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SMOOTHING OF SCATTERPLOTS*

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ABSTRACT

A variable span scatterplot smoother based on local linear fits is described. Local cross-validation is used to estimate the optimal span as a function of obscissa value. A rejection rule is suggested to make the smoother resistant against outliers. Computationally efficient algorithms making use of updating formulas and corresponding FORTRAN subroutines are presented.

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1. Introduction

A smoother is a procedure that operates on bivariate data $(x_1,y_1)...(x_n,y_n)$ and produces a decomposition

$$y_i = s(x_i) + r_i, \quad i=1...n.$$
 (1)

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Here s is a smooth function, often simply called the smooth, and the riar residuals. It is possible to formally define duat constitutes a smooth function, and to define measures of smoot less, but for our purposes an intuitive notion will be sufficient. Smoothers are used to summarize the association between the predictor variable X and the response Y. It was pointed out by Cleveland (***) and is a commonly held belief, that when looking at a scatterplot the eye is distracted by the extreme points in the point cloud, i.e., $\frac{1}{2}$ a fuzzy background, and tends to miss structure in the bulk of the data. Augmentation of the plot by a smooth is a possible remedy. More formally, smoothers can be regarded as curve estimators; one assumes that the response was generated by adding random noise to a smooth function:

$$y_i = f(x_i) + \epsilon_i \tag{2}$$

and considers the smooth s as an estimate for f.

Recently scatterplot smoothers have found a new use in multiple nonparametric regression (Friedman and Stuetzle, 1981). Let $(x_1,y_1)...,(x_n,y_n)$ denote the observations; x_i here is a vector in RP, not just a single number. Assume as above that $y_i = f(x_i) + \epsilon_i$, i = 1...n. Projection pursuit regression constructs an estimate m for f of the form

$$m(x) = \sum_{j=1}^{M} s_j(a_j, x),$$

-2-

where the α_i and suitably chosen unit vectors in RP. For given α_i , s_i (essentially) is found by smoothing the scatterplot of the residuals

$$i-1$$

 $r_j^{i-1} = y_j - \sum_{k=1}^{\infty} s_k(\alpha_k, x_j)$ versus α_1, x_j . The smoother described in this

report is, up to minor modifications, the one used in the current projection pursuit regrossion procedure,

2. Basic Concepts

According to the definition above, any procedure that passes a smooth curve through a scatterplot, for example a procedure that fits a least squares straight line, would be called a smoother. This is not quite what we have in mind. Assume the data are generated according to (2). We are interested in procedures that can approximate f arbitrarily closely, given a dense enough sample, without any conditions on f apart from f being smooth. Such procedures can be based on local averaging. Take $s(x_i)$ to be the average of the responses for those observations with predictor values in a neighborhood N of x_i :

$$s(x_j) = ave(y_j | x_j \in I).$$
 (3)

Here "ave" can stand for the arithmetic mean, the median, or more complicated ways of averaging to be discussed below. A critical parameter to be chosen is the SPAN, the size of the neighborhood over which averaging takes place. It controls the smoothness of s. The bigger the span, the smother s will be. To obtain consistency, i.e., to make sure that s gets arbitrarily close to f as the sampling rate increases one must shrink the diameter of the neighborhood in such a way that the number of observations in the neighborhood still grows to

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infinity. Shrinking the neighborhood makes the systematic or bias component in the estimation error diminish, while increasing the neighborhood sample size guarantees that the variance component of the error goes to zero as well.

An alternative method for nonparametric curve estimation is based on series expansions: make an ansatz for s of the form

$$s(x_k) = \sum_{i=1}^{M} \alpha_i g_i(x_k)$$

where the $g_i(x)$ can, for example, be polynomials or trigonometric functions. The constants α_i are then determined by fitting the series to the data, most commonly by least squares. The role of the span is played here by M, the number of terms included in the model. Trigonometric functions have been used with success in cases where the signal is naturally periodic. If the abscissas x_i are equi-spaced, the fit is particularly inexpensive to compute using the Fast Fourier Transform. Both conditions are usually not fulfilled in the case of scatterplot smoothing, making the method less attractive. The use of polynomials has the drawback that they are not well suited to represent a wide variety of commonly encourtered functions, for example, functions with asymptotes.

There are, of course, connections between smoothing by series expansion and smoothing by local averaging. If the series is fitted by least squares, the fitted values $s(x_i)$ are weighted averages of the responses y_i . Depending on the abscissas and the functions $g_i(x)$, the weights determining $s(x_i)$ might or might not be concentrated on responses with corresponding predictor values close to x_i . If they are, the series

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expansion method behaves like a local averaging method. An example of this is least squares fit of cubic splines which will be further discussed in Section 9.

3. A Simple Nonresistant Smoother

The simplest example for a local averaging type smoother is the moving average, where "ave" in equation (3) denutes the arithmetic mean. The size of the neighborhood is usually specified by the span, the number k of observations to be included in the averaging. Ne will assume k to be odd and the abscissas x_i to be in increasing order. The neighborhood can be chosen either symmetrically, containing k/2 observations to the left of x; and the same number to the right, or it can be chosen to contain the k nearest neighbors of x_i , including x_i . (We assume that k/2 is computed by integer division.) There are no general results on which of thes, two possibilities is better statistically. The nearest neighbors approach generalizes to higher dimensions, but the choice of a symmetric neighborhood is computationally simpler in that exactly one point enters and one point leaves the neighborhood as one moves from observation 1 to observation i+i. We will, in the following, use symmetric neighborhoods. The boundaries, where it is not possible to keep N symmetric, have to be treated specially; a commonly used adjustment is to shrink the neighborhood so that for i=1 and i=n, one averages only over k/2+1observations. With these conventions, the moving average smoother is defined by

$s(x_i) = mean(y_i) max(1-k/2,i) \le y \le min(1+k/2,n).$

Obviously, the mean does not have to be recomputed every time. It can be

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updated, reducing the computation from nk to n. Such updating can be done for all the smoothers we will consider, and is highly desirable because in typical applications k is 5% to 30% of n, and thus the savings are substantial. The simple moving average smoother has some serious shortcomings. One disturbing property is that it does not reproduce straight lines if the abscissa values are not equi-spaced. Another disturbing feature is the bad behaviour at the boundaries. If, for example, the slope of the underlying function f is positive at the right boundary, the estimate for observations close to the boundary will be biased downwards; if the slope is regative, the estimate is biased upwards. Both problems can be alleviated by fitting a least squares straight line L to the observations in the neighborhood instead of fitting a constant and taking the value of the line at x_1 as the smoothed value, This obviously solves both problems mentioned above. For the computation, again updating formulas can be used. The slope β and intercept α of the least squares straight line through a set of points $(x_1,y_1)...(x_m,y_m)$ are given by

$$\alpha = \overline{y}_{m} - \beta \overline{x}_{m}$$
$$\beta = \frac{C_{m}}{V_{m}}$$

with

$$\overline{\mathbf{x}}_{\mathbf{m}} = \sum_{\mathbf{x}} \mathbf{x}_{\mathbf{i}} / \mathbf{m},$$

$$\overline{\mathbf{y}}_{\mathbf{m}} = \sum_{\mathbf{y}} \mathbf{y}_{\mathbf{i}} / \mathbf{m},$$

$$\mathbf{C}_{\mathbf{m}} = \sum_{\mathbf{x}} (\mathbf{x}_{\mathbf{i}} - \overline{\mathbf{x}}_{\mathbf{m}}) (\mathbf{y}_{\mathbf{i}} - \overline{\mathbf{y}}_{\mathbf{m}}),$$

$$\mathbf{V}_{\mathbf{m}} = \sum_{\mathbf{x}} (\mathbf{x}_{\mathbf{i}} - \overline{\mathbf{x}}_{\mathbf{m}}) / \mathbf{v}$$

