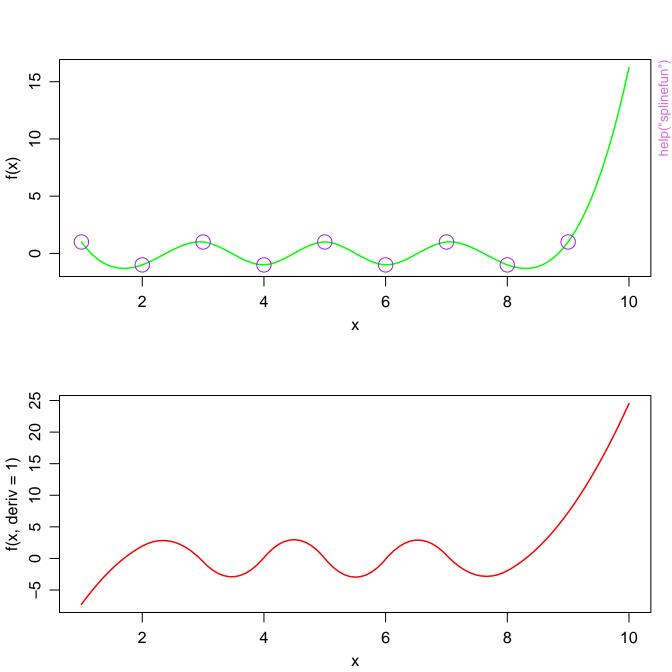


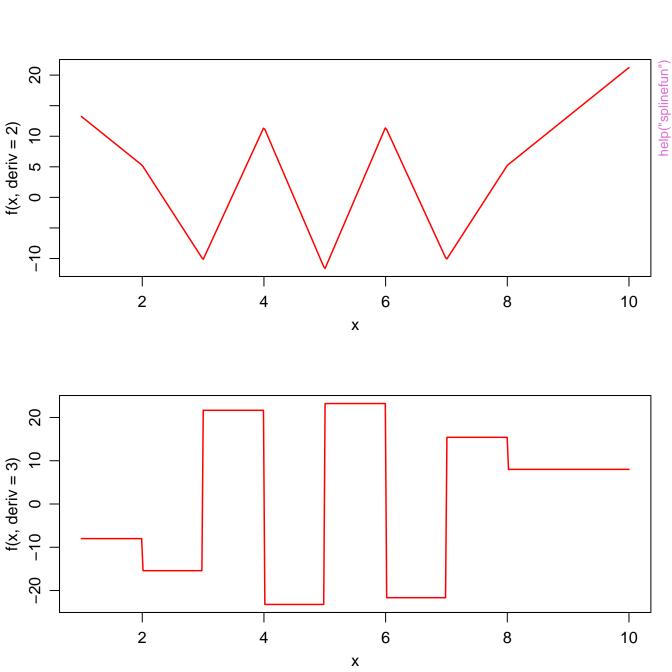


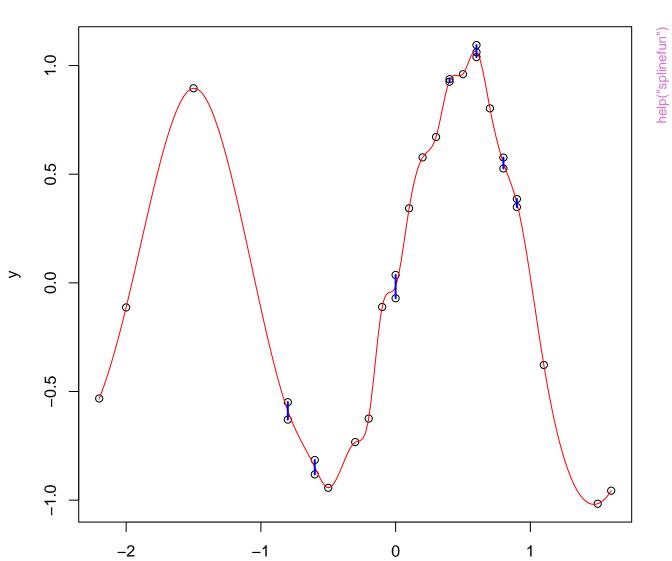
frequency

spline[fun](.) through 9 points help("splinefun" y 0.5 -0.5 2 6 8 Χ

