

Deltares

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Efficient Model Calibration using Sub-models

New development within iPESTP of iMOD

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Status Quo

What

Investigate an innovative solution to support large-scale parameter estimation for groundwater flow models

Clients

LHM (National Model NL, RWS) and Zeeland Model (Regional Model Prov. Zeeland)

Context

Spatial Calibration of the transient LHM- and Zeeland Model (SeaWAT).

Status

Currently available as prototype in iMOD v5.6

Applicability

Regional models might be able to be calibrated more efficiently

Information

iMOD Manual; PPTX of Modflow-and-More conference; paper (Groundwater) in prep.

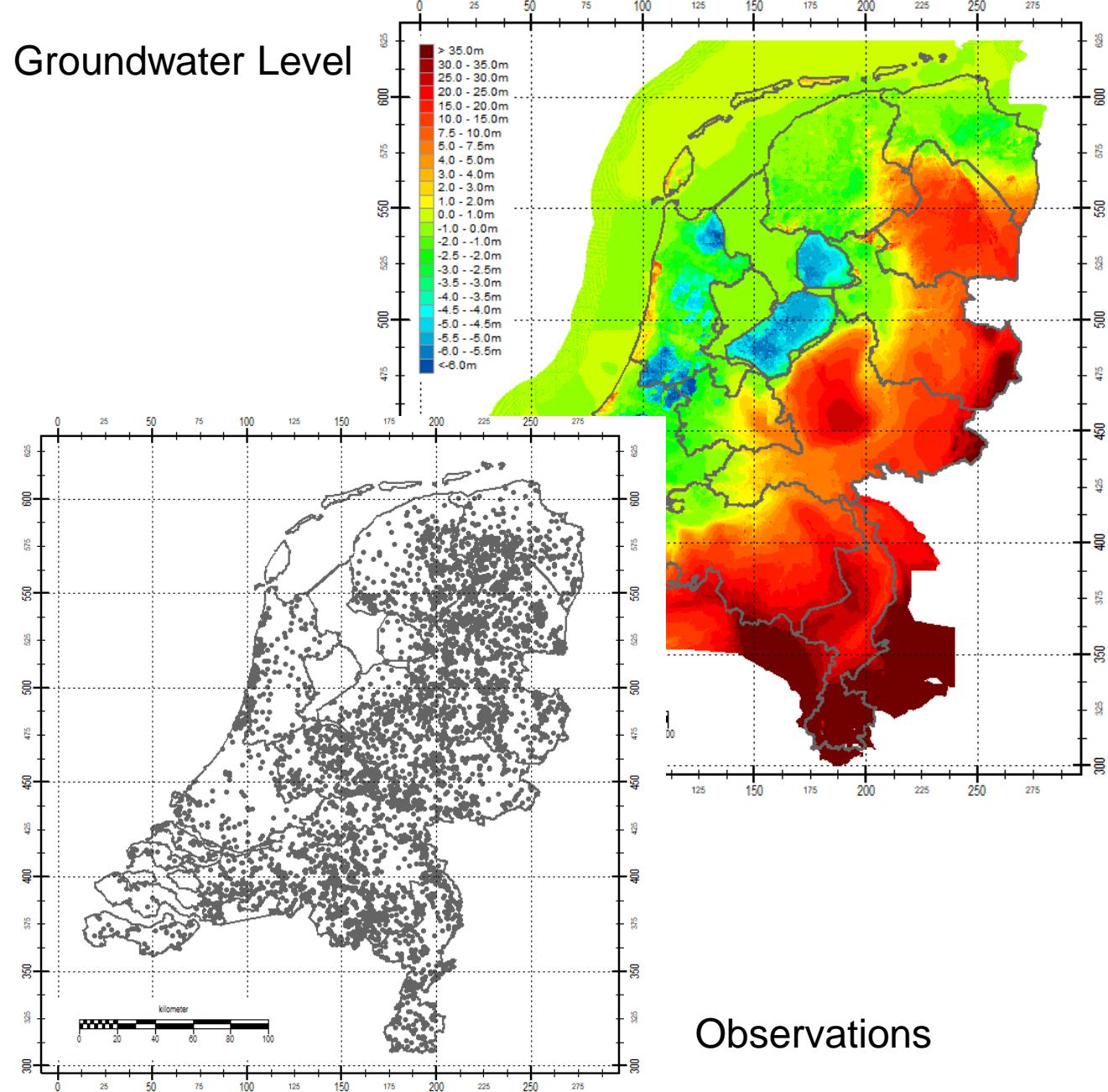
... in highlights some results ...