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When we want to add an observation (x_{m+1}, y_{m+1}) , we can make use of the following easily derived formulas:

$$\bar{\mathbf{x}}_{m+1} = (m\bar{\mathbf{x}}_m + \mathbf{x}_{m+1}) \neq (m+1),$$

$$\bar{\mathbf{y}}_{m+1} = (m\bar{\mathbf{y}}_m + \mathbf{y}_{m+1}) \neq (m+1),$$

$$C_{m+1} = C_m + \frac{m+1}{m} (\mathbf{x}_{m+1} - \bar{\mathbf{x}}_{m+1})(\mathbf{y}_{m+1} - \bar{\mathbf{v}}_{m+1}),$$

$$V_{m+1} = V_m + \frac{m+1}{m} (\mathbf{x}_{m+1} - \bar{\mathbf{x}}_{m+1})^2.$$

Analogous formulas can obviously be used for removal of an observation from the set.

4. Choice of Span

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The most important choice in the use of a local averaging smoother is the choice of the span value. If the smoother is regarded as a curve estimator, then the span controls the trade off between bias and variance of the estimate. We illustrate this for the case of a simple moving average smoother. In this case, the smoothed value at point x_i is given by

$$\begin{array}{rcl}
 1 & i+k/2 \\
 s(x_1) &= - & \sum_{k=1-k/2} y_3. \\
 & & 1-k/2
 \end{array}$$

If we assume that the errors ϵ_1 are 1.1.d., with expected value zero and variance σ^2 , then the expected squared error at point x₁ is

ESE
$$(x_i|k) = (f(x_i) - \sum_{k=1}^{j} f(x_j))^2 + \sigma^2$$
. (4)

Increasing the span will (if $d^2 f / ix^2 \pm 0$) increase the first term, the bias component of the estimation error and decrease the second term, the

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variance component; decreasing the span will have the opposite effect. The span should be chosen such that both compnents of the error are reasonably balanced. Stated more geometrically, a larger span makes the smooth appear less wiggly by more strongly damping high frequency components of the series (x_1,y_1) .

We have, so far, said nothing useful on how to choose the span in practice. The advice given above on balancing bias and variance is not very helpful because both f and the variance of the random error are unknown.

One can estimate the optimal span value in a particular situation as that value that minimizes an estimate for the integrated squared error

$$I^{2}(k) = \int ESE(x|k) dF(x).$$

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Using the average squared residual of the data from the smooth

$$\hat{1}^{2}(k) = -\sum_{n i=1}^{n} [y_{i} - s(x_{i}|k)]^{2}$$

for this purpose is not appropriate since this is always minimized by the span value k=1. A better estimate is provided by a method referred to as "cross-valida" on" (M. Stone, 1974) or "predictive sample reuse" (Geisser, 1975) Each observation is in turn deleted and the value of the smooth $s_{(1)}(x_1|k)$ at x_1 is calculated from the other n-1 observations. The cross-validated estimate of the integrated square error is

$$\hat{I}_{cv}^{2}(k) = -\sum_{n=1}^{n} [y_{1} - s_{(1)}(x_{n}|k)]^{2}.$$
(5)

Clearly, $E[\hat{I}^2_{cv}]$ equals the expected squared error obtained by applying the procedure to a sample of n-1 observations from the same distribution. The cross-validated estimate for the optimal span value is taken to be the value k_{cv} that minimizes (5),

$$k_{cv} = \min^{-1} \hat{I}^2_{cv}$$
 (k).
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Model selection through cross-validation has been remarkably successful in a wide variety of situations (see M. Stone, 1974, Geisser, 1975, Craven and Wahba, 1957, C. Stone, 1981).

For the moving average smoothers discussed in the previous section, the deleted smooth estimates $s_{(i)}(x_i|k)$ are especially easy to compute; each observation is simply deleted from the neighborhood used to compute its local straight line fit. Again, the use of updating formulas makes this computation very rapid. As one moves from observation i to i+1, exactly two observations enter the neighborhood (i and i+k/2+1) and exactly two leave it (i+1 and i-k/2). The (deleted) residual squared

$$r^{2}(i) = [y_{i} - e_{i}] (x_{i}|k]^{2}$$
(6)

is computed for each observation and then averaged over all observations,

$$\hat{1}^{2} cv(k) = -\sum_{n=1}^{n} r^{2}(i), \qquad (7)$$

For small to moderate changes in k, $\hat{1}^2_{cv}(k)$ changes very little so that it is adequate to evaluate it for several (4 to 7) discrete values of k in the range [0 < k/n ≤ 1]. The value of k corresponding to the smallest of these values is then used. This can be accomplished by maintaining several running average smoothers - one for each span value - in the pass over the data, thus keeping the computational cost linear in n.

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So far, we have been assuming that the span is constant over the whole range of predictor values. This is not optimal if either the variance of the random component or the second derivative of the underlying function f change over the range of predictor values. A local increase in error variance would call for an increase in span, whereas an increase in second derivative of f would require a decrease. It is, therefore, desirable to allow the span value to adapt to these changing conditions. This requires that the optimal span value be choosen locally rather than choosing a single global value. Again, the form of moving average smoothers make this especially easy; the (deleted) residual squared (6) -for each of the several k values- is averaged locally in a neighborhood of each observation

$$\hat{I}^{2}_{cv}(k;x_{i}) = -\sum_{k=1}^{1} r^{2}_{(k)}(x_{k}|k)$$
(8)

rather than globally over all observations (7). Note that (8) also has the form of a simple moving average smoother and can therefore be computed rapidly through the use of updating formulas. The value that minimizes (8)

$$\hat{k}_{cv}(x_i) = \min^{-1} \hat{1}_{cv}^2(k_i x_i)$$
 (9)
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is the span value used for the ith observation.

Most often the shape of \hat{r}_{cv}^{2} (k;x;) near its minimum value is shallow and asymmetric, increasing more slowly in the direction of smaller k values. Variability in the estimate $\hat{1}_{cv}^{2}$, therefore, causes \hat{k}_{cv} to be highly variable and biased toward smaller values. Although this has little effect on the quality of the resulting smooth in terms of expected squared error (ESE), it does effect its aesthetic quality since, for comparable ESE, the less smooth solution tends to be selected. This can be remedied by forcing the procedure to take the smoothest solution in these circumstances. Specifically, the largest span value \hat{k}_{ev} for which

$$\hat{1}^{2}_{cv}(k*_{cv};x_{i}) \leq (1+\alpha) \min \hat{1}^{2}_{cv}(k;x_{i})$$
(10)
 $0\langle k \leq N$

is used for the ith observation. Here a loosely controls an upper limit on the fraction of ESE that is to be sacrificed for the goal of smoothness. Values in the range 0.05 $\leq \alpha \leq$ 0.2 are reasonable.

Since the optimal span value is estimated separately for each observation, its size can vary substantially over the range of predictor values. However, since for close abscissa values the neighborhoods overlap considerably, this variation is constrained to be smooth. The degree of smoothness is controlled by the parameter L (8) which can be regarded as a span for smoothing the (deleted) residuals squared from the original smooths. As with the original smoother, its optimal value can be estimated by cross-validation. To the extent that the variation of the sec of derivative of f or the variation in the random component is comparable to the variation of f itself, this second level of crossvalidation may be beneficial. Again, updating formulas make this relatively inexpensive. However, in most circumstances choosing a nominal value for L (0 2n to 0.3n) is adequate.

