Rejoinder to the discussants Nils Lid Hjort and Gerda Claeskens

- August 2003 -

We are honoured to have our work read and discussed at such a thorough level by several experts. Words of appreciation and encouragement are gratefully received, while the many supplementary comments, thoughtful reminders, new perspectives and additional themes raised are warmly welcomed and deeply appreciated. Our thanks go also to JASA Editor Francisco Samaniego and his editorial helpers for organising this discussion.

Space does not allow us answering all of the many worthwhile points raised by our discussants, but in the following we make an attempt to respond to what we perceive of as being the major issues. Our responses are organised by themes rather than by discussants. We shall refer to our two articles as 'the FMA paper' and 'the FIC paper'.

1. The local neighbourhood framework

In our articles we chose to work inside a broad and general parametric framework, which in the regression case corresponds to our using say

$$f_{i,\text{true}}(y) = f(y \mid x_i, \sigma_0, \beta_0, \gamma_0 + \delta/\sqrt{n}); \qquad (1.1)$$

see Section 2 in the FMA and Section 2 in the FIC paper. This draws partial criticism from Raftery and Zheng, who question its realism, as well as from Ishwaran and Rao, who argue that it does not yield a good framework for subset regression problems.

The local neighbourhood framework (1.1) allows one to extend familiar standard i.i.d. and regression models (corresponding to having δ fixed at zero) in several parametric directions (corresponding to $\delta_1, \ldots, \delta_q$ allowed to be non-zero, for different envisaged departures from the start model), as exemplified in our papers. This may in particular be utilised for robustness purposes and sensitivity analyses, and leads to a fruitful theory for model averaging and focussed model selection criteria, as we have demonstrated.

In their Section 4, Raftery and Zheng mention two pro (1.1) arguments, before presenting their scruples. The main argument for working inside (1.1) is however that it leads to natural, general and precise limit distribution results, with consequent approximations for mean squared errors and the like; the key is that variances and squared modelling biases become exchangeable currencies, both of size 1/n. For classes of estimators $\hat{\mu}$ of $\mu(\theta, \gamma)$, including the submodel estimators $\hat{\mu}_S = \mu(\hat{\theta}_S, \hat{\gamma}_S, \gamma_{0,S^c})$, we have

$$E_{\theta,\gamma}\{\widehat{\mu} - \mu(\theta,\gamma)\}^2 = n^{-1}\rho_1(\theta,\sqrt{n}(\gamma-\gamma_0)) + n^{-3/2}\rho_2(\theta,\sqrt{n}(\gamma-\gamma_0)) + \cdots,$$
 (1.2)

for example, under regularity conditions. Such expansions, written out here without the δ that Raftery and Zheng appear to dislike, would typically be valid uniformly over $\|\gamma - \gamma_0\| \le \text{const.}/\sqrt{n}$ balls. We view (1.2) type results as a good reason for developing and presenting

theory in terms of $\delta = \sqrt{n}(\gamma - \gamma_0)$, i.e. using (1.1). Our papers have (in particular) provided formulae for $\rho_1(\theta, \delta)$ here, the limiting risk for $\hat{\mu}$, while expressions for $\rho_2(\theta, \delta)$ are harder to get hold of; see our response below to Tsai's comments. We have also noted, in FIC's Section 5.5, that approximations coming from using the leading term in (1.2) type expansions hold with exactness for finite n, for submodel estimators of means in linear regression.

Thus Raftery and Zheng interpret us a little bit too literally, at the end of their Section 4; as statisticians we do not believe that our model parameter γ changes value when our data set passes from n = 100 to n = 101, but we do believe that limit theorems based on the (1.1) framework provides lucent understanding and useful approximations for the given n. This comment also applies to our BMA investigations (FMA's Section 9), where priors and posteriors for (θ, γ) are transformed to priors and posteriors for (θ, δ) . (A too literal belief in sample-size dependent parameters would clash with Kolmogorov consistency and other requirements for natural statistical models; see McCullagh (2002) and the ensuing discussion.)

A further strand of arguments supporting the view that many questions find their most natural solutions inside $\gamma - \gamma_0 = O(1/\sqrt{n})$ frameworks is related to what we termed 'tolerance radii' in FMA's Section 10.5. How much quadraticity, or variance heteroscedasticity, can the normal regression model tolerate, in the sense that the simpler methods based on standard assumptions still give better results than the more cumbersome ones based on the bigger models? How much autocorrelation can typical i.i.d.-based methods take? Such questions are nicely answered using the sample-size dependent magnifying glass $\delta = \sqrt{n}(\gamma - \gamma_0)$, as touched on in Section 10.5 of the FMA paper. Consider for example the $(\beta_0, \sigma, \beta_1, \lambda)$ model of FIC's Section 4.1. The simple i.i.d. model $Y_i \sim N(\beta_0, \sigma^2)$ can tolerate the presence of a regression coefficient β_1 and a skewness parameter λ , as long as

$$\sqrt{n}|\omega_1\beta_1 + \omega_2(\lambda - 1)| \le (k_{n,1}\omega_1^2 + k_{n,2}\omega_2^2)^{1/2}.$$

This first-order asymptotic answer rests on a framework like (1.1), and depends also on the focus parameter under study via (ω_1, ω_2) ; see FIC's Section 4.1 for examples. Inside the ellipse $k_{n,1}\beta_1^2 + k_{n,2}(\lambda - 1)^2 \leq 1/n$, all estimands will be better estimated using the simple $N(\beta_0, \sigma^2)$ model than using the the formally correct four-parameter model.