Challenge

Optimize

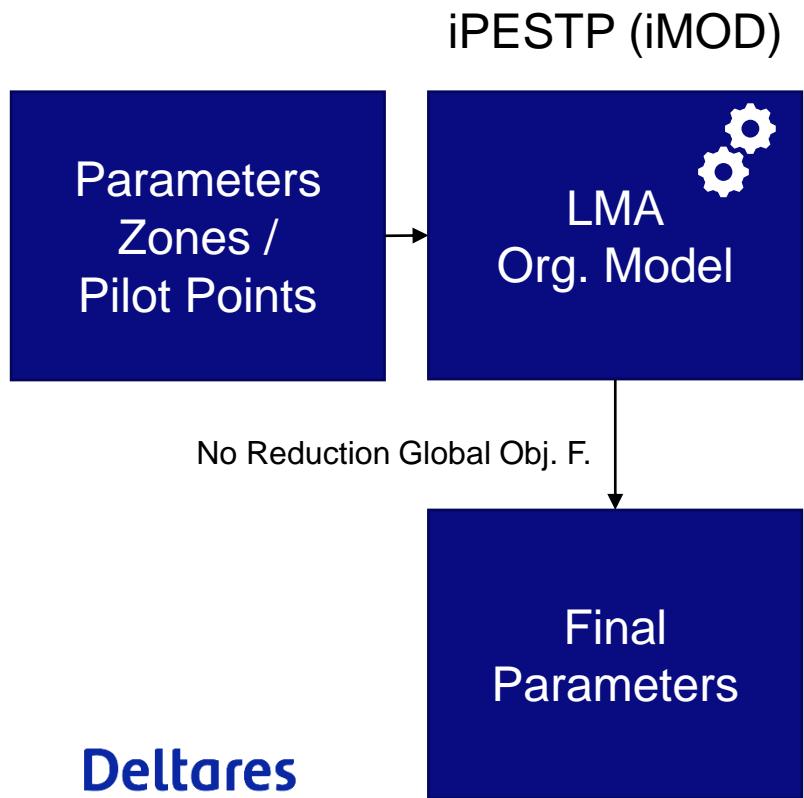
- 3D Groundwater Flow model
 - Transient Model (171 stresses)
 - 12.5 millions nodes
- Totally 2,050 parameters
 - REGIS units in aquifers and aquitards
 - GeoTOP lithology in “dekklaag”
 - Drainage- and river conductance
 - Infiltration factor rivers
 - Storage coefficients
- Observations
 - 0.5 million time-variant observations
 - 60,000 artificial observations GxG

