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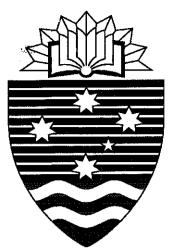
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BOND UNIVERSITY School of Business

DISCUSSION PAPERS

"Diagnostic Testing and Sensitivity Analysis in SAM Construction

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DISCUSSION PAPER NO 51

January 1994

University Drive,

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AUSTRALIA

Diagnostic Testing and Sensitivity Analysis in SAM Construction

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1. Introduction

One of the major issues when constructing a Social Accounting Matrix is that of checking the initial cell estimates provided by the statistician. Statisticians typically protest strongly on this issue, maintaining that all relevant information has been incorporated in the cell estimates and that no further improvement is possible. If the cell estimates are also expected to satisfy various identities then, one obvious check (and adjustment) is to use the identities in testing and balancing modes. Such identities have often been used for balancing - where the adjustment typically depends on producing new estimates which are as close to the initial estimates as possible within the confines of a quadratic norm where the weights are the reliabilities of the initial estimates. Statisticians seem to view such mechanical adjustments with as much enthusiasm as the Royal Navy first viewed the advent of steam. The adjusted estimates satisfy adding up conditions which are accounting identities and the burden of the adjustment falls on the least reliable of the initial estimates.

The dual of constrained estimation is the process of hypothesis testing. The method of classical statistics is to set up a null hypothesis, embodying certain prior information, to test the hypothesis and, if it is not rejected, to incorporate that prior information in the estimation process. The end result is enhanced efficiency. The process of testing is important because hypotheses are often rejected and theory refined, as a result. The procedure of estimating and balancing a SAM has been discussed elsewhere {see Arkhipoff [1969], Stone [1975], Byron [1977] and van der Ploeg [1982] for examples}. However, the process of testing has had no attention in the literature, despite the fact that ad hoc testing and refinement procedures are often used in SAM construction. This paper explores the issue of testing and stems from an earlier paper by Byron, Crossman, Hurley and Smith [1993] which found that constrained estimation or balancing when accompanied by "ad hoc" testing was invaluable in focussing on erroneous initial estimates of cells in a SAM. That experience suggested it might be worthwhile to develop and formalise statistical tests for SAM construction.

2. The Statistical Model

A SAM is simply a table with row and column adding-up conditions, the estimates of the cells (parameters) in the matrix have different levels of reliability, and the adding up conditions are applicable to the true cell values. The row and column adding-up conditions are valid restrictions (identities) applicable to the true cell

¹ The intellectual interest and financial support of the Queensland Treasury is gratefully acknowledged.

values; because the adding up restrictions are valid, they can be used to improve the initial estimates of the cells. If cell estimates are viewed as a random variables with known probability distributions, then traditional statistical theory is applicable and it is easily shown that the use of true prior information must result in more efficient estimates of those cells.

To illustrate, suppose X is a 4x4 SAM, then there are 7 independent row and column restrictions. Let x=vec(X), where the vectorisation is by column and the 4 row restrictions are stacked first, followed by the column restrictions, then the restriction matrix for Gx = h = 0 has one redundant restriction.

SAM balancing is just a procedure to ensure that a set of estimates is produced which satisfy the adding up restrictions and are as close as possible to the statistician's initial estimates of the cells in the context of a quadratic norm. If the condition that all cell members be non-negative is imposed, SAM balancing is identical to the quadratic transportation problem of operations research.

Let x and x be the initial and constrained estimates of the cell members respectively and let μ be the true values. The constrained objective function (with the redundant equation of G deleted) is

(1)
$$(\tilde{x} - \tilde{x})' V^{-1}(\tilde{x} - \tilde{x}) + \lambda' (\tilde{Gx} - h)$$

The first order conditions are

$$(2) \tilde{x} = x - VG' \tilde{\lambda}$$

(3)
$$\lambda = (GVG')^{-1}(Gx - h)$$

(2)
$$\tilde{x} = x - VG' \tilde{\lambda}$$
(3)
$$\tilde{\lambda} = (GVG')^{-1} (Gx - h)$$
(4)
$$\tilde{x} = x - VG' [GVG']^{-1} Gx$$

since h=0.

If the data generation process is that the initial estimates of X are iid with $x = \mu + e$, then Gµ=0, E(e)=0 and GE(x) = 0.

(5)
$$\tilde{x} = (\mu + e) - VG' (GVG')^{-1} G(\mu + e)$$

(6)
$$E(x) = \{I - VG [GVG]^{-1}G\}\mu = \mu$$

Next,

(7)
$$\tilde{x} - \mu = e - VG' (GVG')^{-1} Ge$$

SO

(8)
$$E(\bar{x} - \mu)(\bar{x} - \mu)' = V - VG'(GVG')^{-1}GV$$

which supports the usual result that the constrained estimator is at least as efficient as the initial estimator and because the prior information is correct, \bar{x} is also unbiased.