It is important to note that using cross-validated residuals as a basis for choosing span value is highly sensitive to lack of independence among the ϵ_1 (2) as ordered on x. If there is a large positive (negative) correlation among observations with similar x values,

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substantial under (over) estimates will result. In situations where a high degree of auto-correlation is suspected, these span selection procedures should be used with caution.

Figure 1 illustrates the application of this smoothing algorithm to an artificial data set. (A FORTRAN subroutine implementing this algorithm is listed in Appendix 1.) The data for this example consists of M=200 pairs (x_i, y_i) with the x_i drawn randomly (iid) from a uniform distribution in the interval [0,1]. The y; are obtained from

 $y_{i} = s_{1n}[2\pi(1-x_{i})^{2}] + x_{i}\epsilon_{i}$

with the ϵ_i iid standard normal. The parameter ALPHA [α in (10)] was set to 0.1 and RESPAN [L/n in (8)] was set to 0.25 (see Appendix 1). This example simulates a situation in which both the curvature of f decreases and the variance of the random component increases with increasing x. Figure 1a is a scatterplot of the simulated data. Figure 1b also shows this scatterplot, but with the resulting smooth superimposed. The height of the curve near the bottom indicates the span value chosen at each \mathbf{x}_{s} The span is seen to increase with ix to account for the increasing noise, as well as to take advantage of the decreasing curvature of f. (For X > 0.7, the span has reached the largest value provided in the program, k/N = 0.7.) Figure 1c is the same as Figure 1b except that the curve $Y=f(x)=sin[2\pi(1-x)^2]$ is superimposed for reference. The resulting smooth is seen to estimate the underlying f reasonably well. Note that for a linear function $y=\alpha x+b$ (zero curvature) the smoother will tend to use a constant (maximum) span value regardless of (the variation of) the amplitude of the noise.

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5. <u>Reducing Computation by Binning</u>

In the previous sections, we have described a fairly intricate scatterplot smoother. As an essential building block of projection pursuit regression (Friedman and Stuetzle, 1981), it has performed well. In this context, the smoother is applied to the full data set many times in a single run. In order for such a procedure to be computationally feasible for large data sets, it is necessary that the smoother be as fast as possible. One possibility to increase speed is by binning. Denote the observations for one particular scatterplot by $(x_1,y_1),...(x_n,y_n)$. We assume here that the observations already have been sorted so that the x_i are in increasing order. Choose a bin size, say m, and define new data points $(u_1,v_1)...(u_{n/m},v_{n/m})$ by

> u; = mean (x(i-1)m+1...xim); v; = mean (y(i-1)m+1...yim).

Then apply the smoother described above to the $(u_1,v_1)...(u_{n/m},v_{n/m})$. The smooth for predictor values x_j not among $u_{1 < ... < u_{n/m}}$ can be obtained by linear interpolation or, at the boundaries, by extrapolation.

The computing time for the smoother grows linearly in the number of observations, and so binning reduces the run time of the smoother roughly by a factor of m.

Figure 2 shows the results of applying the smoother to a sample of n=500 observations generated from the same model as the data shown in Figure 1, with the results of applying the binning procedure with m=5 superimposed. The quality of the smooth is seen to suffer very little while the computation has been substantially reduced.

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6. <u>Resistance</u>

As for all data analytic procedures, it is highly desirable for a scatterplot smoother to be resistant against occasional outliers in the data. (All our analysis is conditional on the observed predictor values; nutlier thus means outlier in response.) The smoother described in Sections 4 and 5 clearly is not resistant. One way to overcome this limitation is to first screen the data with a rejection rule identifying outliers, and then apply the smoother to the cleaned data set.

We suggest a rejection rule based on running medians. A running median smoother with span k is defined by

 $s(x_i) = med(y_i-k/2...y_i+k/2)$

The ends of the sequence must be treated specially, most simply by replicating the outermost values defined above. The rejection rule makes five passes over the data:

- 1) Compute a running median smooth s.
- 2) Replace $s(x_i)$ by $s^*(x_i)$ obtained by linearly interpolating between $(x_{i-1}, s(x_{i-1}))$ and $(x_{i+1}, s(x_{i+1}))_{\infty}$. The purpose of this step is to ensure a more realistic estimate of spread in steps (3) and (4) for monotone (sub)sequences, which are exactly reproduced by a running median.
- ?) Smooth the absolute residuals $|r_1| = |y_1 s^*(x_1)|$ by a running median and obtain a sequence $v_1 \dots v_n$ of local spread estimates v^*_1 .

4) Smooth the sequence of local spread estimates by a moving

average with span f*n and obtain smoothed spreads $v*_i$. This makes the spread estimates more stable. The same effect could be achieved by increasing the span of the running medians in Step (3); however, this would be more expensive computationally. In the code given in Appendix 2, the constant f is set to 0.3.

5) Flag all observations for which $|\mathbf{r}_i| \ge c \cdot v *_i$. In the code, c is set to 4.5.

Some details related to the treatment of ties have been omitted. A FORTRAN subroutine implementing this algorithm is listed in Appendix 2.

The span for the running medians in Steps (1) and (3) is chosen to be increasing with the sample size n (see Table 2). A motivation for our particular choices is given in Chapter 7. We use the same span in both steps, although there is no inherent need to do so.

Figure 3a shows the result of applying the rejection rule to an artificial data set. The true underlying function is a sine wave. The predictors are uniformly distributed in $[0,2\pi]$; the random errors are Gaussian with standard deviation 0.3. Outliers occur with probability 0.2; they were generated by adding a Gaussian with standard deviation 2.4 to the original observation. Observations flagged as outliers by the rejection rule are marked by squares. Figure 3b shows the results of applying the rejection rule to a real data set.

7. Resistance of Running Medians

The choice of span k for the running medians in Steps (.) and (3)

gives rise to rather interest ng questions. Somewhat vaguely stated, the rejection rule will be able to detect extreme outliers as long as these running medians do not break down. We will now define precisely how we measure the degree of resistance of a smoother, and give results on the dependence of the resistance of a running median smoother on the span.

Assume we want to smooth a sequence of length n. Responses can be either "good" or "bad", that is, good observations or outliers. We define random variables $b_1 \dots b_n$ by $b_1 = 0$ if y_1 is good, $b_1 = 1$ if y_1 is Assume that $Prob(b_i = 1) = p$ and that the b_i are an outlier. independent. (As noted by Mallows «1980), the latter assumption might not always be realistic; outliers in time series sometimes come in bursts.) A smoothed value s, is called bad if it can be made arbitrarily large by suitable choice of the reponse values for the bad observations. A smoother is said to suffer a breakdown if one or more of the smoothed values s; are bad. The probability that this happens under the above model for the b is called the breakdown probability of the smoother. It will generally depend on p and n. A smoother with breakdown probability (1-p)ⁿ is called nonresistant. (For a different definition of breakdown probability, see Mallows (1980).)