Groundwater Level

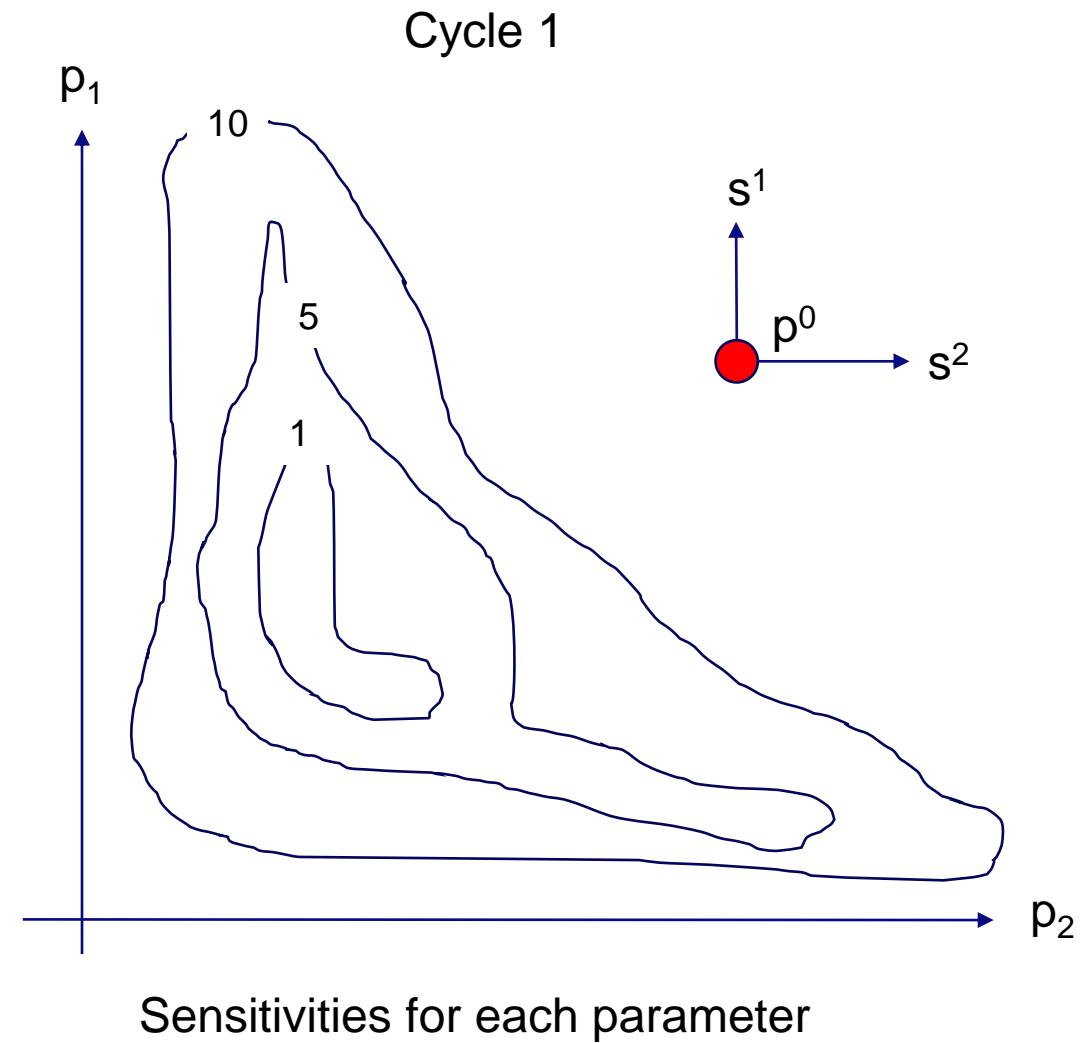
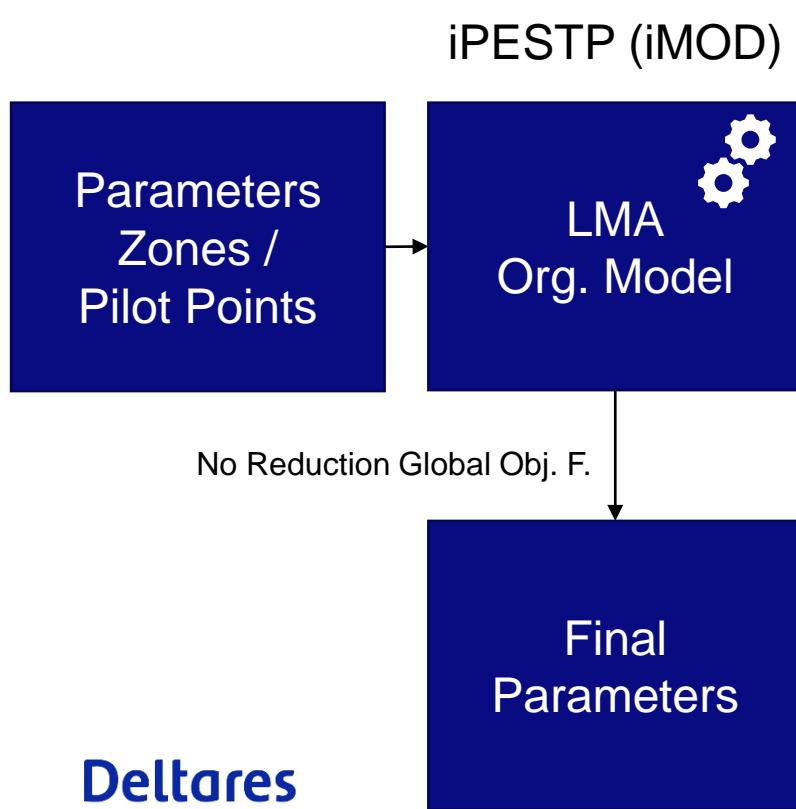