In the traditional statistical model, the role of hypothesis testing is to test the validity of the restrictions. Here the restrictions are identities and must hold for the population parameters. However, what is at issue is whether or not the

initial estimates of the cells are unbiased. Suppose $x = \mu + \delta + e$ where δ is the bias in the initial (unconstrained) estimates of μ . Henceforth, we refer to this is IEB, initial estimate bias. Clearly, if the initial estimate is biased, the constrained estimates of that cell and all linked cells will be biased.

The objective is to isolate IEB by examining the deviations between the restricted and initial estimates of μ and setting up a test statistic based on their deviations using the normality of e. Since

(9)
$$\tilde{x} - \tilde{x} = -VG' (GVG')^{-1} G(\mu + e)$$

and

(10)
$$E(x-x)(x-x)' = VG'(GVG')^{-1}GE\{(\mu+e)(\mu+e)'\}G(GVG')^{-1}GV$$

(11)
$$\Sigma = E(\tilde{x} - x)(\tilde{x} - x)' = VG'(GVG')^{-1}GV$$

under Ho. Furthermore,

(12)
$$E(x-x) = -VG'(GVG')^{-1}G(\mu + E(e)) = 0.$$

since $G\mu = 0$. However, if any element of \hat{x} is biased, such that $\hat{x} = \mu + \delta + e$ then

(13)
$$E(x-x) = -VG'(GVG')^{-1}G\delta.$$

Under the null hypothesis (that $\delta = 0$) the covariance matrix is defined above and the quadratic form provides a Chi-squared statistic

(14)
$$(x-x)' \Sigma^{-1}(x-x) = x' G' (GVG')^{-1} Gx \sim \chi^{2}_{2n-1}$$

The components of the quadratic form can also be set up as standard normal deviates by dividing the difference in the constrained and initial estimates by their standard

deviations. In other words, the validity of the restrictions Gx can be tested separately, under the assumption that the other restrictions are correct. This is one way to isolate the source of the bias.

The initial estimates of the elements of X can be used to provide a test of IEB. If the initial estimates of the elements of X are unbiased, their expected values will satisfy the row and column adding up conditions. If a single estimate is biased, then it will not satisfy its row and column adding-up restrictions and any test statistics on those two restrictions will isolate the offending initial estimate. For example, the

restrictions in the 4x4 case indicate that any bias in x_{11} will be observed in restriction 1 and restriction 5, any bias in x_{12} will be observed in restriction 2 and restriction 5. and so on.

The statistical model is $x = \mu + e$, each restriction contains estimates of different X terms; these initial estimates are statistically independent and since the true values satisfy the adding-up restrictions, if y = Gx, then E(y) = 0 and Var(y) = GCov(x)G'. Since y is normally distributed, a test statistic based on $\frac{y-0}{\sqrt{Var(y)}}$ will have a standard normal or a central t distribution, under H_O depending on whether Var(y) is known or estimated. The key is that any error in an initial estimate of a cell will show up in two of the test statistics, enabling that cell to be **identified**. This is a decomposed Wald test based on the initial estimates.

An "ad hoc" procedure often used is to test if the restricted estimate is within two standard deviation units of the initial estimate. This procedure uses the initial estimate of the variance to provide the standard deviation. Using (4) and setting the 'ad hoc' test up as a quadratic form it is immediately seen to be a likelihood raio test and by substitution becomes a Wald test, which establishes the legitimacy of the procedure.

(15)
$$(x-x)^{1} V^{-1} (x-x) = x^{1} G' (GVG')^{-1} Gx$$

An LM test is also easily constructed based on the distribution of the quadratic form $\lambda' \Sigma^{-1} \lambda$ under H_0 . Substitution yields

(16) $\lambda' \Sigma^{-1} \lambda = \lambda' (GVG') \lambda = x' G' (GVG')^{-1} G x$

(16)
$$\hat{\lambda}' \Sigma^{-1} \hat{\lambda} = \hat{\lambda}' (GVG') \hat{\lambda} = \hat{x}' G' (GVG')^{-1} G\hat{x}$$

with the usual equivalence between the LM and Wald tests. However, in the classical context, where the restrictions are at fault, if only one restriction is incorrect, a test statistic can be set up based on the individual Lagrangians. This has only been done occasionally in the literature before {see Byron [1972], for an example}. In the present context, single Lagrangians provide tests which can pinpoint the source of the bias. The Lagrangians will have a standard normal or central t distribution depending on whether V is known or estimated.

Any discussion of an estimator is incomplete if its large sample properties are not included. In the present context this poses a difficulty. The initial estimates are an informed guess which are assumed to be unbiased. Any errors are independently, identically (and perhaps normally) distributed. The test statistics are based on linear combinations of the underlying random variables. As the size of the system increases the number of random terms increases by the square, whereas the number of restrictions to be tested increases linearly (n² versus 2n-1). The Wald and LM tests can thus be expected to be well behaved as the size of the system increases because they are linear functions of an increasing number of errors. The Difference test, based on the distribution of (x-x) is also well behaved as the size of the SAM increases

because, following (4), the test is a linear function of an increasing number of errors. The upshot is that the three tests should be well behaved as the size of the system increases; under the null hypothesis they should have the assumed distributions with their hypothetical rejection rates and under the alternative hypothesis they should display reasonable power.