Another benefit of our methodology, and the (1.1) type framework, is the ability to compare model selection and model average strategies in a unified way, across situations, so to speak; there is a well-defined limit experiment, characterised by deterministic quantities τ_0 and K, the vector ω which depends on the focus parameter, plus an unknown δ for which one observes $D \sim N_q(\delta, K)$. Inference is then sought for $\psi = \omega^t \delta$. Thus lessons learned for e.g. Poisson regression models can be carried over to e.g. logistic regressions. This is also in the LeCam spirit of asymptotic equivalence of statistical experiments; see e.g. van der Vaart (1998) and Brown (2000) for general discussion.

Note that δ can be big in size, in (1.1) and (1.2), so reading our papers as saying that we only care about γ being close to γ_0 is not correct. Also, when γ happens to be far away from γ_0 , this will be picked up by the data via $\hat{\delta}_{\text{full}} = \sqrt{n}(\hat{\gamma}_{\text{full}} - \gamma_0)$, and most sensible model averaging schemes, including FIC and weighted FIC, will give weights close to 1 for the wide model. As e.g. Johnson hints at, other techniques might be needed to better assess behaviour and properties of model average methods in clearly non-contiguous situations, i.e. when γ is far from γ_0 .

We have seen that the $O(1/\sqrt{n})$ framework is canonical inside general parametric models with independence. This has to do with information increasing linearly with sample size divided by model complexity and variances being proportional to inverse information. Ishwaran and Rao mention Breiman's bagging, which indeed may be viewed in terms of model averaging. Some of the calculations in Section 2 of Bühlmann and Yu (2002) may be seen as special cases of our general FMA theory. Their paper shows that when averaging takes place over a large number of stumps, then (su)bagging is best analysed inside a $O(1/n^{1/3})$ framework. A similar comment applies to some of the goodness-of-fit tests of Claeskens and Hjort (2003), where large classes of alternative models are being searched through.

2. Two uses of regression models

Regression analyses have different aims on different occasions, and even the same data set may be analysed with different goals in mind. We have primarily taken the view that what matters most is quality of predictors and precision of focus estimators. Ishwaran and Rao, in contrast, equate the subset selection regression problem with finding the exact subset of non-zero elements among a vector of coefficients $(\beta_1, \ldots, \beta_K)^t$. As a result they partly criticise our methods for not being optimal for a task they were not set out to perform.

For many applications one would not care much if say $\beta_7 = 0.01$ rather than being exactly zero, and the additional estimation noise caused by including $\hat{\beta}_7$ in the predictor formulae might worsen rather than enhance the precision. Ishwaran and Rao appear to say that it is the duty of any subset selection method to strive for inclusion also of such a small β_7 .

There are of course situations where detecting the non-zero-ness of certain parameters is the main goal of an analysis. This could be a β_j coefficient in a linear regression setting, for example. Our theory works for such a focus parameter too, since it may be expressed as $\beta_j = \mathrm{E}(Y \mid x + e_j, u) - \mathrm{E}(Y \mid x, u)$, with e_j the jth unit vector. In an effort towards making the world slightly less unfair, Hjort (1994a) collected and analysed data from world championships in sprint speedskating, attention focussed on the average difference d between 500 m results reached using last inner track vs. last outer track. The analysis essentially employed a bivariate mixed effects regression model with seven parameters per championship (and it was necessary to fit the full seven-parameter model to make inference for d). Only one parameter mattered to the delegates from 37 nations at the 1994 general

assembly of the International Skating Union. They had to assess the potential non-zeroness of d and its implications, and actually needed to vote for or against the significance of the point estimate (which was $\hat{d}=0.06$ seconds). The Olympic rules for sprint speedskating were in fact changed as a consequence of the statistical analysis; from Nagano 1998 onwards the athletes are forced to skate the 500 m distance twice. See also Hjort and Rosa (1999). This is an example of a sharply defined focus parameter where tools of FMA and FIC might be used.

We disagree with the way Ishwaran and Rao interpret the scope of our machinery for subset selection problems in regression, at the end of their Section 1. The statistician is at the outset required to classify some parameters (say β_j s) as 'protected' and other parameters (say γ_j s) as 'uncertain'; our methods are then geared towards finding the best subsets of γ_j s to include, or to be averaged over. Our methods are certainly not 'restricted to coefficients known to be zero', as Ishwaran and Rao charge in their point (a). First, the methods are well-defined and can be applied regardless of the sizes of the γ_j s. This is also a reply to a comment by Raftery and Zheng, that our (1.1) is 'required by FMA'; FMA methods give algorithms that may be put to work regardless of (1.1) type assumptions. Second, even though the mathematical results we have provided about various methods have utilised the $\gamma_j = \gamma_{0,j} + \delta_j / \sqrt{n}$ framework, the δ_j s may be big in size, as also commented on above. Third, most sensible selection methods or averaging methods will pick out the widest model, in cases where the γ_j s really are far from zero.

Regarding their point (b) (end of Section 1), it is fair to say that statistical modelling is and remains an art demanding skill and experience for its perfect execution, even with the advent of further tools for automatisation and diagnostics. The previous argument indicates that it may be rather harmless if a statistician labels a parameter a ' γ ' when it should rather have been a ' β ' (provided the selection or averaging scheme is among the decently robust ones, with low max risk, see FMA's Section 7); this also serves to counter their point (b). A similar comment applies to Raftery and Zheng's reservations (Section 4), having to do with situations where 'the coefficients for some nuisance variables are substantial, and those for others are small'. In such cases the crafty modeller should take this on board, redesigning nuisance variable coefficients as protected.

In their Sections 1 and 2, Ishwaran and Rao argue that in most regression setups, the γ_0 associated with uncertain (or non-protected) variables must be zero. This is fine, is not surprising, and does not contradict our machinery or methodology. Our theory does allow $\gamma_0 \neq 0$ too, but this would here correspond to known trends, which may be removed from the regression equation. We note that the FIC paper has several examples where the canonical γ_0 is non-zero.