Observations

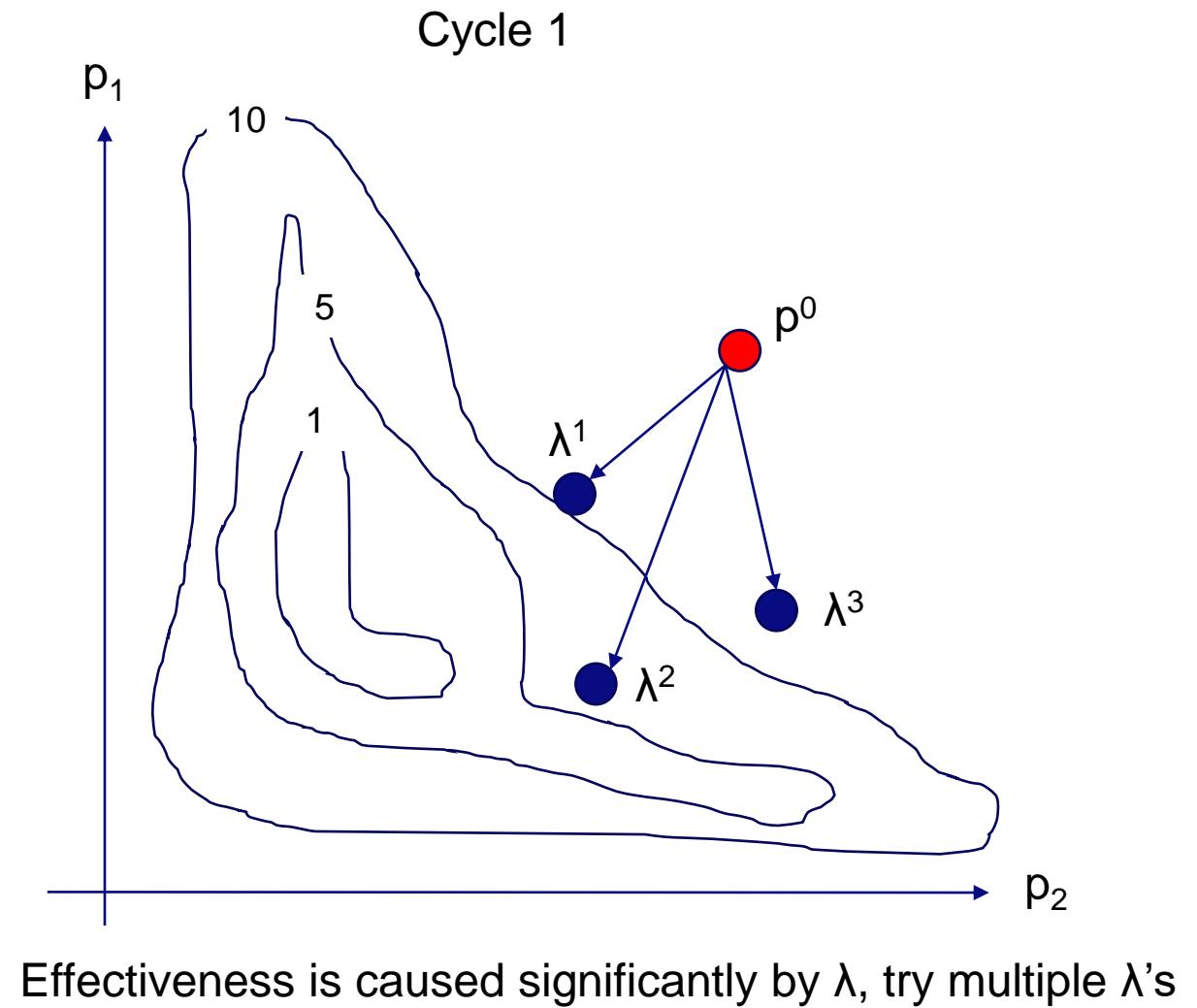
