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Outline

- Introduction
- smod_t2.gms with different data
 - Adjust model data load
 - Rerun previous experiment
- smod_t3.gms
 - Assign equations
 - Check
 - Rerun previous experiment
 - Compare results
- Change nesting structure in smod_t3.gms
 - Data
 - Rerun previous experiment
- Comparing Results

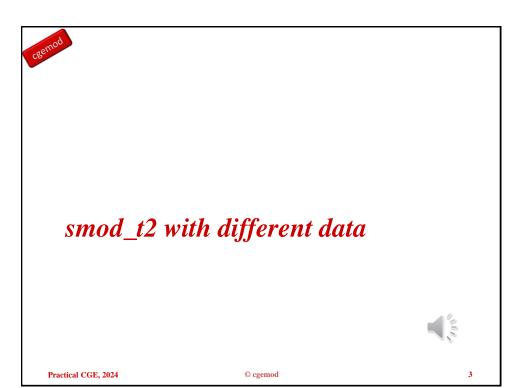


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Data

Well-designed models should be able to run with different data when

- Behavioural relationships are unchanged
- SAM structure is **unchanged**

Review smod t modO6 t3.xlsx

Same SAM structure as in smod_modO6_t2.xlsx

- 10 commodities/activities
- 5 'natural' factors
- 2 households
- 5 Tax instruments

ALL elasticity values are 'place holders'

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Change Data for smod_t2

- 1. Open smod t2.gms in Studio
- 2. In the file smod_t2.gms open the file smod_t2_load.inc (use Ctrl+click on the \$INCLUDE statement)
- Change the \$GDXXRW instruction to load the data file smod t modO6 t3.xlsx
- 4. Run the programme (F10) and check the model is running correctly

The ability to change data used by a 'standard' model is valuable

- When making changes to behaviour in a model starting with an aggregated SAM is useful
- Sometimes a model can be used for different SAMs



Do NOT change data to match a model

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Experiment

Trade policy experiment from Module O5

Experiment File

- 1. Use the file smod t2 exp2.inc used with smod t2,
- 2. Make any necessary changes to the code for saving results
 - i. Extend the set scal to include the additional scalar results and extend the results reported in *levSCAL* and *pcSCAL*.
 - ii. Declare parameters results for the new variables, i.e., lev^{**} and pc^{**} , and assign values to these parameters.
- Add the instruction gdx=[unique file name] in the command line
- 4. Now run the programme (F10) with the same experiment file used for the smod t2 exercises.
- 5. Review the results

For each experiment add the instruction gdx=*** in the command line, where *** identifies different experiments.

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Equation Coding for smod_t3.gms



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smod_t3.gms coding

- Complete data file provided
- · All sets declared and assigned
- All parameters declared and assigned
- All variables declared and initial values assigned
- All equations declared

Your tasks

- 1. Assign equations
- 2. Revise model definition
- 3. Revise factor clearing
- 4. Run trade policy experiment
- 5. Compare results

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Data

Review smod t modO6 t3b.xlsx

Same SAM structure as in smod mod06 t2.xlsx and smod t mod06 t3.xlsx

- 1. What mapping sets are declared?
- 2. What parameters are declared?
- 3. How many commodities, activities, factor and households are declared and assigned?

ALL elasticity values are 'place holders'



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Using \$STOP

GAMS Configuration: set to produce $smod^{***}.gdx$ and $smod^{***}.ref$ files with F10

Progress One 'Baby' Step at a Time

• Use \$stop to set end of code run

Step 1:

- Data Entry
- Data adjustments and scaling
- Data Diagnostics
- Additional set assignments
- Parameter assignments
- Variable initialisation

Checks for

- syntax errors
- data errors
- (some) coding errors

Gets to starting point for model coding

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Equation Assignment – smod_t3

Equations needed are already declared

Transform the algebraic expressions, in the smod_t technical document (Appendix 3), into GAMS code.