We will now devise an approximate formula for the span necessary to guarantee an upper bound on the broakdown probability as a function of n and $p = Prob(b_1=1)$. For that purpose, we define new random variables $s_{1<3}, s_{n-k+1}$ by

$$s_1 = \sum_{j=1}^{j+k-1} b_j,$$

A running median smoother suffers a breakdown if among any consecutive k observations more than k/2 are bad; i.e., if max $s_i > k/2$. This probability does not seem to be simple to evaluate, but it is easy to obtain an upper bound in terms of the binomial probability Prob(B(K,p) > k/2, using the Bonferroni inequality:

Prob(max s; > k/2) \leq (n-k+1) Prob(B(k,p) > k/2). This inequality provides an estimate for the span needed to guarantee a certain upper bound on the breakdown probability for given n and p. Table 1 gives estimates of the necessary span k for breakdown probability bounded by 0.05, and several values of n and p (n=25,50,100,200,400,800; p=0.05,0.1,0.2). For a comparison, we also list the percentage of breakdowns actually observed in thousand randomly generated Bernoulli sequences for the estimated value of k, and the smallest value of k resulting in 50 or fewer breakdowns. The results show that the Bonferroni estimate is close, especially for p=0.05 and p=0.1 where Prob(B(k,p) > k/2) is small; this is in agreement with experience gained in using the Bonferroni inequality in multiple comparisons. The span of the running medians in Steps (1) and (3) of the rejection rule described in Chapter 6 was chosen to guarantee a breakdown probability of less than 0.05 for probability p=0.1 of obtaining an outlier.

Another interesting question is how fast the span k(n) must grow as a function of n with everything else fixed. This question has been answered by P. Erdös and A. Renyi (1970).

<u>Theorem</u>: If k(n) = cln n then

 $\frac{s_i}{r_i - \infty 1 \leq i \leq r_i - k + 1} = \alpha = 1,$

where α is related to c via the equation

$$\frac{-1}{-1} = \ln(p(1-p)) + (1-p-\alpha) (\ln \frac{\alpha}{1-\alpha} - \ln \frac{p}{1-p}),$$

for a > c.

This theorem 1. a special case of Erdös and Renyi's theorem 2. It can be applied to our situation as follows: Choose $\alpha = 1/2 - \epsilon$. Then there exists ar no such that for all $n > n_0$ we have max s; (k/2 for almost all sampling sequences. In addition, Erdös and Renyi show that

- If k grows slower than ln n $(k(n)/\ln n \rightarrow 0,)$, then for a'l but finitely many values of n, max s; = k for almost all sampling sequences, ("k cannot grow slower than ln n".)
- If k grows faster than ln n (k(n)/ln n→∞), then lim max s;=kp for almost all sampling sequences; i,e,, the strong law of large numbers applies. ("k does not have to grow faster than ln n".)

6. An Updating Algorithm for Running Medians

There is a straightforward way to compute running medians: Obtain the median of each consecutive k-tupel by sorting it. That can be substantially improved upon by making use of the fact that the set of responses defining s_{1+1} is almost the same as the set of responses defining s_i ; only $y_{1:n} = y_{1+k/2+1}$ has to be added, and $y_{out} = y_{1-k/2}$ has to be deleted. The following rules are easy to verify:

- If $y_{1n} = s_i$, then $s_{i+1} = s_i$.

- If $y_{1n} > s_1$ and $y_{out} > s_2$ or if $y_{1n} < s_1$ and $y_{out} < s_2$, then $s_{1+1} = s_1$.

So in the case of random data, we have to do nothing but make these tests half the time.

- If $y_{in} > s_i$ and $y_{out} < s_i$, then let k⁺ denote the number of observations in the new span that are bigger than s_i . If K⁺ < k/2, then $s_{i+1} = a_i$, else s_{i+1} is the smallest observation strictly bigger than s_i . The analog of that is true if $y_{in} < s_i$ and $y_{out} > s_i$.
- If $y_{in} > s_i$ and $y_{out} = s_i$, then define k⁺ as above. If k⁺ = k/2, then s_{i+1} is the smallest observation in the span strictly bigger than s_i ; else s_{i+1} is the smallest observation in the span that is bigger than or equal to s_i . The analog of that is true if $y_{in} < s_i$ and $y_{out} = s_i$.

In appendix 2, we give a FORTRAN subroutine that implements the algorithm outlined above.

For random data, the algorithm will take O(nk) operations. It is possible to reduce that to O(n log k) by organizing the observations in the span into a binary tree which is kept balanced as observations are moved in and out (AVL-tree: see Knuth (1973), pp 451). Unfortunately, for the range of k that we have in mind (about 20), log k is not enough smaller than k to compensate for the increased overhead.

9. Discussion

Cleveland (1979) has suggested a scatterplot smoother also based on local linear fits. It differs from the one described in this report mainly in three respects:

- It does not use variable span.

- In the fit of the local straight line determing the smooth $s(x_1)$ for

predictor value x_i , the observations are weighted according to their distance from x_i : observations towards the extremes of the span receive lower weights than observations with predictor values close to x_i . Asymptotic calculations suggest that assigning unequal weights should reduce the error of the curve estimate, but there is no evidence that it makes a substantial difference for sample sizes occurring in practice. It does, however, produce a smoother looking estimate.

- The procedure derives its resistance properties not from data screening with a rejection rule. Instead, each of the local straight lines is fitted, not by least squares, but by a resistant fitting procedure.

Updating formulas cannot be used in this scheme, making it comparatively expensive in terms of computing. To reduce computation, Cleveland suggests evaluating the smooth only for every m-th predictor value. The parameter m here plays a similar role as our bin size; it would be chosen as a fraction of k. We developed our smoothing procedure because variable span is often important, and because the use of updating formulas dramatically reduces computation.

Another class of procedures suggested for smoothing are procedures based on splines. A spline function s of order k with knots at $z_1...z_k$ is a function satisfying the following two conditions:

- In each of the intervals $(-\infty, z_1), (z_1, z_2) \dots (z_{k-1}, z_k), (z_k, \infty)$, s is a polynomial of degree l = 1;
- s has £ 2 continuous derivatives.

One way to use spline functions in scatterplot smoothing is to fit a spline function with knots $z_1, ..., z_k$ to the data $(x_1, y_1) ..., (x_n, y_n)$, either by least squares or by some resistant method. The degree of smoothness is determined by the number and position of the knots. A major disadvantage of this method is that k+1 parameters must be chosen: the number and the positions of the knots. Usually some heuristic procedure is used to place the knots once k has been fixed (Jupp, 1978). This leaves the number of knots to be determined. This number plays the role of the span in determining the degree of smoothing, Unfortunately, the output of the smoother can depend on k in a very nonlinear way; it is easy to construct examples where the addition of one more knot substantially decreases the residual sum of squares, whereas further knots hardly make any difference. This makes k more difficult to choose than the span in a local averaging smoother. Furthermore, least squares fit of splines is substantially slower so that choosing k through crossvalidation is usually too expensive.

Another way is to use smoothing splines in the sense of Reinsch (1967). A smoothing spline is of order 22 for smoothing parameter λ is the function that minimizes

$$\sum (y_{1} - f(x_{1}))^{2} + \lambda \int_{x_{1}}^{x_{n}} f(x_{1})^{2}(x) dx$$

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among all functions f with \mathcal{L} derivaties. The solution really turns out to be a spline function of order 2 \mathcal{L} with knots $x_{12} \dots x_n$; the name is thus justified. The larger λ is chosen, the smoother s becomes; thus, λ here plays the role of the span. Computation of the spline for given λ requires the solution of a banded $n \neq n$ linear system. A drawback of the method, as described here, is that it is impossible to ortain an intuitive feeling for the choice of λ in a given example. So, one usually fixes not λ , but the residual sum of squares around the smooth. The corresponding value of λ then has to be found iteratively by repeatedly solving the minimization problem. This substantially increases the necessary amount of computation. Algorithms to determine the optimal λ by cross-validation usually require computation of the singular value decomposition of an nxn matrix; they are expensive and infeasible for sample sizes larger them 200-300.