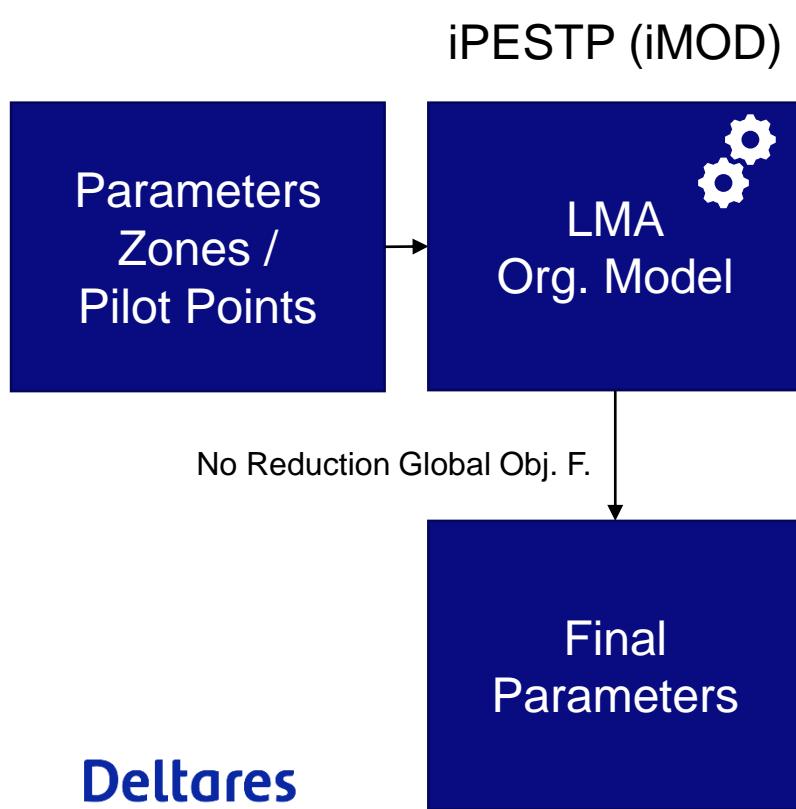
LMA Optimization Method