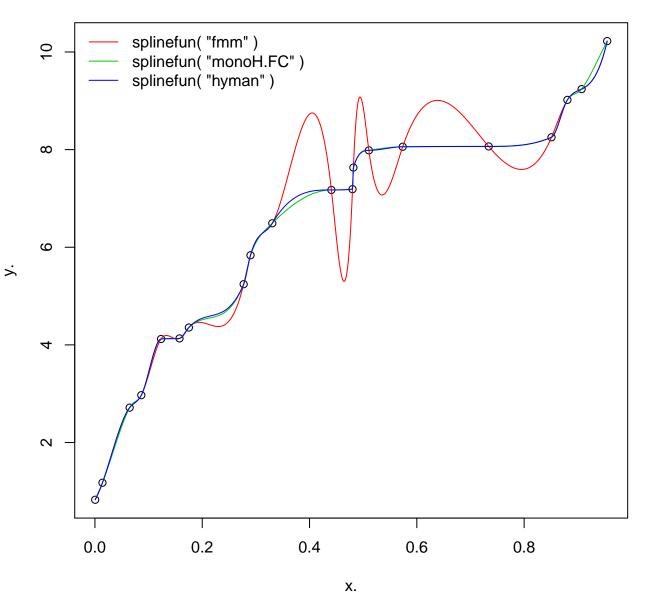


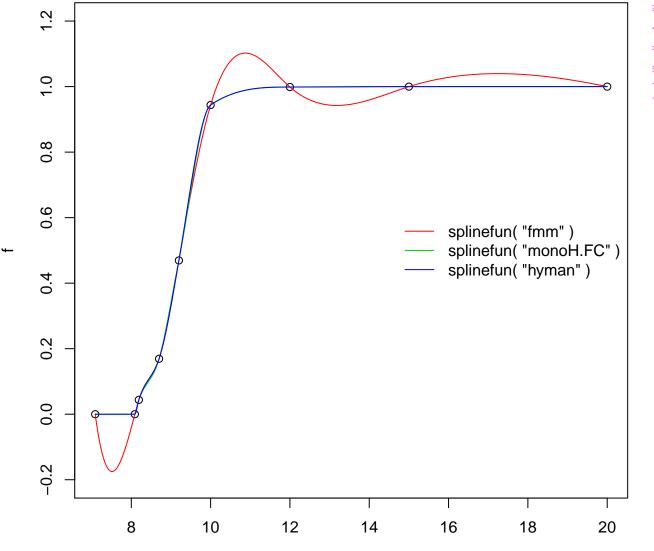






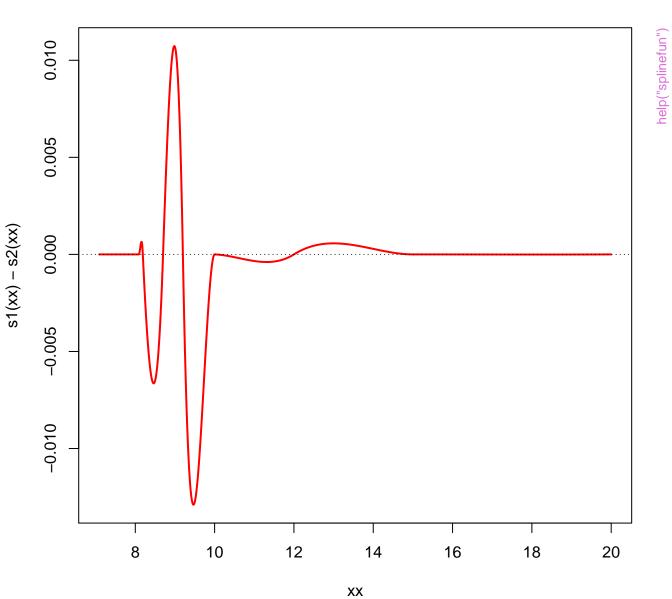
Χ



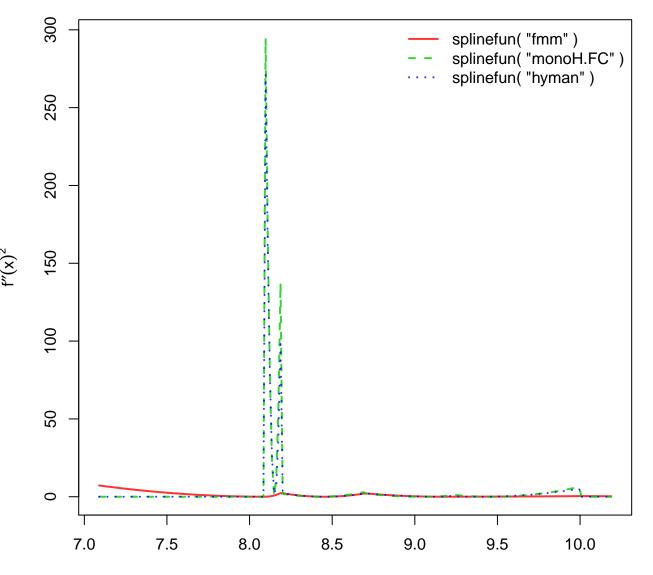


х.