3. Estimating model order

In some settings there is a natural order of complexity among candidate models, as with e.g. polynomial regression. Ishwaran and Rao (Section 3) study the problem of estimating

the actual underlying order of the true model, that is, the unknown number k_0 where the coefficient vector is $(\beta_1, \ldots, \beta_{k_0}, 0, \ldots, 0)$ and the first k_0 are strictly non-zero. As mentioned above, in many situations estimating k_0 might not be a vital issue. Their Theorem 1 contrasts backwards and forwards selection schemes under some conditions, and is of interest. We believe their theorem should and can be extended to more general settings, however.

We do think the assumption about finiteness of fourth moments may be softened, although this is not crucial. The primary problem we see is their assumption that $\Sigma_n = n^{-1}X^tX = n^{-1}\sum_{i=1}^n x_ix_i^t$ must equal the identity matrix I; this appears too restrictive. One may transform a regression model to achieve such orthogonality, but this would typically inflict a different ordering of new coefficients, losing the original motivation of nested-ness. This makes it difficult to keep track of the original k_0 . To avoid the problem one may keep the original model, and consequently keep the k_0 as defined by the untransformed β vector, but accept the weaker assumption that Σ_n tends to a general positive definite Q.

Under weak Lindeberg type conditions, see e.g. Hjort and Pollard (1994) (in particular, it does not appear necessary to assume finite fourth moments), we then have $\sqrt{n}(\hat{\beta}-\beta) \to_d N_K(0,\sigma^2Q)$, with consequent $\sqrt{n}(\hat{\beta}_k-\beta_k) \to_d N(0,\sigma_k^2)$, say, where $\sigma_k = \sigma(Q_{k,k})^{1/2}$. Thus there is simultaneous convergence $Z_{k,n} = \sqrt{n}\hat{\beta}_k/\hat{\sigma}_k \to_d Z_k$, say, for $k \geq k_0+1$, where these are standard normals with correlations inherited from Q. Also, $|Z_{k,n}|$ flees to infinity in probability for $k \leq k_0$. Defining the backwards and forwards model order estimates as in Ishwaran and Rao, one may now show that $\hat{k}_B \to_d k_B$ and $\hat{k}_F \to_d k_F$, where

$$\Pr\{k_B = k\} = \begin{cases} 0 & \text{for } k \le k_0 - 1, \\ \Pr\{Z_{k_0+1} \in J_{k_0+1}, \dots, Z_K \in J_K\} & \text{for } k = k_0, \\ \Pr\{Z_k \notin J_k, Z_{k+1} \in J_{k+1}, \dots, Z_K \in J_K\} & \text{for } k \ge k_0 + 1 \end{cases}$$

and

$$\Pr\{k_F = k\} = \begin{cases} 0 & \text{for } k \le k_0 - 1, \\ \Pr\{Z_{k_0+1} \in J_{k_0+1}\} & \text{for } k = k_0, \\ \Pr\{Z_{k+1} \in J_{k+1}, Z_{k_0+1} \notin J_{k_0+1}, \dots, Z_k \notin J_k\} & \text{for } k \ge k_0 + 1. \end{cases}$$

Here $J_k = (-z_{\alpha_k/2}, z_{\alpha_k/2})$ is the acceptance interval for $Z_{k,n}$, with limit probability $1 - \alpha_k$. Ishwaran and Rao's Theorem 1 corresponds to the case of a diagonal Q matrix, where the Z_k s become independent.

In practice the $|Z_{k,n}|$ s for $k \leq k_0$ have not quite had time to flee to infinity, for finite n, as $Z_{k,n}$ has mean value about $\sqrt{n}\beta_k/\sigma_k$. The approximations afforded by the limit theorem above are easily too crude, particularly when β_k s are small. It is again natural to use the local neighbourhood parametrisation, with say $\beta_k = \delta_k/\sqrt{n}$. The limit distributions for \hat{k}_B and \hat{k}_F may be derived. There will in particular be positive probabilities for values $k \leq k_0 - 1$. One finds in fact

$$\Pr\{k_B = k\} = \Pr\left\{\frac{\delta_k}{\sigma_k} + Z_k \notin J_k, \frac{\delta_{k+1}}{\sigma_{k+1}} + Z_{k+1} \in J_{k+1}, \dots, \frac{\delta_K}{\sigma_K} + Z_K \in J_K\right\}$$

for k = 1, ..., K, where $\delta_k = 0$ for $k \ge k_0 + 1$, and where $Z_1, ..., Z_K$ are standard normals with correlations coming from Q. There is a corresponding result for k_F . This creates a different picture than with Ishwaran and Rao's Figure 1, which has been produced under conditions corresponding to having $|\delta_k|$ of infinite size for $k \le k_0$ (and Q diagonal).

Ishwaran and Rao comment that the model order parameter k_0 is not a smooth function of β , and as such falls outside the standard regularity conditions used in our FMA and FIC papers. The resulting predictors, say $\hat{\mu}_B = x^t \hat{\beta}_B$ and $\hat{\mu}_F = x^t \hat{\beta}_F$ for given covariate position x, are however amenable to our methods, viewed as estimators of $\mu = x^t \beta$. The backwards and forwards predictors are model average methods and can be analysed using the FMA methodology. Limit distributions are non-linear mixtures of biased normals, and their performances may in particular be compared to that of AIC and FIC, as per Section 7 in the FMA paper. We also note that the arguments and results alluded to here should generalise without serious difficulties to e.g. generalised linear models.

Ishwaran and Rao 'have always wondered about' whether it is better to use forward or backward stepwise regression. They might perhaps be encouraged to continue their fruitful wondering. Even in cases when \hat{k}_F is more successful than \hat{k}_B as an estimator of k_0 (where, as we argue above, the analysis and conclusion is less clear-cut than what it appears to be in their discussion), the backwards performance would still be better than forwards performance for predicting $x^{\dagger}\beta$, in significant portions of the parameter space.