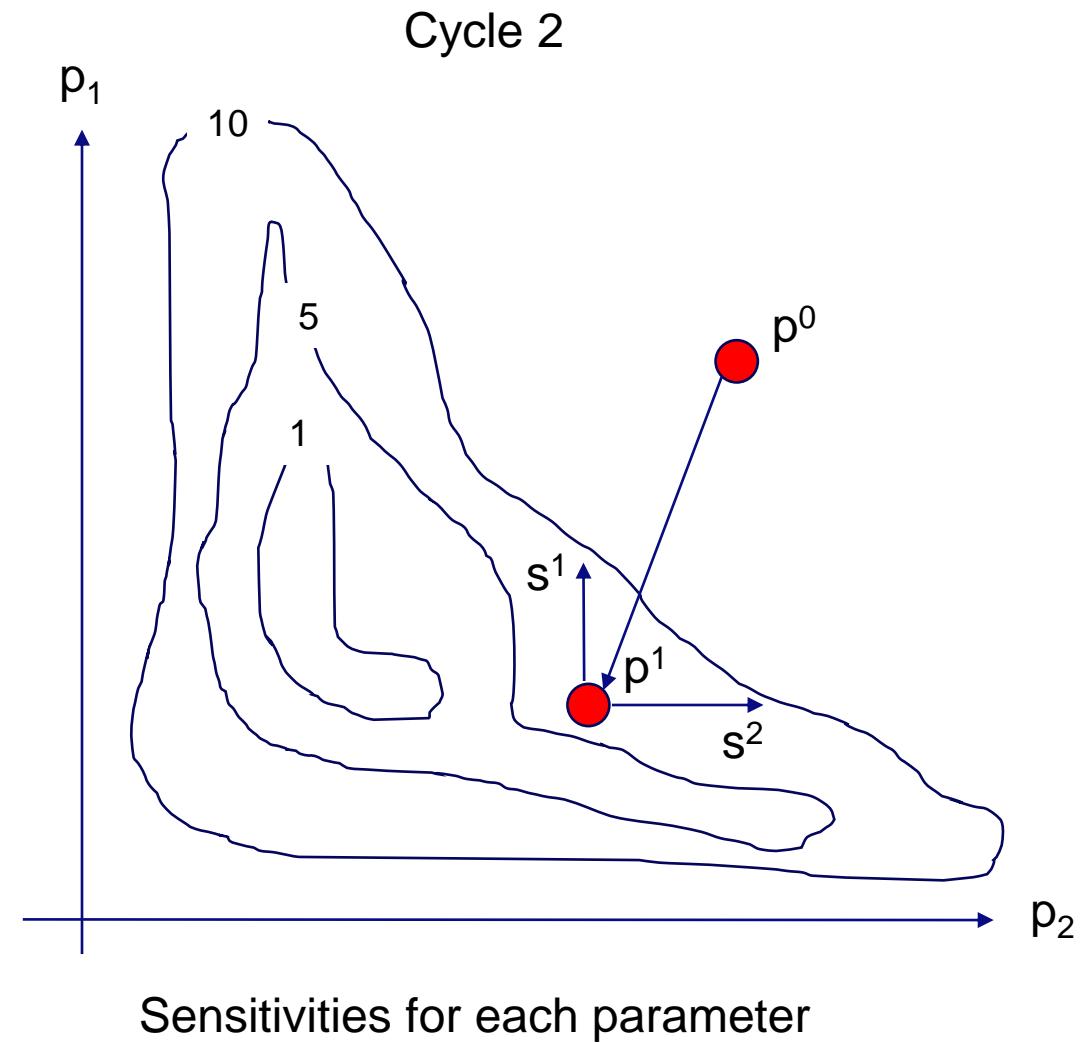
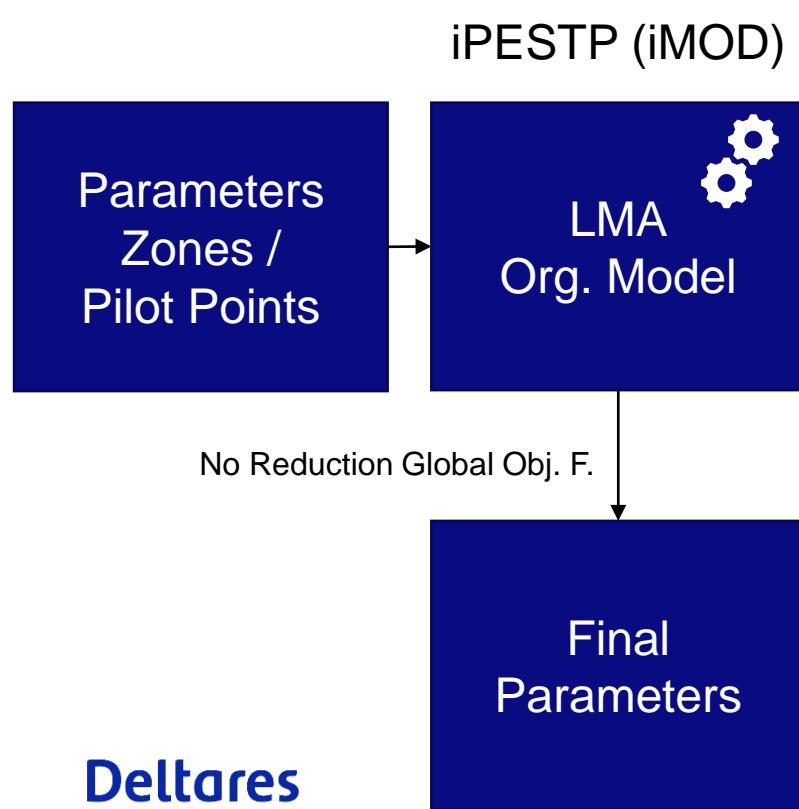
LMA Optimization Method