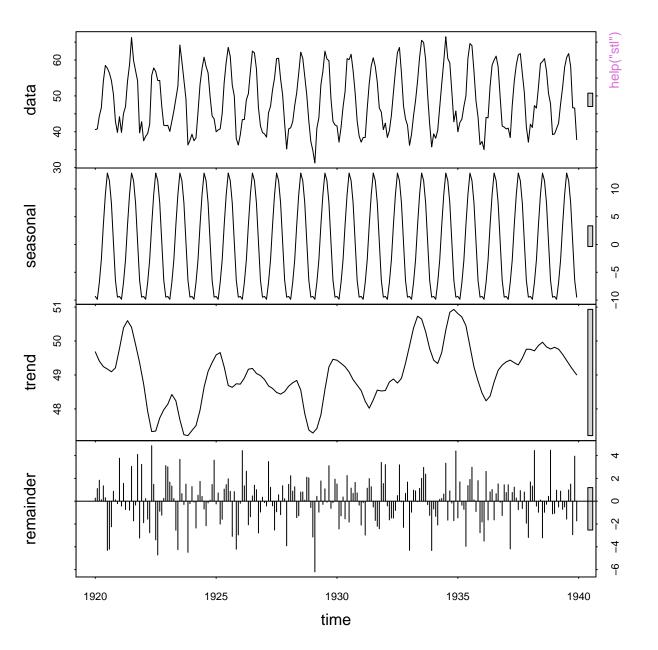
Difference monoH.FC - hyman

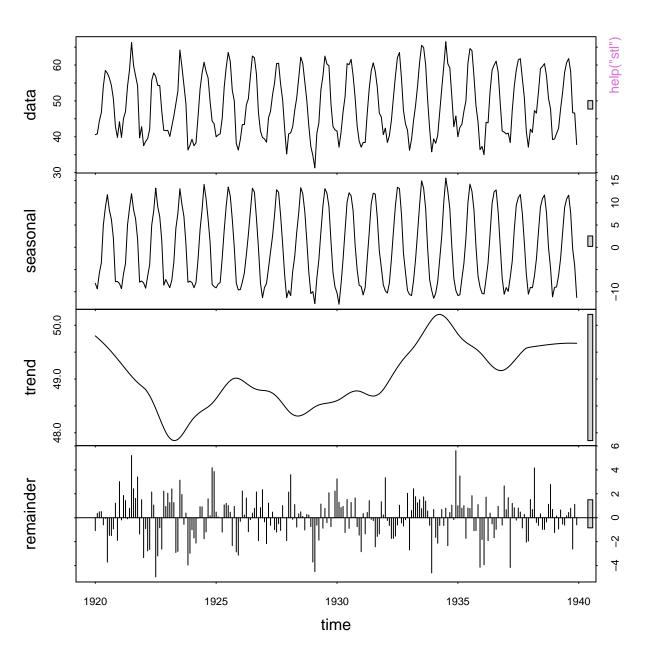


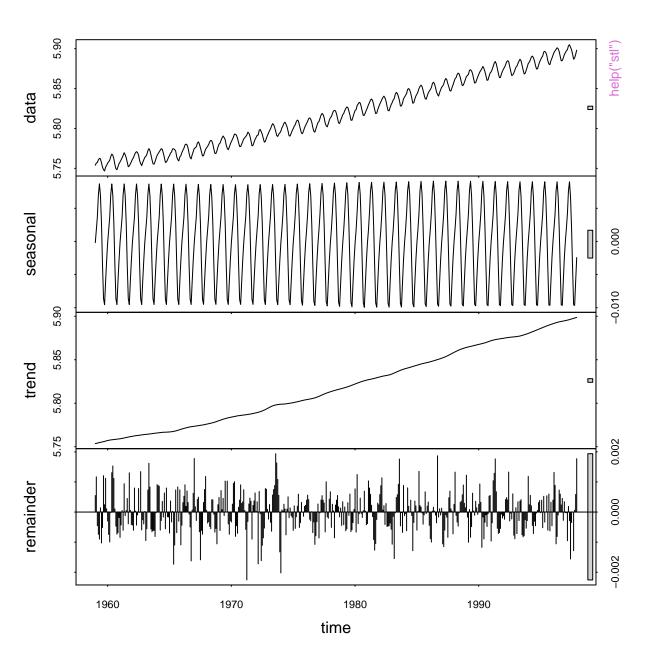
 $f''(x)^2$ for the three 'splines'