We use this opportunity to nod in agreement to comments made by Shen and Dougherty (Section 3), that it is very useful when the list of candidate models can be restricted a priori, for both FIC and FMA. In situations with a nested sequence of models, as above, this means reducing the number of candidates from 2^q to q+1. On the other hand the list should be broad enough to reflect real modelling information, as viewed in conjunction with focus parameters. One possibility for shortening the queue of suitors is via suitable thresholding and re-weighting, e.g. including only the ten most promising models as monitored by the FIC scores, or by the posterior probabilities inside a BMA setup. Our FMA theory continues to be applicable also for such strategies.

4. FMA versus BMA

Raftery and Zheng come dressed as BMA's witnesses and deliver a strong case. In their earnest zeal they perhaps inadvertently risk classifying or portraying our FMA work as being anti-Bayesian, in spirit, intent or result. That would be a case of incorrect classification. Our FMA bag comprises not only the compromise estimators of FMA's Section 4, but also averages of the generalised ridge estimators developed in Section 8, and these again are close relatives to BMA methods, as explained in Section 9. When developing our FMA methodology our points of motivation indeed included our wish to understand better the behaviour of BMA strategies.

Realising that both 'BMA' and 'FMA' are big bags of methods, then, it is a little oversuggestive when Raftery and Zheng say that 'BMA [was] generally found to have better performance' and that 'FMA itself does not appear to yield optimal methods'. Some BMA regimes are better than others, and some FMA schemes have optimality properties. One may e.g. work with model selection schemes that post model selection use estimators that are minimax over say $\delta^t K^{-1} \delta \leq c$ type regions, using for this step methods similar to Blaker's (2000). Also, as mentioned above, some of the generalised ridge versions of FMA correspond (to the first order) to BMA schemes.

Several model average schemes may be added to the annotated list given in FMA's Sections 5 and 7. Ishwaran and Rao took up backwards and forwards model selection procedures and as we explained above these may be analysed inside the FMA framework; in particular, their behaviour may be analysed using FMA's Theorem 4.1. Johnson might consider having his former life prolongated by revisiting his and his colleagues' robust Bayesian estimation methods, using the FMA apparatus to understand performance.

Cook and Li discuss sliced inverse regression and central subspaces methods. Such methods are geared more towards dimension reduction than selection of subsets, and may be compared to principal components regression (see e.g. Mardia, Kent and Bibby, 1979, Ch. 8) and to partial least squares regression (see e.g. Helland, 1990). With some work we believe versions of these dimension reduction methods may be characterised and analysed as FMA methods. In these situations it would be more natural to compare performances in terms of suitably averaged prediction accuracy, see our next section.

As far as performance is concerned, Raftery and Zheng are perhaps right to arrest us for not paying enough attention to the existing BMA literature. They provide references to and give a summary of three main strands of results: general Bayes theory (along with studies of robustness to prior specifications); simulations; and cross-validation type predictive performance. See also the concise and useful discussion in Clyde and George (2003). What we intended to point to in our introduction to the FMA paper was the surprising lack in the literature of what one may think of as 'the fourth strand of results', namely limit distribution statements. In mathematical statistics we are not quite satisfied with simulations and cross validation and indications of good performance; we need precise limit distribution results. This is not only dictated by tradition and aesthetics, but gives practical mathematics, providing good approximations for precision measures as well as a tool for comparing performances, say of different BMA schemes. What is logistic regression without results about limiting behaviour of likelihood methods? What is years of hands-on experiences with averages without the central limit theorem?

Shen and Dougherty stress, along with Johnson and with Raftery and Zheng, like we have done, the necessity of securing a well-defined interpretation of focus parameters (or variables) across models. In our framework this is taken care of via $\mu = \mu(f)$, where f belongs to suitable submodels of the widest $f(y, \theta, \gamma)$ model. This requirement, when boomeranged back to BMA's watchtower, becomes the issue we raise in FMA's Section 1.1, that BMA typically entails mixing together conflicting prior opinions about focus parameters. Our discussants do not take up this point.

5. Average quality of predictors

We appreciate Raftery and Zheng's additional comments to and extended analysis of the low birth weight data set. Our own analysis of this data set was primarily intended as an illustration of the developed methods, as opposed to a full scientific report on low birth weights. This was also why we chose somewhat simple parameters as foci. Let us write these as $p_1 = p(z_1)$, $p_2 = p(z_2)$ and $\rho = p_2/p_1$, where z_1 and z_2 are the average covariate vectors for white and black mothers, respectively. We may agree with Johnson and with Raftery and Zheng that there are yet other parameters to focus on, with perhaps higher socio-biological relevance; again, our parameters were chosen for illustration and simplicity. We still believe that ρ has some merit, though. A litmus test for 'being of interest' might be whether one can imagine a newspaper or magazine publishing a story about a finding concerning the parameter in question; here a news story sentence like 'the average black mother has a 50% greater chance than the average white mother of giving birth to too small children' would appear to pass the test. Of course one should with such a finding attempt to investigate further, including aspects of the covariate distributions.

Comments from Cook and Li as well as from Raftery and Zheng point to the usefulness of developing the FIC and FMA apparatus to assess prediction quality when averaged in suitable ways, rather than for one focus parameter at a time. We touch on this in the FIC paper's Sections 5.6 and 7.2. Such averaging is particularly natural in regression models, where focus might be on the behaviour of say $\hat{\mu}(x,u)$ for a regression surface $\mu(x,u)$ for particular sub-regions of u for fixed x, and so on. We note that the theory and arguments also invite suitable weighted generalisations of the AIC.