To summarize, the local averaging smoother described in this report has two desirable properties that set it apart om other scatterplot smoothers: it is very fast to compute and the value of the parameter that controls the amount of smoothing is optimized locally (through crossvalidation), allowing it to change over the range of predictor values,

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APPENDIX I

The following is a complete listing of a FORTRAN subroutine implementing the smoothing procedure described in this paper.

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فدعهم وعمدوتهم سنتعقق بكلا للتزمية تسلاكها فلمعتر مند معطيه مكترما فتحارف ولمتدعين

SUBROUTINE SUPSMU (N,X,Y,W, IPER, ALPHA, RESPAN, IBIN, SMO, SC) C----С C SUPER SMOOTHER (FRIEDMAN AND STUETZLE, 1982). C C CODED BY: J. H. FRIEDMAN AND W. STUET2LE С DEPARTMENT OF STATISTICS AND С STANFORD LINEAR ACCELERATOR CENTER С STANFORD UNIVERSITY С STANFORD CA. 94305 С C INPUT: N : NUMBER OF OBSERVATIONS (X, Y - PAIRS). С С X(N) : ORDERED ABSCISSA VALUES. С Y(N) : CORRESPONDING ORDINATE (RESPONSE) VALUES, С W(N) : (OPTIONAL) WEIGHT FOR EACH (X,Y) OBSERVATION. С W < 0 0 => ALL OBSERVATIONS HAVE EQUAL WEIGHT. С IPER : PERIODIC VARIABLE FLAG. С IPER=1 => X IS ORDINARY ORDERED VARIABLE. С IPER=2 => X 1S A PERIODIC (CIRCULARLY DEFINED) VA' ABLE. С ALPHA : FRACTIONAL SMOOTHNESS PENALITY (SEE (10) SECTION 4). RESPAN : FRACTIONAL SPAN FOR RESIDUAL SMOOTHING С С (L/N, SEE (8) SECTION 4). C, RESPAN .Lf. 0 => FIXED SPAN SMOOTHER WITH FRACTIONAL C_{i} SPAN = ABS(RESPAN). IBIN : BINNING FACTOR (M, SEE SECTION 7). С C OUTPUT: С SMO(N) ; SMOOTHED ORDINATE (RESPONSE) VALUES. C SCRATCH: С SC(3,N) : INTERNAL WORKING STORAGE. С C NOTE: ALPHA=0.1 AND RESPAN=0.25 AR" REASONABLE VALUES. FOR RESPAN > 0 С SMOOTHER OUTPUT IS COMPLEMELY CROSS-VALIDATED; X(I), Y(1), AND C W(I) ARE NOT USED IN THE CALCULATION OF SMO(1). THEREFORE, С C THE AVERAGE SQUARED RESIDUAL C N C $ASR = SUM \quad W(I) * (Y(I) - SMO(I)) * *2$ C I = 1С CAN BE USED AS A GOODNESS-OF-FIT MEASURE FOR THE PURPOSE OF SELECTING OPTIMAL VALUES FOR SMOOTHING LARAMETERS BY C С REPEATED APPLICATION. С С FOR SMALL SAMPLES (N < 40) OR IF THERE ARE SUBSTANTIAL SERIAL CORRELATIONS BETWEEN OBSERATIONS CLOSE IN X - VALUE, THEN С A PRESPECIFIED FIXED SPAN SMOOTHER (RESPAN <)) SHOULD BE С C USED. REASONABLE SPAN VALUES ARE 0.3 .GE, ABS(RESPAN) .GE, 0.5. C C-DIMENSION X(N), Y(N), W(N), SMO(N), SC(3, N)DOUBLE PRECISION SX(5), SY(5), SXX(5), SXY(5), SUM(5), FBW(5)DIMENSION IBW(5), RESQUE(5,101), SMOQUE(5,101) INTEGER IN, OUT DATA IBW1, SPEMAX /3,0,35/ DATA BIG, EPS /1 0F20, 1,0E-03/

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-25-
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IF (W(1).LT.0.0) GO TO 20 DO 10 I=1,N SC(3,I)=W(I)10 CONTINUE GO TO 40 DO 30 I=1,N 20 SC(3, I) = 1. CONTINUE 30 IF (X(N).GT.X(1)) GO TO 70 40 SX(1) = 0.0FBW(1)=SX(1)DO 50 J=1,N SX(1)=SX(1)+SC(3, J)*Y(J)FBW(1) = FBW(1) + SC(3, J)50 CONTINUE A=SX(1)/FBW(1)DO 60 J=1,N SMO(J) = A60 CONTINUE RETURN 70 I=N/4J = 3 * ISCALE = X(J) - X(I)IF (SCALE.GT.0.0) GO TO 90 80 IF (J.LT.N) J=J+1 1F (I GT 1) I = I - 1SCALE = X(J) - X(I)GO TO 80 VSML=(EPS*SCALE)**2 90 IF (IBIN.LE.1) GO TO 11J NA=0 SX(1)=0.0SY(1) = SX(1)FBW(1)=SY(1)DC 100 J=1, N $S_{\lambda}(1) = S_{\lambda}(1) + X_{\lambda}(J) * S_{\lambda}(3, J)$ SY(1) = SY(1) + Y(J) * SC(3, J)FBW(1) = FBW(1) + SC(3, J)IF (MOD(J,IBIN),NE.0) GO TO 100 NA=NA+1 SC(1, NA) = SX(1) / FBW(1)SC(2, NA) = SY(1) / FBW(1)SC(3, NA) = FBW(1)SX(1) = 0.0SY(1) = SX(1)FBW(1) = SY(1)100 CONTINUE IF (MOD(N, IBIN), EQ.0) GO TO 130 NA=NA+1SC(1, NA) = SX(1)/FBW(1)SC(2 NA) = SY(1) / FBW(1)SC(3, NA) = FBW(1)GU TU 130 110 NA=N DO 120 1-1,N

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	SC(1, J) = X(J)
100	SC(2, J) = Y(J)
120	
120	1BW(1)=1BW1
	1001401=2.5
	IBW(T) = MINO(NA/2, IBW(T-1) + IDELTA)
140	CONTINUE
	DO 150 I=1.5
	SX(I) = 0.0
	SY(I) = SX(I)
	SXX(I) = SY(I)
	SXY(I) = SXX(I)
	FBW(I) = CXY(I)
	SUM(I) = FBW(I)
	IF $(MOD(1BW(1), 2) \cdot EQ \cdot O)$ $IBW(1)=IBW(1)+1$
150	CONTINUE
	IF (RESPAN.GE.0.0) GO TO 160
	IBW(1)=0.5 ABS(RESPAN) NA IBW(1)=1BW(1) > 1
	IC (MOD(IBW(1), 2), EQ.0) IBW(1)=IBW(1)+1 IDW(E)=IBW(1)
	$\frac{10}{10} = 10$
160	18WS=5
170	IF (IPER.NE.2) GO TO 220
	IT=NA-IBW(5)+1
	IH=IBW(5)-1
	DO 190 J=IT, NA
	DO 180 I=1,IBWS
	IF $(J.LT.NA-IBW(I)+1)$ GO TO 180
	XT = SC(1, J)
	YT = SC(2, J)
	WT = SC(3, J)
	SX(1) = SX(1) + XT * WT
	SA(T)=SA(T)+AL.M. SA(T)=SA(T)+AL.M.
	SXX(1)~SXX(1)+X1*X1*W1 CVV/()~CVV/()+Y#####
	$FEW(T) = FRW(T) + W\Omega$
180	CONTINUE
190	CONTINUE
	DO 210 J=1,1H
	DO 200 I=1, IBWS
	IF (J.GT. BW(I)-1) GO TO 200
	XT=SC(1,J)
	YT = SC(2, J)
	WT = SC(3, J)
	$SX(T) = SX(T) + XT^*WT$
	<pre>cvv(t)=cvv(t)+t1*Wt 21(1)=51(1)+11*Wt </pre>
	SXX(1)=SXX(1)+XT*XT*XT SXY(1)=SXY(1)+XT*YT*WT
	FBW(1) = FBW(1) + WT
200	CONTINUE
210	CONTINUE
	GO TO 250
220	$1T = 2 \le 1BW(5) = 1$