Consider the Lagrangian (3), $\tilde{\lambda} = (GVG')^{-1}(Gx - h)$. If the null hypothesis is true $G\mu = 0$ and $\tilde{\lambda} = (GVG')^{-1}Ge$. However, given G is a matrix whose row vectors consist of unit elements and zeros plim $\frac{Ge}{n} = 0$. It is easily shown that $\lim_{n \to \infty} \frac{GVG'}{n} = K$, where K is a constant matrix, so the estimator of λ is consistent.

To see this, consider the form of GVG' when n=4 and V is diagonal with, for convenience of exposition, all diagonal elements equal. Referring to G on page 2

$$GVG' = \begin{bmatrix} 4v & 0 & 0 & 0 & v & v & v \\ 0 & 4v & 0 & 0 & v & v & v \\ 0 & 0 & 4v & 0 & v & v & v \\ 0 & 0 & 0 & 4v & -v & -v & -v \\ \hline v & v & v & -v & 4v & 0 & 0 \\ v & v & v & -v & 0 & 4v & 0 \\ v & v & v & -v & 0 & 0 & 4v \end{bmatrix}$$

with the notation that the common diagonal elements are v. Since n=4, GVG' when divided by n cannot increase in magnitude as the size of the SAM increases. Hence λ is consistent.

Given this consistency, it appears that test statistics based on a linear combination of the random variables (the Wald, LM and Diff procedures) will be valid whatever the size of the SAM. The large sample properties of test statistics can be described by the behaviour of their confidence intervals and a consistent estimator provides a well behaved test [see Bickel and Docksum, ch.6]. The simulations below offer further evidence of the validity of the test procedures.

3. Empirical Evidence

In what follows an exploratory Monte Carlo experiment is set up with 1000 replications. In the first pass, the estimates of μ are simulated to vary normally around the true μ with a variance consistent with the initial assumptions made. There are 16 parameters to be estimated in the 4x4 case resulting in 8 test statistics corresponding to the row and column restrictions.

The true values of the cell members and their (true) variances are

$$X = \begin{pmatrix} 10 & 2 & 5 & 17 \\ 5 & 2 & 8 & 15 \\ 9 & 3 & 9 & 21 \\ 24 & 7 & 22 & 53 \end{pmatrix} \text{ and } V = \begin{pmatrix} 5 & 1 & 2 & 4 \\ 2 & .01 & 1 & 3 \\ 2 & 1 & 2 & 4 \\ 3 & 1 & 4 & 4 \end{pmatrix}.$$

The first row in Table 1 provides the N ratio for Gx when the null hypothesis is true, and all the cell estimates are unbiased. The average of these ratios is always close to zero.

Table 1

Mean Values of Wald Test Statistics
Resulting from Biased Cell Estimates - 1000 Replications

bias	n1	n2	n3	n4	n5	n6	n7	n8
H _o true		0.001.6	0.0000	0.0004	0.04.70	0.0470	0.04.04	0.04.4-
	-0.0308	-0.0316	0.0239	0.0096	-0.0159	-0.0679	-0.0101	-0.0148
H _o false								
x1	5.7427	-0.0316	0.0239	0.0096	5.7576	-0.0679	-0.0101	-0.0148
x2	-0.0308	3.2317	0.0239	0.0096	2.2935	-0.0679	-0.0101	-0.0148
х3	-0.0308	-0.0316	2.6906	0.0096	2.2935	-0.0679	-0.0101	-0.0148
x4	-0.0308	-0.0316	0.0239	3.4737	-3.4800	-0.0679	-0.0101	-0.0148
x5	1.1239	-0.0316	0.0239	0.0096	-0.0159	2.2377	-0.0101	-0.0148
х6	-0.0308	-0.0152	0.0239	0.0096	-0.0159	-0.0448	-0.0101	-0.0148
x7	-0.0308	-0.0316	1.3573	0.0096	-0.0159	2.2377	-0.0101	-0.0148
x8	-0.0308	-0.0316	0.0239	1.1643	-0.0159	-2.3734	-0.0101	-0.0148
x9	2.2786	-0.0316	0.0239	0.0096	-0.0159	-0.0679	2.6566	-0.0148
x10	-0.0308	1.6001	0.0239	0.0096	-0.0159	-0.0679	1.3233	-0.0148
x11	-0.0308	-0.0316	2.6906	0.0096	-0.0159	-0.0679	2.6566	-0.0148
x12	-0.0308	-0.0316	0.0239	4.6284	-0.0159	-0.0679	-5.3434	-0.0148
x13	-4.6496	-0.0316	0.0239	0.0096	-0.0159	-0.0679	-0.0101	4.1164
x14	-0.0308	-4.9265	0.0239	0.0096	-0.0159	-0.0679	-0.0101	3.0836
x15	-0.0308	-0.0316	-5.3094	0.0096	-0.0159	-0.0679	-0.0101	4.1164
x16	-0.0308	-0.0316	0.0239	-4.6092	-0.0159	-0.0679	-0.0101	-4.1460

Next, a bias was introduced, by adding $4xVar(x_{ij})$ to each cell estimate, one at a time. For the first cell, this should result in a rejection or high N-value for the first and fifth restrictions; a bias in the estimate of the second cell (vectorised columnwise) should result in a rejection of restrictions 2 and 5, and so on. The mean values of the test statistics on the restrictions with the offending parameter estimates tends to be much larger than 2, and the pattern supports the previous conjecture. In some cases the tests do not appear to isolate offending initial estimates, particularly in relation to x_{22} ; however, an examination of the rejection rates associated with the test statistics will be more meaningful.