To indicate how the machinery can be developed further, consider a linear regression setup with $Y_i = x_i^t \beta + u_i^t \gamma + \varepsilon_i$ for i = 1, ..., n, where the ε_i s are i.i.d. with mean zero and standard deviation σ , and where $\gamma = \delta/\sqrt{n}$. The x_i s are protected while elements of the u_i s may or may not be taken into the finally selected model. We study the average weighted prediction error

$$\mathcal{E}_n = n^{-1} \sum_{i=1}^n (\hat{\xi}_i - \xi_i)^2 w(x_i, u_i), \tag{5.1}$$

where $\xi_i = \mathrm{E}(Y | x_i, u_i) = x_i^{\mathrm{t}} \beta + u_i^{\mathrm{t}} \gamma$ and $\widehat{\xi}_i$ an estimator thereof, with w(x, u) a suitable weight function. We shall see that $n\mathcal{E}_n$ has a limit distribution, under reasonable conditions.

Let

$$\Sigma_n = n^{-1} \sum_{i=1}^n \begin{pmatrix} x_i \\ u_i \end{pmatrix} \begin{pmatrix} x_i \\ u_i \end{pmatrix}^{\mathsf{t}} = \begin{pmatrix} \Sigma_{n,00} & \Sigma_{n,01} \\ \Sigma_{n,10} & \Sigma_{n,11} \end{pmatrix},$$

of size $(p+q) \times (p+q)$, assumed to be of full rank. Its inverse Σ_n^{-1} has blocks denoted Σ_n^{ij} , and similarly for the smaller $(p+|S|) \times (p+|S|)$ matrix $\Sigma_{n,S}$ with inverse $\Sigma_{n,S}^{-1}$. We assume that $\Sigma_n \to \Sigma$ as n increases, also of full rank. We let $L_n = \Sigma_n^{11}$, along with

 $L_{n,S} = (\pi_S L_n^{-1} \pi_S^{\rm t})^{-1}$ and $H_{n,S} = L_n^{-1/2} \pi_S^{\rm t} L_{n,S} \pi_S L_n^{-1/2}$. The matrices $L_n, L_{n,S}, H_{n,S}$ have limits L, L_S, H_S .

For the S subset estimator

$$\begin{pmatrix} \widehat{\beta}_S \\ \widehat{\gamma}_S \end{pmatrix} = \sum_{n,S}^{-1} n^{-1} \sum_{i=1}^n \begin{pmatrix} x_i \\ u_{i,S} \end{pmatrix} Y_i,$$

we have

$$\sqrt{n} \begin{pmatrix} \widehat{\beta}_S - \beta \\ \widehat{\gamma}_S \end{pmatrix} \to_d \begin{pmatrix} C_S \\ D_S \end{pmatrix} = \Sigma_S^{-1} \begin{pmatrix} \Sigma_{01} \delta + M \\ \Sigma_{11} \delta + N_S \end{pmatrix},$$

where $(M, N) \sim N_{p+q}(0, \sigma^2 \Sigma)$ and $N_S = \pi_S N$. We may write

$$\mathcal{E}_{n} = n^{-1} \sum_{i=1}^{n} (x_{i}^{t} \widehat{\beta}_{S} + u_{i,S}^{t} \widehat{\gamma}_{S} - x_{i}^{t} \beta - u_{i}^{t} \gamma)^{2} w(x_{i}, u_{i})$$

$$= n^{-1} \sum_{i=1}^{n} \left[\binom{x_{i}}{u_{i}}^{t} \binom{\widehat{\beta}_{S} - \beta}{\binom{\widehat{\gamma}_{S} - \gamma_{S}}{-\gamma_{S^{c}}}} \right]^{2} w(x_{i}, u_{i})$$

$$= \binom{\widehat{\beta}_{S} - \beta}{\widehat{\gamma}_{S} - \gamma_{S}}^{t} \Omega_{n} \binom{\widehat{\beta}_{S} - \beta}{\widehat{\gamma}_{S} - \gamma_{S}}^{t},$$

$$-\gamma_{S^{c}}$$

where Ω_n is the w-weighted version of Σ_n above. Thus, if only $\Omega_n \to_p \Omega$,

$$n\mathcal{E}_n \to_d \mathcal{E} = \begin{pmatrix} C_S \\ D_S - \delta_S \\ -\delta_{S^c} \end{pmatrix}^{\mathrm{t}} \Omega \begin{pmatrix} C_S \\ D_S - \delta_S \\ -\delta_{S^c} \end{pmatrix}.$$

Expressions for the mean of \mathcal{E} may be found using tools of the FIC paper.

When w = 1 in (5.1) we have $\Omega_n = \Sigma_n$ and a corresponding simplification for \mathcal{E} . The limiting risk using S can be shown to become

$$E(\mathcal{E}) = (p + |S|)\sigma^2 + \delta^t L^{-1/2} (I - H_S) L^{-1/2} \delta,$$

using arguments as in FIC's Section 7.2. Let $D_n = \sqrt{n} \hat{\gamma}_{\text{full}}$, which goes to a $N_q(\delta, \sigma^2 L)$. An unbiased risk estimator is

$$\widehat{\text{risk}}_S = (p + |S|)\widehat{\sigma}^2 + \text{Tr}\left[L^{-1/2}(I - H_S)L^{-1/2}(D_n D_n^{\text{t}} - \widehat{\sigma}^2 L)\right]
= (p - q + 2|S|)\widehat{\sigma}^2 + D_n^{\text{t}} L_n^{-1} D_n - D_n^{\text{t}} L_n^{-1/2} H_{n,S} L_n^{-1/2} D_n,$$

where $\hat{\sigma}^2$ is the usual unbiased estimator of variance, using the full model. This leads to the following selection criterion: choose the subset with smallest value of

ave-FIC(S) =
$$\widehat{\sigma}^2 \{ 2|S| + n\widehat{\phi}^{t}(I - H_{n,S})\widehat{\phi}/\widehat{\sigma}^2 \},$$

where $\hat{\phi} = L_n^{-1/2} \hat{\gamma}_{\text{full}}$. This appears to be related to both Mallows' C_p as well as to Cook and Li's eq. (2) (worked out there for the case of p = 1, $x_i = 1$, and $\sum_{i=1}^n u_i = 0$). Note that $\sqrt{n}\hat{\phi}/\hat{\sigma} \to_d N_q(\phi, I)$, where $\phi = L^{-1/2}\delta$. For other extensions of the Mallows criterion, and theory, see Birgé and Massart (2001).