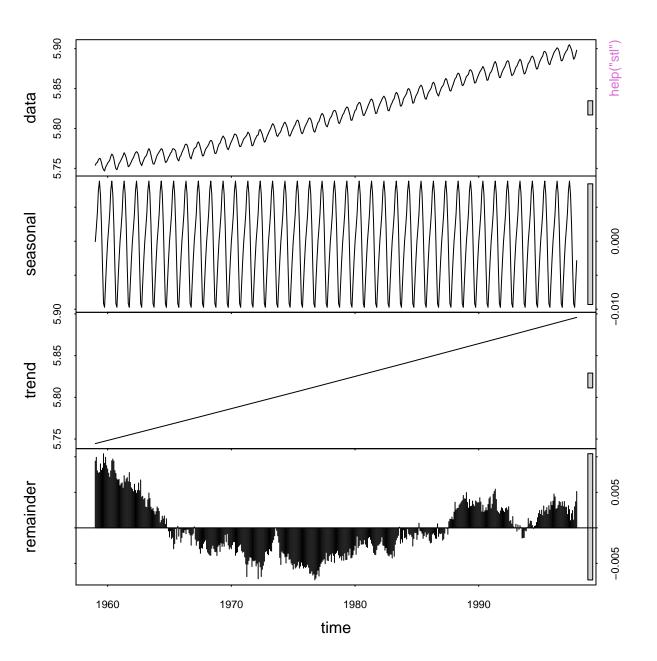


Χ

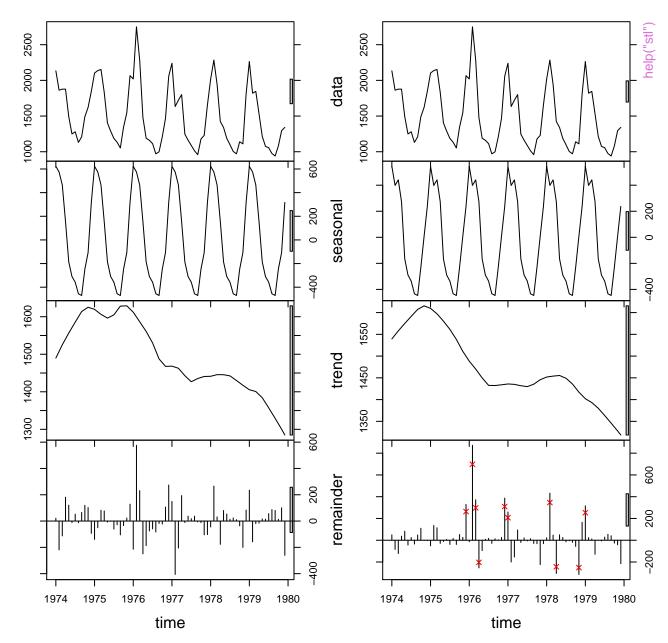




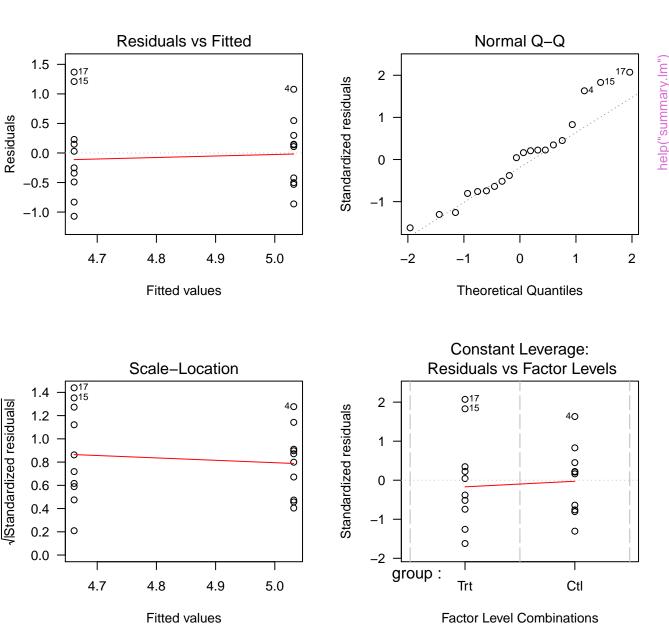


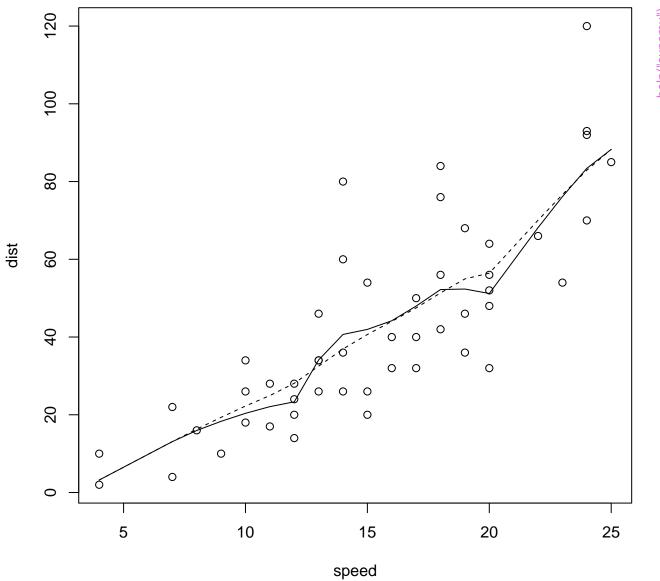


stl(mdeaths, s.w = "per", robust = FALSE / TRUE)



lm(weight ~ group)





group

termplot(glm(formula = $y \sim ns(x, 6) + z$) . termplot(glm(formula = $y \sim ns(x, 6) + z$) . help("termplot") Partial for ns(x, 6) 0.5 0.5 Partial for z -0.5 -0.5 -1.5 -1.5 0 20 40 60 80 100 Α В С D Χ Z 1.5 Partial for ns(x, 6) 0.5 0.5 Partial for z -0.5 -0.5 -1.5 -1.5