The mean results for the LM tests are given in Table 2 and are similar to the Wald test results in Table 1. The well known equality W>LR>LM is observed once again [see Berndt and Savin(1977)]. Usually the performance of a test can be improved by increasing the sample size or improved variance estimation; but it is not obvious how either can be exploited here.

Table 2

Mean Values of LM Test Statistics
Resulting from Biased Cell Estimates - 1000 Replications

	11	12	13	14	15	16	17
x 1	2.9162	-1.0963	-0.8588	1.0062	3.7092	-0.1389	0.1651
x2	-0.9296	2.4322	-0.5830	0.6515	1.8593	0.4632	0.3008
х3	-0.8697	-0.5772	1.8593	0.5323	1.9778	-0.0503	0.1048
x4	1.0937	0.7266	0.4524	2.2083	-2.6688	0.1722	0.4991
x5	0.6673	0.0407	-0.3886	0.2810	-0.1327	2.1389	0.0925
х6	0.0076	0.0306	-0.0223	-0.0050	-0.0004	0.0742	0.0375
x7	-0.3543	-0.0335	0.7181	0.3487	0.1147 °	2.1262	0.1011
x8	0.2975	0.0057	0.3421	0.6429	-0.0165	-2.0346	0.1594
x9	1.5363	-0.2561	-0.4942	0.7700	-0.2836	0.0045	2.2507
x10	-0.2776	1.3105	-0.3703	0.5077	0.0458	0.2718	1.2499
x11	-0.5068	-0.4045	1.7191	0.9054	0.2111	-0.0208	2.2679
x12	0.7287	0.6179	0.8894	2.2014	-0.0247	0.1536	-3.6728
x13	-5.8402	-1.3438	-1.7067	2.0755	3.2949	1.9027	2.4367
x14	-1.2227	-5.2956	-0.9408	1.1725	1.9067	0.7074	1.5165
x15	-1.7540	-1.0472	-6.1333	1.8047	2.3055	1.9533	2.4022
x16	2.0968	1.3182	1.7186	-5.8627	-2.7080	-1.8107	-3.0860

Table 3 presents the N-ratios for the differences between the constrained and the initial estimates divided by their standard errors when H_0 is false; that is, as a result of the biases introduced on the 16 coefficients. The approach works, the diagonal N-values should the large and significant, the off-diagonal elements insignificant. This is the observed pattern.

Further results on the same 4x4 SAM are given below. Firstly, based on 1000 replications, it is verified in Table 4, that the test statistics are (individually) standard normal under the null hypothesis. The Kolmogorov-Smirnov test is satisfied at the 5% level in all cases except one, that being one of the difference tests, and even there the rejection is only marginal. Empirical critical values are calculated for these distributions based on 2.5% of the distribution under H_0 being under either tail. These critical values are then used to establish the rejection rates under H_1 in Table 5. The perturbations are those mentioned previously and as can be seen, when a bias is introduced into the (1,1) element, the rejection rates on the corresponding row and column of the Wald and LM tests increase dramatically. The results for the difference tests (expressed in terms of the coefficients rather than the restrictions) also increase substantially for the offending coefficient and the accompanying row and column. These elements are highlighted in Table 5.