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```
DO 240 J=1,IT
     DO 230 I=1, IBWS
     IF (J.GT.2*IBW(I)-1) GO TO 230
     XT = SC(1, J)
     YT = SC(2, J)
     WT = SC(3, J)
     SX(I) = SX(I) + XT * WT
     SY(I) = SY(I) + YT * WT
     SXX(I)=SXX(I)+XT*XT*WT
     SXY(I) = SXY(I) + XT*YT*WT
     FBW(I) = FBW(I) + WT
230 CONTINUE
240
    CONTINUE
250
     KBW=MINO(101,2*INT(0.5*RESPAN*NA+0.5)+1)
     KBWO2 = KBW/2+1
     IH=0
     JT=IH
     JM≕JT
     DO 370 J=1,NA
      RESMIN=BIG
      IF (J.LT.KBWO2) GO TO 260
      JT≈JT+1
      JM0=JM
      JM = MOD(JM, KBW) + 1
260 IH=MOD(IH,KBW)+1
      DO 310 I=1,1BWS
      IF (IBWS.NE.5) GO TO 270
      XT = SC(1, J)
      YT=SC(2,J)
      WT = SC(3, J)
      SX(I) = SX(I) - XT*WT
      SY(I) = SY(I) - YT * WT
      SXX(I) = SXX(I) - XT * XT * WT
      SXY(1) = SXY(1) - XT * YT * WT
      FBW(I) = FBW(I) - WT
270
      OUT = J - IBW(I)
      IN=J+IBW(I)-1
      IF ((IPER.NE.2).AND. (OUT.LT. I.OR. IN.GT.NA)) GO TO 280
      IF (OUT.LT.1) OUT=NA+OUT
      IF (IN GT NA) IN=IN-NA
      XT = SC(1, OUT)
      YT = SC(2, OUT)
      WT = SC(3, OUT)
      SX(I)=SX(I)-XT*WT
      SY(I) = SY(I) - YT * WT
      SXX(1) = SXX(1) - XT * XT * WT
      SXY(I) = SXY(I) - XT * YT * WT
      FBW(1) = FBW(1) - WT
      XT = SC(1, IN)
      YT = SC(2, IN)
      WT = SC(3, 1N)
      SX(1) = SX(1) + XT * WT
      SY(I) = SY(I) + YT * WT
      SXX(1) = SXX(1) + XT * XT * WT
      SXY(1) = SXY(1) + XT * YT * WT
```