0

20

40

Χ

60

80

100

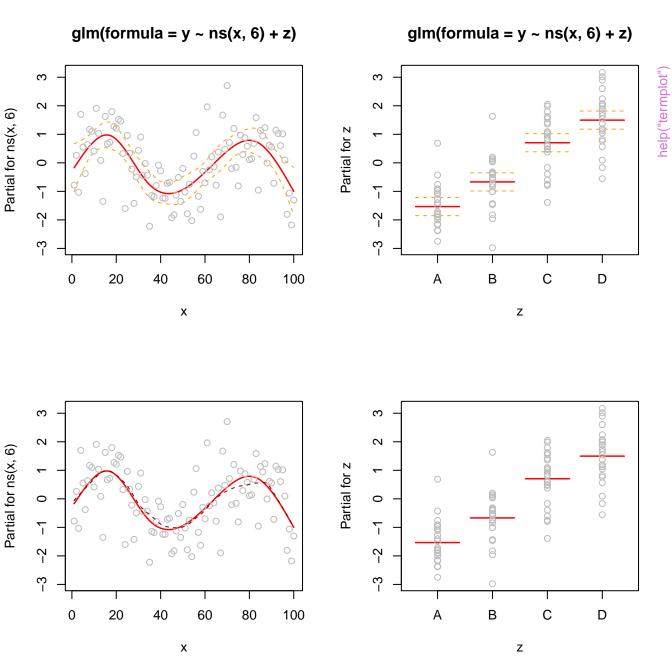
Α

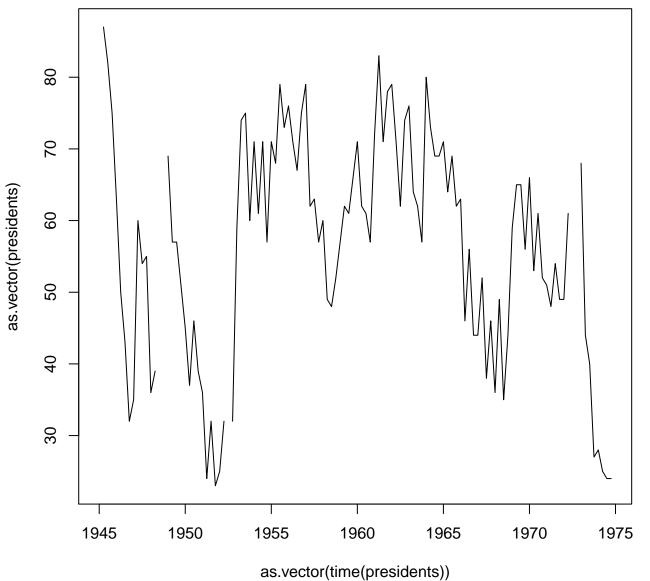
В

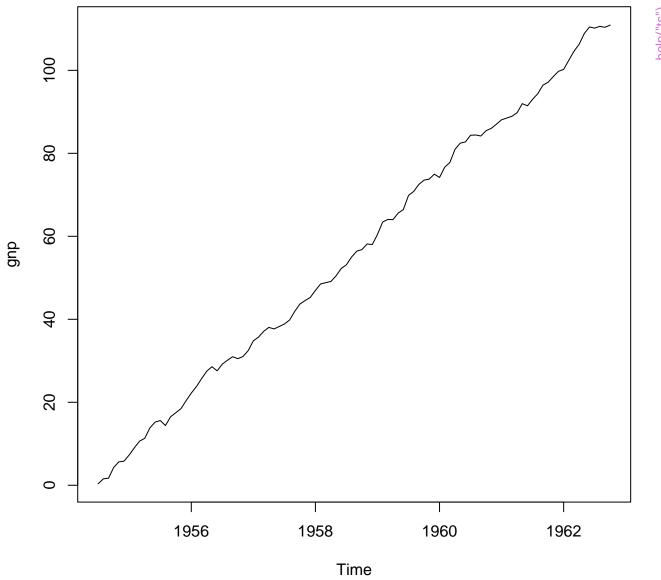
Z

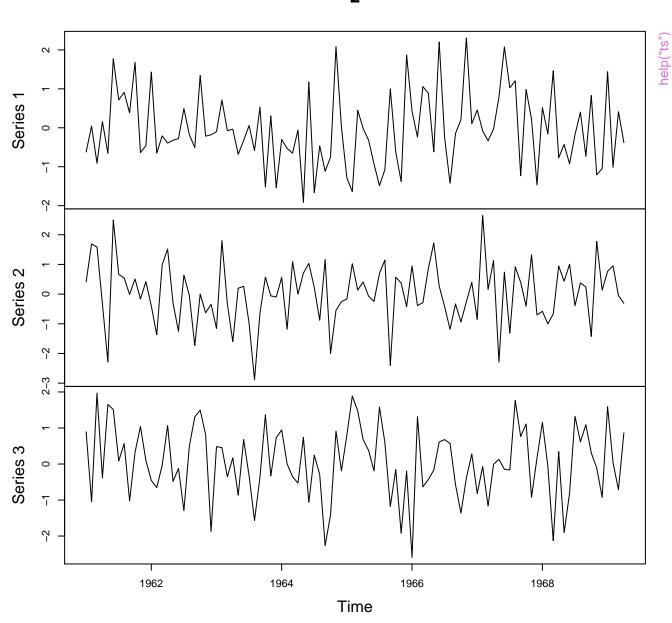