Table 3

Standardised Differences resulting from Perturbations Coefficients

Pe	rt.															
	1	2	3	4	5	6	7	8	9	10	11 1	2	13	14	15	16
1	-7.09	-2.02	-2.64	2.77	-1.52	0.83	0.67	-0.75	-2.52	0.88	0.63	-0.85	2.91	-1.11	-0.82	0.99
2	-1.08	-3.79	-1.2	1.26	0.11	-1.84	-0.1	0.07	0.59	-2.22	0.26	-0.37	-0.94	2.41	-0.54	0.63
3	-1.28	-1.08	-3.47	1.49	0.6	0.44	-1.06	-0.39	0.73	0.49	-1.63	-0.47	-0.88	-0.59	1.9	0.51
4	1.83	1.54	1.99	-4.69	-0.77	-0.55	-0.43	-1.14	-1.32	-0.93	-0.81	-1.62	1.08	0.71	0.49	2.19
5	-0.53	0.09	0.43	-0.38	-2.52	-1.85	-1.91	1.99	-0.56	-0.02	0.3	-0.25	0.66	0.02	-0.35	0.26
6	0.0	-0.01	-0.01	0.02	-0.04	-0.04	-0.05	0.05	0.06	0.03	0.04	-0.04	0	0.01	0.02	-0.03
7	0.25	-0.05	-0.77	-0.19	-1.91	-1.79	-2.56	1.94	0.3	0.04	-0.65	-0.3	-0.36	-0.05	0.76	0.33
8	-0.28	0.03	-0.32	-0.6	1.94	1.82	1.86	-2,47	-0.31	-0.05	-0.39	-0.51	0.29	-0.01	0.38	0.62
9	-1.26	0.49	0.66	-0.98	-0.86	0.19	0.31	-0.47	-3.41	-1.53	-1.66	1.67	1.53	-0.27	-0.45	0.75
10	.25	-1.24	0.25	-0.41	-0.07	-0.99	-0.04	-0.05	-0.9	-2.04	-0.82	0.82	-0.29	1.29	-0.33	0.49
11	.30	0.22	-1.75	-0.61	0.36	0.31	-1	-0.58	-1.71	-1.42	-3.56	1.57	-0.52	-0.42	1.76	0.88
12	-0.71	-0.54	-0.8	-2.05	-0.54	-0.47	-0.68	-1.16	3.1	2.57	2.79	-6.22	0.72	0.6	0.93	2.18
13	2.4	-1.45	-1.51	1.37	1.52	-0.8	-0.87	0.7	2.63	-0.76	-0.81	0.64	-5.85	-1.36	-1.67	2.05
14	-0.83	3.39	-0.93	0.83	0.04	2.64	-0.13	0.01	-0.36	3.34	-0.59	0.48	-1.23	-5.31	-0.9	1.15
15	-0.73	-0.91	3.3	0.63	-0.92	-1.03	1.75	0.91	-0.79	-0.99	2.99	0.86	-1.76	-1.06	-6.09	1.78
16	0.84	1.02	0.9	2.78	0.66	0.83	0.8	1.56	1.37	1.5	1.52	2.06	2.09	1.3	1.76	-5.88

Table 4

Simulation Results under Ho

Wald Test

means	and	standard	deviations

1	2	3	4	5	6	7	8
-0.0308	-0.0316	0.0239	0.0096	-0.0159	-0.0679	-0.0101	-0.0148
1.0089	1.0268	1.0029	0.9882	1.0001	0.9801	1.0406	1.0293
				typ	e I error		
44	54	41	52	47	49	58	51
]	Kolmogor	ov-Smirno	v Test	
0.0281	0.0413	0.0187	0.0171	0.0341	0.0361	0.0238	0.0263
			cr	itical value	e at 95% =	: 0.0430	

LM Test

means and standard deviations

 11	2	3	4	5	6	7	
-0.0022	-0.0206	0.0431	-0.0151	-0.0175	-0.0723	-0.0194	
1.0486	1.0092	1.0138	1.0011	1.0158	1.0056	1.0380	
				type I e	rror		
68	54	43	49	55	46	62	
			Kolmog	orov-Smir	nov Test		
0.0212	0.0404	0.0266	0.0165	0.0203	0.0377	0.0288	

Difference Test

means and standard deviations

1	2	3	4	5	6	7	8	
0.0216	0.0337	-0.0223	-0.0033	0.0749	0.0757	0.0468	-0.0648	
1.0061	1.0344	0.9960	0.9866	0.9681	0.9850	0.9906	0.9914	
9	10	11	12	1	3 1	4	15	16
0.0210	0.0334	-0.0182	- 0.0071	-0.0022	-0.0206	0.0431	-0.0151	
1.0116	1.0501	1.0289	1.0255	1.0486	1.0092	1.0138	1.0011	
				type I e	rror			
41	62	44	42	46	46	52	47	
53	67	57	54	68	54	43	- 49	
			Kolmo	ogorov-St	nirnov Tes	st		
0.0429	0.0412	0.0273	0.0210	0.0374	0.0483	0.0289	0.0348	
0.0259	0.0343	0.0202	0.0261	0.0212	0.0404	0.0266	0.0165	

Table 5
Rejection Rates on Test Statistics resulting from Biased Initial Estimates (H1)
Wald and LM Tests