```
FBW(I) = FBW(I) + WT
280
    D=SXX(I)-SX(I)**2/FBW(I)
     VAR=D/FBW(I)
     A=0.0
     IF (VAR.GT.VSML) A = (SXY(I) - SX(I) + SY(I) / FBW(I)) / D
     SM=A*SC(1, J)+(SY(1)-A*SX(1))/FBW(1)
     IF (IBWS.NE.1) GO TO 290
     SMO(J) = SM
     GO TO 320
290
    RES=SC(3,J)*(SC(2,J)-SM)**2
     IF (J.GT.KBW) SUM(I)=SUM(I)-RESQUE(I,IH)
     SUM(I) = SUM(I) + RES
     RESQUE(I, IH)=RES
     SMOQUE(I, IH)=SM
     IF (VAR.LT.VSML.AND.I.LT.5) SMOQUE(I,IH)=BIG
     IF (J.LT.KBWO2) GO TO 300
     SUM(I) = SUM(I) - RESQUE(I, JM)
     IF (JT.GT.1) SUM(I)=SUM(I)+RESQUE(I, JMO)
     IF (SUM(1).GT.RESMIN.OR.SMOQUE(I,JM).GE.BIG) GO TO 300
     RESMIN=SUM(I)
     IS=I
300
     XT = SC(1, J)
     YT=SC(2,J)
     WT = SC(3, J)
     SX(I) \approx SX(I) + XT * WT
     SY(I) = SY(I) + YT * WT
     SXX(I)=SXX(I)+XT*XT*WT
     SXY(I) = SXY(I) + XT*YT*WT
     FBW(I) = FBW(I) + WT
310 CONTINUE
320
     IF (IBWS.EQ.1) GO TO 370
     IF (J.GE.KBWO2) SMO(JT)=SMOQUE(IS,JM,
     IF (ALPHA.LE.O.O.OR.J.LT.KBWO2.OR.IS.GE.5) GO TO 370
     RESMIN=(1.0+ALPHA)*RESMIN
     I = 5
     GO TO 340
330
     I = I + (-1)
     IF ((-1)*((1)-(IS)).GT.0) GO TO 350
340
     IF (SUM(I).GT.RESMIN) GO TO 330
350
    IF (I.GE.5) GO TO 360
     A = (RESMIN-SUM(I)) / (SUM(I+1)-SUM(I))
     SMO(JT) = (1 \circ 0 - A) * SMOQUE(1, JM) + A * SMOQUE(I+1, JM)
     GO TO 370
360
     SMO(JT) = SMOQUE(I, JM)
370
     CONTINUE
     IF (IBWS.NE.5) GO TO 440
     JT=JT+1
     DO 430 J=JT, NA
     IH=MOD(IH,KBW)+1
     RESMIN=BIG
     JMU-JM
     JM=MOD(JM, KBW)+1
     DO 380 [=1,5
     SUM(1) = SUH(1) - RESQUE(1, IH) + RESQUE(1, JMO) - RESQUE(1, JM)
     IF (SUM(I).GT.RESMIN.OR, SMOQUE(I, JM).GE.BIG) GO TO 380
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RESM_N=SUM(I)
     IS=I
380
     CONTINUE
     SMO(J)=SMOQUE(IS,JM)
     IF (ALPHA.LE.O.O.OR.IS.GE.5) GO TO 430
     RESMIN=(1.0+ALPHA)*RESMIN
     I=5
     GO TO 400
390
     I = I + (-1)
400
     IF ((-1)*((I)-(IS)).GT.0) GO TO 410
     IF (SUM(I).GT.RESMIN) GO TO 390
410
     IF (I.GE.5) GO TO 420
     A = (RESMIN-SUM(I)) / (SUM(I+1)-SUM(I))
     SMO(J) = (1.0-A) * SMOQUE(I, JM) + A * SMOQUE(I+1, JM)
     GO TO 430
420
     SMC(J) = SMOQUE(I, JM)
430
     CONTINUE
440
     IT=NA-1
     S2=SMO(1)
     IF (IPER.NE.2) GO TO 450
     A=S2
     SMO(1)=0.25*(SMO(NA)+2.0*S2+SMO(2))
     GO TO 460
450
     SMO(1)=0.25*(2.0*S2+3.0*SMO(2)-O(3))
460
    DO 470 J=2,1T
     S1=S2
     S2=SMO(J)
     SMO(J)=0.25*(S1+2.0*S2+SMO(J+1))
470
     CONTINUE
      IF (IPER.NE.2) GO TO 480
     SMO(NA) = 0.25*(A+2.0*SMO(NA)+S2)
     GO TO 490
     SMO(NA)=0.25*(2.0*SMO(NA)+3.0*S2-S1)
480
490
     IF (IBINALEAL) GO TO 550
     DO 500 1=1,NA
     SC(2, 1) = SMO(1)
500
     CONTINUE
     XUP = SC(1, 1) - 1.
     J=0
     DO 540 1=1, N
     (1) X = 1 X
      IF (XI.LE.XUP) GO TO 530
     J = J + 1
     XLOW = SC(1, J)
     XUP = SC(1, J+1)
     YLOW = SC(2, J)
     YUP = SC(2, J+1)
     IF (XLOW.NE.XUP) GO TO 510
     SLOPE=0.
     GO TO 520
510
     SLOPE = (YUP - YLOW) / (XUP - XLOW)
520
     IF (J+1, EQ, NA) XUP=X(N)
530
     SMO(1)=YLOWr(X1-XLOW)*SLOPE
540
     CONTINUE
550
     J=1
```

560	J0=J
	SY(1) = SMO(J)
	IF (W(1).LE.O.O) GO TO 570
	SY(1) = W(J) * SMO(J)
	FBW(1) = W(J)
570	IF (J.GE.N) GO TO 610
580	IF $(X(J+1).GT.X(J))$ GO TO 610
	J=J+1
	IF (W(J).GT.0.0) GO TO 590
	SY(1)=SY(1)+SMO(J)
	GO TO 600
590	SY(1)=SY(1)+W(J)*SMO(J)
	FBW(1) = FBW(1) + W(J)
600	IF (J.LT.N) GO TO 580
610	IF (J.LE.JO) GO TO 630
	IF $(W(1).LE.0.0)$ FBW(1)=J-J0+1
	SY(1)=SY(1)/FBW(1)
	DO 620 I=J0,J
	SMO(I)=SY(1)
620	CONTINUE
630	J=J+1

IF (J.LE.N) GO TO 560

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RETURN
END
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APPENDIX II

The following is a complete listing of a FORTRAN subroutine implementing the rejection rule described in this paper.

SUBROUTINE REJECT (PRED, RESP, N, WEIGHT, SCRAT) C----С C REJECTION RULE FOR SMOOTHING (FRIEDMAN AND STUETZLE, 1982) С C CODED BY: J. H. FRIEDMAN AND W. STUETZLE C DEPARTMENT OF STATISTICS AND C STANFORD LINEAR ACCELERATOR CENTER C STANFORD UNIVERSITY C STANFORD, CA. 94305 C C INPUT: С PRED(N) :ABSCISSA VALUES IN INCREASING ORDER :CORRESPONDING ORDINATE (RESPONSE) VALUES RESP(N) Ċ J :NUMBER OF OBSERVATIONS (X, Y-PAIRS) N Ç C OUPUT: С WEIGHT(N) : REJECTION FLAGS. С WEIGHT(I)=0 IF OBSERVATION I IS CONSIDERED AN OUTLIER C WEIGHT(I)=1 OTHERWISE Ċ C SCRATCH: C SCRAT(N,2): INTERNAL WORKING STORAGE С Ċ C NOTE: REJECT USES SUBROUTINE RUNMED (SEE BELOW) \mathbb{C} ŝ C----DIMENSION PRED(N), RESP(N), WEIGHT(N), SCRAT(N, 2) DATA FACT/4.5/ DATA RELSPA/0.3/ IF (N.GT.25) GO TO 10 IBAND=7 GO TO 50 10 IF (N.GT.100) GO TO 20 IBAND=9 CO TO 50 20 IF (N.GT.400) GO TO 30 IBAND=11 GO TO 50 IF (N.GT.800) GO TO 40 30 IBAND=13 GO TO 50 40 IBAND=15 50 CALL RUNMED (RESP, WEIGHT, N, IBAND) IFIRST=IBAND/2+1 [LAST=N-IBAND/2 DO 60 I=1, IFIRST SCRAT(I,1) = WEIGHT(I)CONTINUE 60 DO 70 I=ILAST, N SCRAT(I, 1) = WEIGHT(I)70 CONTINUE

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DO 90 I=IFIRST, ILAST
     IMI = I - I
     IP1=I+1
     IF (PRED(IM1).NE.PRED(IP1)) GO TO 80
     SCRAT(I,1)=0.5*(WEIGHT(IM1)+WEIGHT(IP1))
     GO TO 90
80
     SCRAT(I,1)=WEIGHT(IM1)+(WEIGHT(IP1)-WEIGHT(IM1))*(PRED(I)-PRED(IM1))
    1))/(PRED(IP1)-PRED(IM1))
90
     CONTINUE
     I=0
100
     IF (1.GE.N-1) GO TO 150
     I=I+1
     MO=I
110
    IF (PRED(I+1).GT.PRED(I)) GO TO 120
     I = I + 1
     IF (I.LT.N) GO TO 110
     IF (I.EQ.MO) GO TO 100
120
     NTIE=I-MO+1
     R=0.
     DO 130 J=M0,I
     R=R+SCRAT(J,1)
130
     CONTINUE
     R=R/NTIE
     DO 140 J=M0,I
     SCRAT(J, 1) = R
140 CONTINUE
     GO TO 100
150 DO 160 I=1,N
     WEIGHT(I)=ABS(RESP(I)-SCRAT(I,1))
160 CONTINUE
     CALL RUNMED (WEIGHT, SCRAT(1,1), N, IBAND)
     IS2=N*RELSPA/2.
     SUM=0.
     DO 170 I=1, IS2
     SUM=SUM+SCRAT(I,1)
170 CONTINUE
     ISEFF=IS2
     DO 200 I=1,N
     IF (I.GT.N-IS2) GO TO 130
     SUM=SUM+SCRAT(I+IS2,1)
     ISEFF=ISEFF+1
180
     1F (1.LE.IS2+1) GO TO 190
     SUM=SUM-SCLAT(I-1S2-1,1)
     ISEFF=ISEFF-1
190
     SCRAT(1,2"=SUM/ISEFF
200
     CONTINUE
     0 = 1
210
     IF (I.GE.N-1) GO TO 260
     I=I+1
     MO=I
220 IF (PRED(I+1).GT.PRED(I)) GO TO 230
     I = I + I
     1F (1.LT.N) GO TO 220
230
     IF (1.EQ.MO) GO TO 210
     NTIE = I - MO + I
```