С

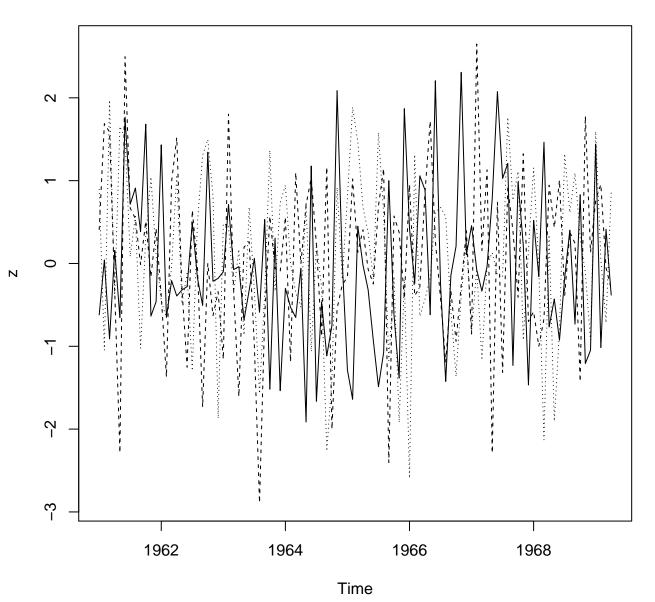
D



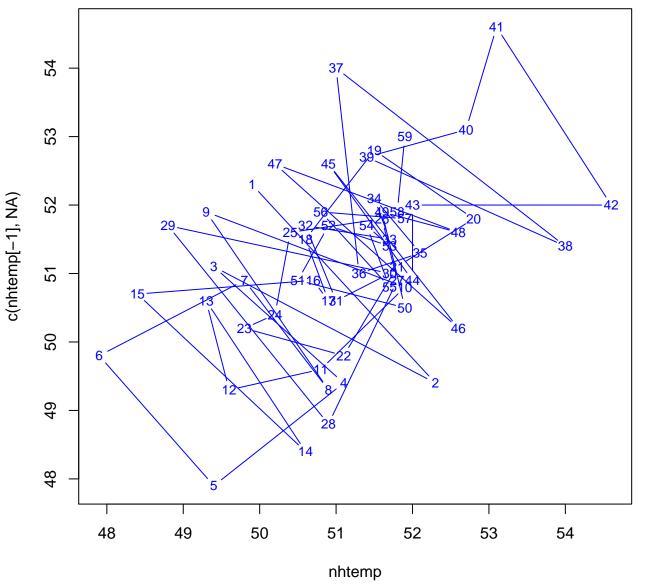


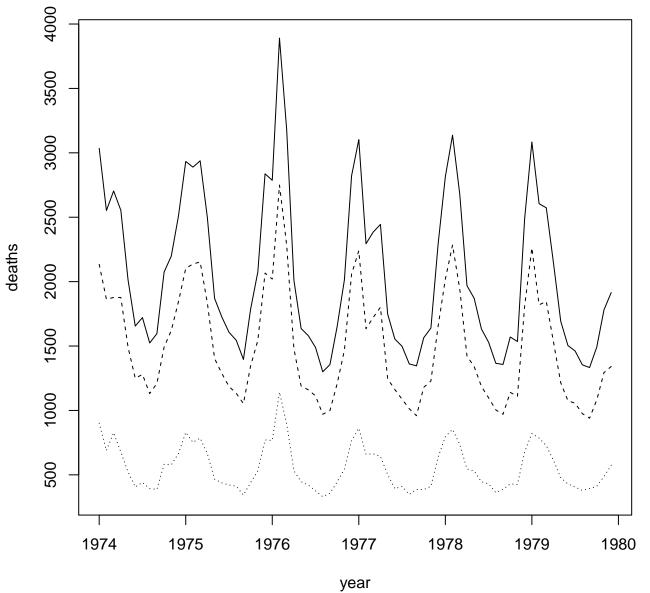






Lag plot of New Haven temperatures





lm(weight ~ group)