We note that the above ideas and arguments may be used to find precise limit distributions for average prediction error variables of the type $\sum_{i=1}^{n} {\{\hat{\mu}(x_i, u_i) - \mu(x_i, u_i)\}^2 w(x_i, u_i)}$, in quite general regression models and for quite general model average estimators. Such results may in particular be used for model and subset selection purposes. One is quite free to choose weight schemes appropriate for the purpose. If one wishes to assess predictor quality for a fixed x_0 , when averaged over u, one may insert $w(x_0, u)$ proportional to an estimate of the conditional density of u given x_0 . This might be a multinormal density, or a kernel-smooth over a window around x_0 .

We think that developments as above might lead to useful 'focussed regression diagnostics' of different types. The comments of Cook and Li also point in such directions.

6. Second order corrections

In our papers we have determined the limit distribution of $\Lambda_{n,S} = \sqrt{n}(\hat{\mu}_S - \mu_{\text{true}})$ (as well as for more general estimators, like the post model selection estimator). This gives the approximation

$$\operatorname{risk}_{n}(S, \delta) = n \operatorname{E}(\widehat{\mu}_{S} - \mu_{\text{true}})^{2} \doteq \operatorname{E}\Lambda_{S}^{2}, \tag{6.1}$$

where Λ_S is the limit variable. In FMA's Section 10.7 and FIC's Section 7.6 we mentioned the potential for suitable finite-sample corrections to the first order results of type (6.1). We are glad that Tsai has taken up this challenge, providing what he terms 'improved' and 'corrected' versions of the FIC.

The exact bias and variance of $\hat{\mu}_S$ would often depend in complicated ways on the model and sample size; see e.g. Dukić and Peña (2003) for finite-sample analysis of some particular post-selection estimators in Gaußian models. Sometimes expansions for these might be worked out, however. Suppose in general terms that

$$E\Lambda_{n,S} = B_1(S,\delta) + B_2(S,\delta)/\sqrt{n} + B_3(S,\delta)/n + o(1/n),$$

$$Var \Lambda_{n,S} = V_1(S) + V_2(S,\delta)/n + o(1/n),$$

for suitable coefficients. Lemma 3.3 in the FMA paper gives in fact expressions for the leading terms $B_1(S, \delta)$ and $V_1(S)$, and hence for the leading term $E\Lambda_S^2 = B_1(S, \delta)^2 + V_1(S)$ in (6.1). In then follows that

This shows that the second order term to catch (and estimate) is $2B_1(S,\delta)B_2(S,\delta)/\sqrt{n}$. This necessitates finding an expression for $B_2(S,\delta)$.

Following Tsai, this requires taking the delta method one step further, using a 2nd order Taylor expansion. We do this in a somewhat different way. Starting with

$$\widehat{\mu}_S - \mu_{\text{true}} = \mu(\widehat{\phi}_S, \gamma_{0,S^c}) - \mu(\phi_{0,S}, \gamma_{0,S^c}) + \mu(\theta_0, \gamma_0) - \mu(\theta_0, \gamma_0 + \delta/\sqrt{n}),$$

we may split $\Lambda_{n,S}$ into two parts, with leading terms

$$(\partial \mu/\partial \phi)^{t} \sqrt{n} (\widehat{\phi}_{S} - \phi_{0,S}) + \frac{1}{2} \sqrt{n} (\widehat{\phi}_{S} - \phi_{0,S})^{t} \mu_{11,S} (\widehat{\phi}_{S} - \phi_{0,S})$$

and

$$-(\partial \mu/\partial \gamma)^{t}\delta - \frac{1}{2}\delta^{t}\mu_{22}\delta/\sqrt{n}.$$

In our notation, $\mu_{11,S}$ is the $(p + |S|) \times (p + |S|)$ matrix of 2nd order derivatives of $\mu(\theta, \gamma_S, \gamma_{0,S^c})$ w.r.t. (θ_S, γ_S) , while μ_{22} is the $q \times q$ matrix of 2nd order derivatives of $\mu(\theta, \gamma)$ w.r.t. γ . These derivatives are evaluated under the narrow model (θ_0, γ_0) .

To come further we need

$$\mathbb{E}\sqrt{n}(\widehat{\phi}_S - \phi_{0,S}) = J_S^{-1} \begin{pmatrix} J_{01} \\ \pi_S J_{11} \end{pmatrix} \delta + m_S(\delta)/\sqrt{n} + n_S(\delta)/n + \cdots,$$

with suitable (but often cumbersome) expressions for $m_S(\delta)$ and $n_S(\delta)$ obtainable from work touched on by Tsai; see also Barndorff-Nielsen and Cox (1994, Chapters 5–6). This leads to

$$B_2(S,\delta) = (\partial \mu / \partial \phi_S)^{t} m_S(\delta) + \frac{1}{2} \text{Tr}(\mu_{11,S} J_S^{-1}) - \frac{1}{2} \delta^{t} \mu_{22} \delta.$$

To summarise,

$$\operatorname{risk}_{n}(S,\delta) = \operatorname{E}\Lambda_{S}^{2} + 2B_{1}(S,\delta)B_{2}(S,\delta)/\sqrt{n} + o(1/\sqrt{n})$$
(6.2)

provides a 2nd order corrected version of (6.1).