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R=0.
      DO 240 J=M0,I
      R=R+SCRAT(J,2)
 240 CONTINUE
      R=R/NTIE
      DO 250 J=M0,I
      SCRAT(J, 2) = R
 250 CONTINUE
      GO TO 210
 260
     DO 280 I=1,N
      IF (WEIGHT(I).LE.FACT*SCRAT(I,2)) GO TO 270
      WEIGHT(I)=0.
      GO TO 280
 270 WEIGHT(I)=1.
 280 CONTINUE
      RETURN
      END
C----
С
C
      SUBROUTINE RUNMED (SEQ, SMO, N, IBAND)
C --
                                          С
C FAST RUNNING MEDIAN FINDER (FRIEDMAN AND STUETZLE, 1982).
С
C CODED BY : J. H. FRIEDMAN AND W. STUETZLE
Ċ
             DEPARTMENT OF STATISTICS AND
С
             STANFORD LINEAR ACCELERATOR CENTER
С
             STANFORD UNIVERSITY
C
             STANFORD, CA. 94305
С
С
C INPUT:
С
             :RESPONSES IN ORDER OF INCREASING PREDICTOR VALUES
     SEQ(N)
              :NUMBER OF OBSERVATIONS
С
     N
C
              :SPAN OF RUNNING MEDIANS (HAS TO BE ODD AND <=21)
     IBAND
C
C OUTPUT:
С
     SMO(N)
             :SMOOTHED RESPONSES
С
C NOTE:
     THE MAXIMAL SPAN CAN BE INCREASED BY INCREASING THE DIMENSION
С
С
     OF THE ARRAYS SCRAT AND ITAG
С
C ----
          DIMENSION SEQ(N), SMO(N)
      DIMENSION SCRAT(21), ITAG(21)
      DATA RINE/1.E20/
      DO 10 I=1, IBAND
      SCRAT(I) - SEQ(I)
      1 TAG(I) = 1
      CONTINUE
 1 () I
      RMIN=SCRAT(1)
      IMIN=1
      DO 20 I=2, IBAND
```

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	IF (SCRAT(I).GE.RMIN) GO TO 20 RMIN=SCRAT(I)
	IMIN=I
20	CONTINUE
	TEMP = SCRAT(1)
	SCRAT(1) = RMIN
	SCRAT(IMIN)=TEMP
	TTAG(1) = 1MIN
	1TAG(1MIN)=1
	1=3 no mo 40
20	
30	1=1+1
40	IF ((1), GT, (IBAND)) GO TO 60
	IF (SCRAT(1), GE, SCRAT(1-1)) GU TU 30
	IEMP=SCRAI(I)
50	
10	$I_{\text{TAC}}(J) = I_{\text{TAC}}(J-J)$
	$\frac{1}{12} = \frac{1}{12} $
	SCRAT(U=1):SI:IEMP = GO = IO = JO
	TTAG(.T) = TTEMP
	GO TO 30
60	(BAND2=(BAND/2+1)
	RMED=SCRAT(IBAND2)
	DO 70 $I=1$, IBAND2
	SMO(I)=RMED
70	CONTINUE
	IFIRST=2
	ILAST=IBAND+1
	ISMO=IBAND2+1
	TMED=RMED
80	YIN=SEQ(ILAST)
	YOUT=SEQ(IFIRST-1)
	IF (YIN.GE.RMED) GO TO 180
	IF (YOUT.GE.RMED) GO TO 90
	RNEW=RMED
	GO TO 290
90	IF (YOUT.LE.RMED) GO TO 120
	KMINUS=0
	RNEW=-RINF
	DO 110 I=IFIRST, ILAST
	SI = SEQ(1)
	IF (SI<&MED) GO TO 100
100	CO TO TTO
100	TE (CI TE DNEW) CO MO 110
	RNEW=SI
110	CONTINUE
	IF (KMINUS.GE, IBAND2) GO TO 290
	RNEW=RMED
	GO TO 290
120	KM1NUS=0
	3TS=-RINF

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20

RSE=-RINF DO 160 I=IFIRST, ILAST SI = SEQ(I)IF (SI.LE.RMED) GO TO 130 GO TO 160 IF (SI.GE.RMED) GO TO 150 130 KMINUS=KMINUS+1 IF (SI.LE.RTS) GO TO 140 RTS=SI 140 IF (SI.LE.RSE) GO TO 160 RSE=SI GO °O 160 150 RSE=SI CONTINUE 160 IF (KMINUS.NT.IBAND2) GO TO 170 RNEW=RTS GO TO 290 170 RNEW=RSE GO TO 290 180 IF (YIN.LE.RMED) GO TO 280 IF (YOUT.LE.RMED) GO TO 190 RNEW=RMED GO TO 290 190 IF (YOUT.GE.RMED) GO TO 220 KPLUS=0 RNEW=RINF DO 210 I=IFIRST, ILAST SI = SEQ(I)IF (SI.GT.RMED) GO TO 200 GO TO 210 200 KPLUS=KPLUS+1 IF (SI.GE.RNEW) GO TO 210 RNEW=SI 210 CONTINUE IF (KPLUS.GE.IBAND2) GO TO 290 RNEW=RMED GO TO 290 220 KPLUS=0 RTB=RINF RBE=RI DO 260 FIRST, ILAST SI = SEQ(1)IF (SI.GE.RMED) GO TO 230 GO TO 260 IF (SI.LE.RMED) GO TO 250 230 KPLUS=KPLUS+1 IF (S1.GE.RTB) GO TO 240 RTB=SI 240 IF (SI.GE, RBE) GO TO 260 RBE=SI GO TO 260 250 RBE=SICONTINUE 260 IF (KPLUS.NE.IBAND2) GO TO 270

RNEW=RTB

2

-- -

	GO TO 290			
270	RNEW=RBE			
	GO TO 290			
280	RNEW=RMED			
290	RMED=RNEW			
	SMO(ISMO)=RMED			
	IFIRST=IFIRST+1			
	ISMO=ISMO+1			
	ILAST=ILAST+1			
	IF (ILAST.LE.N)	GO	TO	80
	DO 300 I=ISMO,N			
	SMO(I) = RMED			
300	CONTINUE			
	RETURN			

END

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Figure Captions

- Figure 1a: Two hundred observations (points) drawn from the model $Y=sin[2\pi(1-X)^2] + X\epsilon$ with ϵ iid standard normal.
- Figure 1b: The data of Figure 1a with the computed smooth superimposed. The height of the bottom curve is proportional to the span value employed at the corresponding abscissa value.
- Figure 1c: Same as Figure 1b with the addition of the curve $Y=sin[2\pi(1-X)^2]$
- Figure 2: Five hundred observations from the same model as Figure 1, with the computed smooths for both m=1 and m=5.
- Figure 3a: Output of rejection rule applied to artificial dat. set. Rejected observations are marked by squares.

Figure 3b: Output of rejection rule applied to real data set.



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FIGURE 1b



FIGURE 1c



FIGURE 2

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FIGURE 3a

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FIGURE 3b

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TABLE	I	

n	Pbad	ĸ	%bds	к
25	0.05	5	1.0	5
50	0.05	7	0.5	5
100	0.05	7	1.0	7
200	0.05	7	2.0	7
400	0.05	9	1.0	7
800	0.05	9	2.0	9
25	0.1	9	0.5	7
50	0.1	9	2.3	9
100	0.1	11	1.4	9
200	0.1	13	0.5	11
400	0.1	13	1.0	11
800	0.1	15	0.0	13
25	0.2	15	1.9	13
50	0,2	21	0.7	15
100	0.2	23	1.4	19
200	0.2	27	1.8	21
400	0.2	31	1,9	27
800	0.2	33	2.1	31

- n: length of sequence
- Pbad: probability of an outlier
- K: Bonferroni estimate of span necessary to guarantee breakdown probability ≤ 0.05
- %bds: Percentage of breakdown actually observed in 1000 Monte Carlo trials for span \widetilde{K}_{\circ}
- K: Span necessary to guarantee breakdown probability & 0.05 (estimated from 1000 Monte Carlo trials),

.

TABLE II

n	к
\$ 25	7
\$100	9
\$400	11
\$800	13
>800	15

- n: length of sequence
- K: span of running medians in steps (1) and (3) of rejection rule

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