	1	2	3	4	5 5	6	7	8	
11 Wald	1000	49	49	49	999	49	49	49	
LM	790	188	161	211	951	62	46	. 42	
21 Wald	49					49	49	40	
		870	49	49	626			49	
LM	134	649	112	130	429	92	49		
31Waid	49	49	796	49	626	49	49	49	
LM	118	88	502	104	479	54	47		
1Wald	49	49	49	940	946	49	49	49	
LM	164	104	97	641	744	59	55	72	
LIVI	104	104	91	071	/	39	33		
2 Wald	208	49	49	49	49	660	49	49	
LM	95	50	80	65	51	619	47		
2 Wald	49	49	49	49	49	50	49	49	
			51					47	
LM	50	50		49	50	52	50	40	
3 Wald	49	49	298	49	49	660	49	49	
LM	56	52	146	73	46	616	47		
l Wald	49	49	49	215	49	642	49	49	
LM	56	51	83	129	50	560	46		
			40	40	40				
Wald	629	49	49	49	49	49	745	49	
LM	319	55	93	153	52	51	494		
Wald	49	286	49	49	49	49	262	49	
LM	57	229	79	98	49	67	165		
Wald	49	49	796	49	49	49	745	49	
								49	
LM	67	66	440	188	48	54	506		
Wald	49	49	49	997	49	49	1000	49	
LM	108	86	176	637	52	61	948		
Wald	00=	40	40	40	40	40	40	007	
Wald	997	49	49	49	49	49	49	987	
LM	999	268	439	597	885	528	578		
Wald	49	997	49	49	49	49	49	864	
LM	205	998	179	250	452	137	225		
Wald	49	49	999	49	49	49	49	987	
				496		545	564	207	
LM	368	175	1000		619			000	
Wald	49	49	49	996	49	49	49	982	
LM	513	230	440	1000	756	466	876		
			,	The Differ	rence Test				
	1	2	33	4	.01100 1 030	. 1	2	3	4
	1000	442	754	824	12	93	51	77	82
	360	127	110	104	12	744	463	448	526
	698 700	148	92 161	134		88 95	53 50	62 80	56 65
	790	188	101	211		75	30	6U	63
	219	949	242	281	22	50	49	49	49
	47	454	50	52	22	51	50	51	51
	88	536	61	62		51	50	50	50
	134	649	112	130		50	50	51	49
	202	150	000	051	20	c -	51	120	~0
	283	150	933	371	32	65	51	132	58
	89	73	150	61		523	439	705	509
	111	84	361	69		59	50	100	53
	118	88	502	104		56	52	146	73
	502	369	503	999	42	67	50	64	105
	126	90	51	18		473	429	459	678
	244	131	126	361		64	52	72	74
	164	104	97	641		56	51	83	129
	282	73	109	178	41	694	262	342	322
	146	46	63	62		322	133	112	105
	920	274	371	408		749	104	126	112
	319	55	93	153		999	268	439	597
	66	187	58	84	42	160	914	172	144
	51	172	50	48		46	757	50	49
	149	453	127	137		70	896	91	90
	57	229	79	98		205	998	179	250
								4.1	
}	66	64	432	111	43	126	122	913	106
	62	53	138	79		169	177	416	146
	383	241	933	365		128	140	821	148
	200		440	188		368	175	1000	496
	67	66				-			
	67	66	•••						
	124	71	135	565	44	157	187	150	824
					44	157 100	187 127	150 139	824 355
	124	71	135	565	44				

The results for Tables 6 and 7 are based on a 4x4 SAM with all initial cells and their variances of the same magnitude (equal to 1 and 0.2 respectively). The final row and column variances were then the sum of their rows and columns and the weighted quadratic form respectively. The story is much the same as observed previously

			Ta	able 6			
		Simulati	on Resul	ts under l	Но: Туре	1 Error	
				Wald Test			
1	2	3	4	5	6	7	8
43	51	45	43	43	47	56	67
			Kolmogo	rov-Smir	nov Test		
1	2	3	4	5	6	7	8
0.0286	0.0286	0.0160	0.0185	0.0111	0.0240	0.0314	0.0397
			K-S Criti	ical Value	0.0430		
				LM Test			
	1	2	3	4	5	6	7
	46	56	53	48	55	56	52
	•		Kolmogo	rov-Smiri	nov Test		
	0.0299	0.0290	0.0313	0.0303	0.0217	0.0201	0.0471
			Di	ff Test			
48	51	46	49	44	53	45	48
40	53	45	55	46	56	53	48
			Kolmogo	rov-Smiri	nov Test		
0.0263	0.0185	0.0196	0.0164	0.0283	0.0288	0.0140	0.0208
0.0238	0.0269	0.0317	0.0348	0.0299	0.0290	0.0313	0.0303