The treatment above is related to but not fully equivalent to what Tsai does. He studies nonlinearity aspects in his Section 2 and bias of likelihood estimators in Section 3. It appears to us, from the arguments above, that it is necessary to combine both these 2nd order aspects. If not one risks catching one or two of the terms making up $B_2(S, \delta)$, but not all three, and a partial reparation might be worse than no reparation.

We would perhaps hesitate to affix the labels 'improved' and 'corrected' too firmly on Tsai's modified FICs. It is clear from the above that there are several possibilities for such 2nd order approximations to the mean squared error of estimators. Also, one needs indirectly or directly to estimate $B_1(S,\delta)B_2(S,\delta)$ from data, where there are several paths to follow, e.g. regarding wide versus narrow estimation of partial derivatives. Furthermore, this estimation step might cause additional variability that might take away the intended benefit. Such phenomena are well-known in mathematical statistics. A 2nd order Edgeworth expansion might not be a genuine improvement over a 1st order Edgeworth

expansion, for example, or perhaps there is improvement only for very large sample sizes. All this serves to indicate that further studies are required before a general-purpose 2nd order FIC can be established.

We note that Tsai's work, and presumably also the development above, is relevant also when it comes to assessing behaviour of model average estimators.

7. Estimators from other likelihoods

Tsai points out that γ parameters sometimes are in focus, and we agree. Our FIC and FMA apparatus is nicely able to handle this, since it covers all smooth $\mu(\theta, \gamma)$ parameters; Tsai appears to claim otherwise. With focus on γ_j we find $\omega = -e_j$, with $e_j = (0, \dots, 1, \dots, 0)^t$ being the jth unit vector. We are free to form general model average estimators $\hat{\gamma}_j = \sum_S c(S \mid D_n) \hat{\gamma}_{j,S}$, where incidentally terms with S not touching j will be equal to zero. Using FMA's Theorem 4.1, we find

$$\sqrt{n}\{\widehat{\gamma}_j - (\gamma_{0,j} + \delta_j/\sqrt{n})\} \to_d \widehat{\delta}_j(D) - \delta_j \text{ for } j = 1, \dots, q,$$

and so on. The FIC can also be applied, and one may study simultaneous estimation of the full γ vector. It is also natural to include the goodness-of-fit measure

$$\widehat{\delta}^{t}\widehat{K}^{-1}\widehat{\delta} = n(\widehat{\gamma} - \gamma_0)^{t}\widehat{K}^{-1}(\widehat{\gamma} - \gamma_0)$$

in the data analysis. It is a $\chi_q^2(\delta^t K^{-1}\delta)$ in the limit.

There might be situations where the ordinary likelihood apparatus cannot be used, or can be expected to perform poorly, and where variations like profile likelihoods, empirical likelihoods and quasi-likelihoods may be helpful. This would call for extensions of our work. We do not think, however, that profiling is necessary, or that it leads to new results, inside our parametric $f(y, \theta, \gamma)$ framework. We are therefore puzzled with Tsai's elaborations in this regard; under weak conditions the S-model profile likelihood estimator of μ will simply be our old maximum likelihood estimator $\hat{\mu}_S$. Tsai's intricate definition of a new 'random parameter' $\mu_{\text{prof,true}}$ does not correspond to our more naturally defined μ_{true} .

In Hjort and Claeskens (2003) we report on extensions of our FIC and FMA work, for model selection and model averaging inside the semiparametric Cox regression model. Focus parameters could take the form $\mu(\beta, H, z)$, involving the parametric as well as the nonparametric part of the model, as with the median time to survival for a patient with given covariates z. Our existing theory will be seen to go through without essential modifications as long as μ is a function of β and covariates only, whereas such modifications are called for when it also involves H. Similarly, extensions may be envisaged for use in spatial models with covariates, inside particular formats of parameter estimation; we in particular have in mind the pseudo-likelihood method of Besag (1974, 1977) for Markov random fields; the quasi-likelihood of Hjort and Omre (1994, Section 3) for spatial correlation models; and various methods for observed and aggregated point processes reviewed in Richardson (2003).

8. Supplementary comments

8.1. When p and q become big. Our methodology has been developed under the classic asymptotics scenario where the number of parameters stays bounded when the sample size increases. Shen and Dougherty point out (in their Section 4) that the results might need modifications to apply when p + q is big, as will happen in many potential applications. We agree. This needs further mathematical developments. We do believe, however, that our asymptotics results will continue to provide adequate descriptions and approximations even when p + q grows with n, but slowly enough to have $p + q = o(\sqrt{n})$. Establishing such results would need further work, but might use methods similar to those used in e.g. Portnoy (1988).

We use the opportunity to opine that if p + q becomes too big, it should be reduced. If one has 1,000 covariates per patient, one does good to compress and synthesise these, using substantive prior knowledge along with statistical techniques, before throwing the data set to a regression selector or averager. Also, methods like principal components and partial least squares regression might easily perform better than subset finding schemes.

8.2. Loss functions and aspects of costs. Cook and Li point out that using limiting mean squared error will not always suffice for making the relevant conclusions, regarding e.g. model selection; see also comments by Shen and Dougherty. In some cases there is a cost k(S) associated with observing future data for regressors in index set S. With loss functions that suitably combine precision with cost, like $n(\hat{\mu} - \mu)^2 + \alpha k(S)$, we would have

$$E loss_n(S) = n E(\widehat{\mu}_S - \mu_{true})^2 + \alpha k(S) \rightarrow E\Lambda_S^2 + \alpha k(S).$$

This might be estimated using a slight extension of the FIC, after which an optimal subset may be extracted.

We have favoured limiting mean squared error as performance criterion, but might also have worked e.g. with L_1 loss, leading however to more complicated expressions for and estimators for $E|\Lambda_S|$ and so on.