Tricky parts

- use of LHS and RHS \$ control at different levels of the production nest
- use of \$ control in parameter calibration

'new' variables already initialised

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Some Questions

- Why are the sets ff(sac) and f(ff) necessary? Note the following
 - What is the difference between the members of fag and f?
 - Factor incomes (YF) are only received for natural factors, f. Why?
 - Do the aggregate factors, members of fag, have 'real world' identities? If not, what are their roles in the system?
- What are the roles of the \$ controls on equations for QVAFOC(ff,a) and FDPRODFN(ff,a) and FDFOC(ff,ffp,a) (mapping sets run TO FROM)
 - map_va_ff(ff,a)
 - map_fagg_ff(ff,ffp,a)
- What are the roles of the RHS \$ controls in QVAPRODN(a), QVAFOC(ff,a) and FDPRODFN(ff,a) and FDFOC(ff,ffp,a)
 - map_va_ff(ff,a) and map_fagg_ff(ff,ffp,a)
 - deltava(ffp,a) and deltafd(ff,ff,a)
- Why are the following statements that fix variable values needed
 - FD.FX(f,a)\$(NOT SAM(f,a)) = 0.0;
 - WF.FX(ff)\$fag(ff) = WFO(ff) * CPI.L;

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Model Definition & Closure Conditions

Defining the Model

The new equations need to be added to the model definition. This has been done for you in the template.

The model closure file - smod_t3_cl_base.inc - updated

- Compare this closure file with the closure file smod t2 cl base.inc
- Explain why the changes are needed



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Check the Model

- 1. Check the data in the model are the intended data.
- 2. Check that the value for VAR WALRAS is zero.
- 3. Check that the basic prices (PE, PD, PM) are equal to one.
- 4. Check that all entries in ASAM1CHK are equal to zero
- 5. Check that all entries in ASAM2CHK are equal to one
- 6. Check the LHS values.
- 7. Numéraire check.

Only after these checks have been passed should you move on to using the model.

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A Simple Experiment



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Experiment

Trade policy experiment from Module O5

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Comparing Results



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Comparing Results

Comparison of smod t2.gms with smod t3.gms

Comparison of the Results

- 1. To compare the results across the 2 different assumptions about factor market clearing, use the GDX MERGE utility. Open the file compare.gms and save it as compare**.gms.
- 2. Edit the call statement to refer to the gdx files generate by each run of the model.
- 3. Run compare**.gms (make sure compare**.gms is the Main File before running the model). Review the merged file, MERGE.GDX.

Analyse the results

Emphasise explanation not simple reporting of the results.

- 1. Factor demands (FD)
- 2. Factor prices (WF and WFDIST) and factor incomes (YH)
- 3. Value added (QVA) and intermediate inputs (QINT)
- 4. Production structure (*QX* and *QXC*)
- 5. Domestic quantities (QD and QQ)
- 6. Household incomes (YH) consumption (HEXP & QCD) & welfare (EV)

7. etc.

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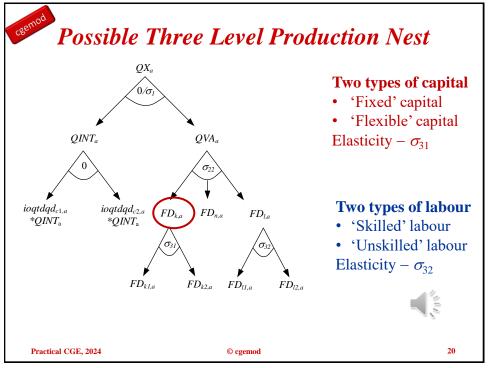
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Tasks

- Extend the sets ff and fag to include aggregate labour
 - Make changes in an Excel file
 - Extend the set *sac*
- Modify the maps that control the nesting
 - map_va_ff(ff,a)
 - map_fagg_ff(ff,ff,a)
- Checks (use \$stop?)
 - data intended are being loaded
 - parameters are correctly assigned
 - model runs correctly
- Run the experiment (different model)
- Compare the results across different models with the same data

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Comparing Results



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Comparing Results

smod_t2.gms with variants of smod_t3.gms

Comparison of the Results

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The End

Nested Functions Exercises 2



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