Table 7

		_			M tests: 1	Rejection			
11 377 11		1 220	2 49	<u>3</u> 49	4	1	. 2	3	4
11 Wald	ļ.	320			49	357	49	49	49
LM		198	54	55	62	242	49	49	
12 Wald	l	49	310	49	49	357	49	49	49
LM		70	187	55	62	242	49	49	
31 Wald	1	49	49	339	49	357	49	49	49
LM		70	54	187	62	242	49	49	
11 Wald	Į.	49	49	49	722	992	49	49	49
LM		91	97	88	180	959	49	49	
		220	40	40	40	10	***	40	
21 Wald LM		320 198	49 54	49 55	49 62	49 49	297 213	49 49	49
22 Wald		49	310	49	49	49	297		40
								49	49
LM		70	187	55	62	49	213	49	
3 Wald		49	49	339	49	49	297	49	49
LM		70	54	187	62	49	213	49	
4 Wald		49	49	49	722	49	993	49	49
LM		91	97	88	180	50	963	49	
3 Wald		320	49	49	49	49	49	295	. 49
		198	54						+9
LM				55	62	49	49	226	
3 Wald		49 70	310	49	49	49	49	295	49
LM		70	187	55	62	49	49	226	
3 Wald		49	49	339	49	49	49	295	49
LM		70	54	187	62	49	49	226	
3 Wald		49	49	49	722	49	49	992	49
LM		91	97	88	180	50	49	964	
4 337-1.2		noa.	40	40	40	40	40	40	649
Wald!		992	49	49	49	49	49	49	648
LM		997	104	110	180	242	213	226	
4 Wald		49	989	49	49	49	49	49	648
LM		120	996	110	180	242	213	226	
Wald		49	49	992	49	49	49	49	648
LM		120	104	994	180	242	213	226	
4 Wald		49	49	49	1000	49	49	49	1000
LM		532	498	481	1000	959	963	964	
1	1 482	2 129	3 132	220	Difference 21	1 117	<u>2</u> 55	<u>3</u> 67	<u>4</u> 45
_	153	57	51	49		537	131	141	205
	152	49	45	53		152	49	45	53
	198	54	55	62		198	54	55	62
l	117	499	132	220	22	53	129	67	45
	57	131	51	49		153	450	141	205
	64	158	45	53		64	158	45	53
	70	187	55	62		70	187	55	62
		440	405		,			400	
	117	129	487	220	23	53	55	132	45
	57	57	141	49		153	131	504	205
	64	49	164	53		64	49	164	53
	70	54	187	62		70	54 .	187	62
	815	765	811	997	24	67	69	79	82
	85	703	82	119	201	829	805	788	997
	85 90	95	62 100	97		90	95	100	997 97
	90 91	95 97	88	97 180		90 91	95 97	88	97 180
	71	71	συ	100		71	71	υυ	100
	117	55	67	45	14	815	69	79	106
	153	57	51	49		829	77	82	104
	507	158	164	184		818	95	100	101
	198	54	55	62		997	104	110	180
		4.6-							
	53	129	67 51	45	24	67	765	79	106
	57	131	51	49		85	805	82	104
	152	528	164	184		90	746	100	101
	70	187	55	62		120	996	110	180
	53	55	132	45	34	67	69	811	106
ı		57	141	49	٥.	85	77	788	104
i	57		511	184		90	95	778	101
•	57 152	158		107					101
	152	158 54				120	104	994	180
	152 70	54	187	62		120	104	994	180
	152 70 67	54 69	187 79	62 82	44	390	382	398	564
	152 70 67 85	54 69 77	187 79 82	62 82 119	44	390 427	382 391	398 382	564 547
3	152 70 67	54 69	187 79	62 82	44	390	382	398	564

The single test results in Tables 6 and 7 suggest that examination of the Lagrange or Wald test associated with the individual restrictions can enable the identification of biased initial estimates. However, a more thorough approach would be to examine rejection rates for the row-column interaction when both tests **simultaneously** reject. In Table 8 this is done and the joint rejections turn out to be quite low. This highlights the point that the actual Wald or LM test should be a χ^2 test (with two degrees of freedom) on the joint restrictions. The test, of course, is just $\begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\ \lambda & \lambda \end{pmatrix} = \lambda \begin{pmatrix} \lambda & \lambda \\$

				Table	8			
		Reject	ion Rates	s under H	I ₁ (4 sect	or SAM)	}	
		•			Diff tests			
	W1	W2	W12	L1	L2	L12	Diff	
1	320	291	117	193	213	21	464	
2	325	291	121	201	213	20	447	
3	361	291	129	240	213	24	526	
4	712	991	707	166	958	150	994	
5	320	338	132	193	265	26	521	
6	325	338	143	201	265	31	490	
7	361	338	137	240	265	41	559	
8	712	995	708	166	978	153	998	
9	320	296	115	193	233	26	494	
10	325	296	130	201	233	31	472	
11	361	296	135	240	233	31	504	
12	712	992	708	166	961	152	997	
13	994	753	751	0	0	0	995	
14	990	753	751	0	0	0	997	
15	995	753	749	0	0	0	1000	
16	1000	1000	1000	0	0	0	1000	
		Rejection	on Rates u	ınder H1 (10 sector	SAM)		
			Wald, I	LM and D	iff tests			
1	135	114	18	116	128	9	287	
2	140	114	17	127	128	11	208	
3	140	114	7	126	128	6	255	
4	130	114	12	137	128	9	209	
5	121	114	19	105	128	11	204	
6	165	114	17	146	128	9	256	
7	146	114	19	130	128	12	223	
8	141	114	17	126	128	11	259	
9	151	114	22	132	128	13	237	
10	704	1000	704	200	1000	200	1000	
11	135	141	25	116	135	14	203	
12	140	141	21	127	135	16	238	

W1 and W2 refer to the individual Wald (row and column) tests. W12 refers to the joint outcome of the two individual tests. The surprising result is that the joint tests (done individually) reject much less freqently than the Difference test. The W12 and L12 columns refer to the rejection count on both restrictions simultaneously. (Note the LM test was not calculated for the last 4 restrictions.) In the second half of the table, the first 12 rows (from X11 to X22) illustrate the same point in the context of a 10 sector SAM. However, the comparisons really are irrelevant unless the appropriate χ^2 tests are done.

In Table 9 the joint test results are presented for the 4 and 10 sector models. The results for the 10 sector model are not reported under H_0 ; briefly, there are no problems with the distributions of the test statistics under H_0 for either the 4 or 10 sector models. Only the first three rows are reported for the 4-sector SAM and only the first row is reported in the 10 sector case. The results are fully representative. The Difference test (size adjusted) has better power than the Wald test in every situation, thought the joint test now performs at a similar level to the Difference test. The fact that the Difference test (size adjusted) correctly rejects more frequently than the Wald (or LM) tests, in every case, suggests the existence of an underlying inequality favouring the Difference test. .