8.3. Handling corner parameters. Shen and Dougherty discuss a general four-parameter model where rate measurements are of the form $V(x_1, x_2 | \beta_1, \beta_2, \beta_3, \beta_4)$ plus observation error, with

$$V = \frac{\beta_1 x_1}{\beta_2 (1 + \beta_3 x_2) + x_1 (1 + \beta_4 x_2)}.$$

The case of $(\beta_3, \beta_4) = (0,0)$ is the so-called Victor-Michaelis-Menten model for enzyme mediated reactions. In fisheries research it is also well known as the spawner-recruit model, dating back to an influential paper of Beverton and Holt (1957); see Gavaris and Ianelli (2002) and the engaging discussion in Smith (1994, Ch. 8). Shen and Dougherty discuss aspects of modelling the V, in particular looking at the four possibilities in-in, in-out, out-in, out-out for (β_3, β_4) . This cannot be studied well without a clearer understanding

of the error structure involved. That this is non-trivial and vital, and will vary widely with context, is clear from Ruppert, Cressie and Carroll (1989). Shen and Dougherty allude to pre-test methods, which decide on in- or exclusion of β_3 and β_4 on the basis of tests for their presence. We note that such schemes are again model average methods, and fall inside our developed theory.

There might sometimes be situations where it is known a priori that e.g. $\beta_3 \geq 0$, $\beta_4 \geq 0$. The theory we have developed presupposes that (β_3, β_4) is an inner point of the parameter space. To handle 'corner problems', like here, one needs somewhat more intricate methods, which would depend more on the specifics of the problem. See Hjort (1994b) for one such example, concerned with compromise estimators when the t family is used as an extension of the normal in e.g. regression settings. Similar problems emerge in models with variance components. Methods of Vu and Zhou (1997) appear relevant when attempting to generalise our results to corner parameters.

An opinion perhaps too rarely expressed, which we share, is that statisticians should be more eager to help develop good non-linear regression models, as here. The comfort and ease with which we reach moderately adequate approximations and inference precision using the flexible machinery of (generalised) linear models may sometimes take the edge out of our professional modelling creativity.

8.4. Non-nested models. We have for the most part stayed inside a framework where the biggest model is thought to be correct. Cook and Li mention the problem of non-nested models. The simplest answer, perhaps, from a principled point of view, is that one might search for a bigger model formulation that encompasses both. Consider estimating the median, for example, including under view both the gamma and the log-normal models. One may then work with estimators of the type $\hat{\mu} = W \hat{\mu}_{\text{gam}} + (1 - W) \hat{\mu}_{\text{logn}}$, with weights somehow dictated by data, e.g. via goodness-of-fit measures, or via closeness of the two estimates involved to the nonparametric $\hat{\mu}_{\text{nonpm}}$, i.e. the sample median. Behaviour and performance may be studied using our methods.

There are examples in science where non-nested and somehow conflicting statistical theories are not easily resolved, of course. A controversy of some fame inside fisheries research, and that has perhaps not yet been solved to satisfaction despite having been pondered over for about a hundred years, is the Dannevig vs. Hjort case. It is concerned with models for spawning, recruitment, migration and development of fish populations. Dannevig essentially believed in a deterministic relationship between the number of recruits and the number of yolk-sack codfish larvae, whereas Hjort argued that it is the environmental conditions during the critical phases of development that play the more important roles. He was able to develop year-class assessment methods, collect relevant data and utilise actuarial mathematical methods of the time to substantiate and refine his theories; cf. e.g. Hjort (1914). See http://www.math.ntnu.no/~ingeol/bemata/, where a study of structured stochastic models has been launched, involving computer-intensive

inference in biological marine systems, and the interesting discussion in Smith (1994) and Secor (2002). This may be an example where model averaging might be useful, in a non-nested setup, mixing predictions of e.g. next season's abundance (perhaps as a function of quota thresholds) using elements of both scientific models.

8.5. Interpreting FIC numbers. The FIC scores have been developed as estimates of n times mean squared error of subset estimators (modulo an additive constant), and as such depend on the scale used. They may be made scale-independent via say

$$\operatorname{FIC}^*(S) = \widehat{\operatorname{FIC}}(S)/\widehat{\omega}^{\operatorname{t}}\widehat{K}\widehat{\omega},$$

as in FMA's Section 5.3. This would make comparison and interpretation easier across applications. We would in particular have

$$\operatorname{FIC}^*(\operatorname{full}) = 2$$
 and $\operatorname{FIC}^*(\operatorname{narrow}) = n\{\widehat{\omega}^{\operatorname{t}}(\widehat{\gamma}_{\operatorname{full}} - \gamma_0)\}^2 / \widehat{\omega}^{\operatorname{t}} \widehat{K} \widehat{\omega}.$

8.6. When is ω equal to zero? We have seen that the behaviour of model average estimators is critically determined by $\omega = J_{10}J_{00}^{-1}\frac{\partial\mu}{\partial\theta} - \frac{\partial\mu}{\partial\gamma}$. In particular, if $\omega = 0$, then all subset and model average estimators are asymptotically equivalent to the narrow model estimator; $\sqrt{n}(\hat{\mu} - \mu_{\text{true}}) \to_d N(0, \tau_0^2)$ for all reasonable competitors. The typical situation leading to $\omega = 0$ is when the parameter does not depend on γ and in addition θ and γ are orthogonal parameters, in the sense that their full model estimators are independent in the limit, i.e. $J_{01} = 0$. Johnson asks whether ω may be zero also in other situations. Here is one example, in the framework of the exponential-within-Weibull model of FMA's Section 4.4. Assume we wish to estimate the α -quantile $\mu = \nu^{1/\gamma}/\theta$, where $\nu = -\log(1-\alpha)$. Then calculations give $\omega = (\nu/\theta) \{-(1-r) + \log \nu\}$. For estimating the $\alpha = 0.7826$ -quantile, therefore, ω happens to be equal to zero.

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