Table 9 Wald $\chi^2(2)$ and Difference Tests (4 sector SAM) 1000 replications

					Di	istri	bution	unde	r H	0						
				Reje	ection	Ra	ite at 5	% lev	el: 1	type	1 error	•				
			11	12	13	14	21	22	23	24	31	32	33	34		
		w	50	54	49	54	47	52	49	55.	45	53	49	55		
		D	41	52	46	52	59	48	54	49	47	55	56	48		
				Kol	moge	orov	/ Smir	nov te	st							
	_11	12	13	14	2	21	22	23		24	31	3	32	33	34	
W	.0272	.0272	.0302	.0185	.02	17	.0169	.0291	J. J	228	.0396	.01	91	.0285	.0327	cv(1 sided) = .0386
·D	.0460	.0329	.0252	.0355	.03	72	.0263	.0232	2 .0	218	.0332	.02	229	.0181	.0296	cv(2 sided) = .0430

	Distribution under H1: Rejection Rate at 5% level											
	11	12	13	14	21	22	23	24	31	32	33	34
W	367	349	376	988	373	348	375	985	411	378	390	990
Ð	515	479	562	996	453	485	471	997	556	490	501	998

Wald χ^2	(2) and D	ifferenc	e Tests (1	0 sector S	SAM)
Distrib	ution unde	er H1: R	ejection Ra	te at 5% le	evel
13	14	15	16	17	18

	11	12	13	14	15	16	17	18	19	110
W1	163	173	178	166	186	180	185	171	179	1000
D1	249	242	239	197	259	249	206	203	226	1000
W2	152	158	163	159	169	141	177	195	176	1000
D2	182	196	218	1 79	201	213	221	207	245	1000
W3	195	178	184	163	174	173	176	176	175	1000
D3	244	188	270	240	206	226	217	214	213	1000
W4	167	179	171	177	172	160	175	186	172	1000
D4	227	206	190	262	242	224	224	248	222	1000
W5	173	176	158	170	162	153	158	162	167	1000
D5	206	238	197	216	222	193	215	230	185	1000
W6	157	154	158	179	177	162	168	168	169	1000
D6	214	190	228	211	229	225	218	221	176	1000
W7	159	152	153	154	167	168	160	176	153	1000
D7	206	185	227	192	251	238	197	223	205	1000
W8	182	181	182	160	182	163	202	182	172	1000
D8	261	209	226	240	257	248	254	190	264	1000
W9	148	149	157	146	159	149	152	151	163	1000
D9	1 7 9	226	226	273	170	252	187	202	248	1000

A further Monte Carlo study allowed the initial variances (V) to vary randomly as well as X. The same results were observed as above. Recognising that V has to be estimated and allowing for randomness in that operation (a variation of about 40% based on a uniform distribution) made little difference to the results under either the null or the alternative hypothesis.

4. Conclusions

The results are a vindication of common sense. The Difference test works well and does not appear to be adversely affected by the size of the problem. The statistical theory for the test procedures is straightforward and the tests are properly behaved under the null and alternative hypotheses. Monte Carlo simulations bear out the intuition and the creator of a SAM can choose between Wald, LM and Difference tests in trying to pinpoint initial estimate bias. The simplest procedure, the Difference test, works well, and its use for diagnostic investigation in data construction should be encouraged.

The importance of this is a validation of procedures being used in the field and ammunition to those who have been arguing against those government statistical offices which are unwilling to conceed that the figures they release are only estimates and are subject to random errors, as all estimates are. It has implications for national accounting, trade flow data, input-output analysis as well as SAM construction.

5. References

- 1. Arkhipoff, O., (1969), "Les Circuits Financiers au Cameroun", Compatilite Nationale 1965/66, etude speciale no 2; Direction de la Statistique, Yaounde.
- 2. Berndt, E.R. and Savin, N.E., (1977), "Conflict among Criteria for Testing Hypotheses in the Multivariate Regression Model", *Econometrica*, 45, pp.1263-1278.
- 3. Bickel, P.J. and Doksum, K.A., (1977), *Mathematical Statistics: Basic Ideas and Selected Topics*, Holden-Day, Oakland.
- 4. Byron, R.P., (1972), "Testing for Misspecification in Econometric Systems using Full Information", *International Economic Review*, 13, pp.745-756.
- 5. Byron, R.P., P.J.Crossman, J.E.Hurley and S.C.E.Smith, (1993), Balancing Hierarchical Regional Accounting Matrices, *International Conference in memory of Sir Richard Stone*, Certosa di Pontignano, Siena, ITALY, October 17-20, 1993.
- 6. Stone, R. (1975), Direct and Indirect Constraints in the Adjustment of Observations, in P.J.Bjerve (ed.), *Nasjonalregnskap, Modeller og Analyse*, Statistisk Sentralbyra, Oslo.
- 7. van der Ploeg, F. (1982), Reliability and the Adjustment of Sequences of Large Economic Accounting Matrices, *Journal of the Royal Statistical Society, Series A* 145, pp. 169-194.