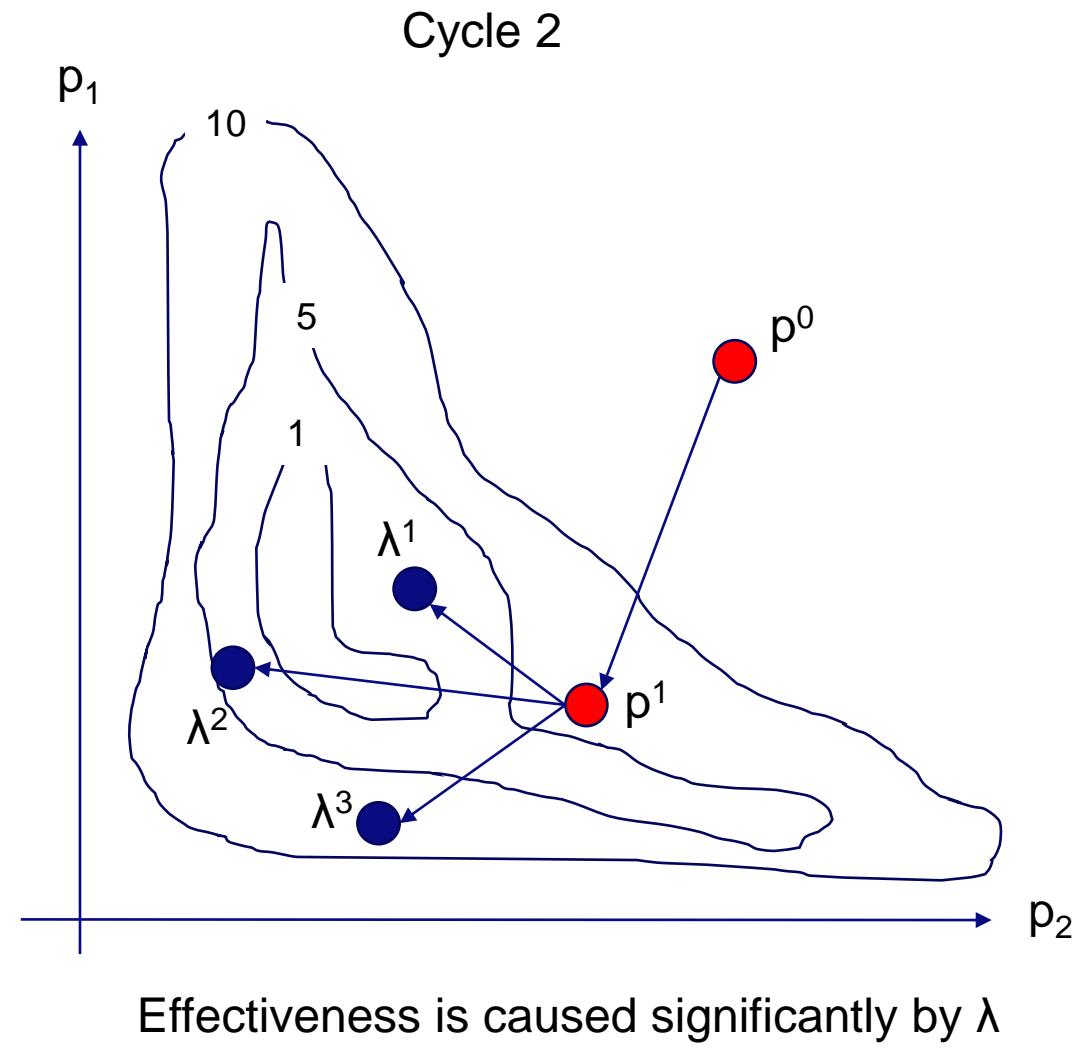
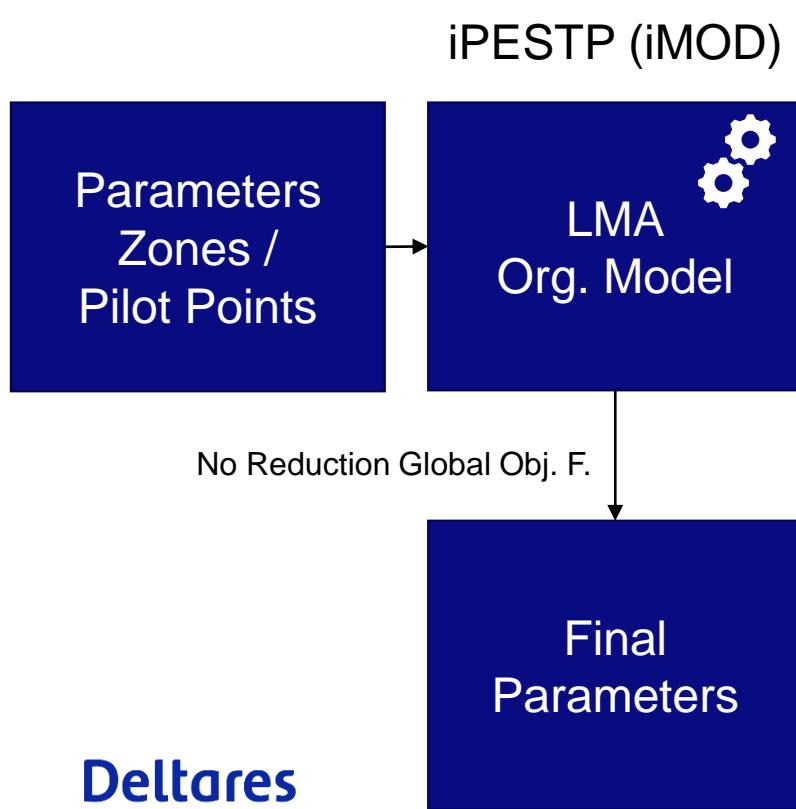
LMA Optimization Method



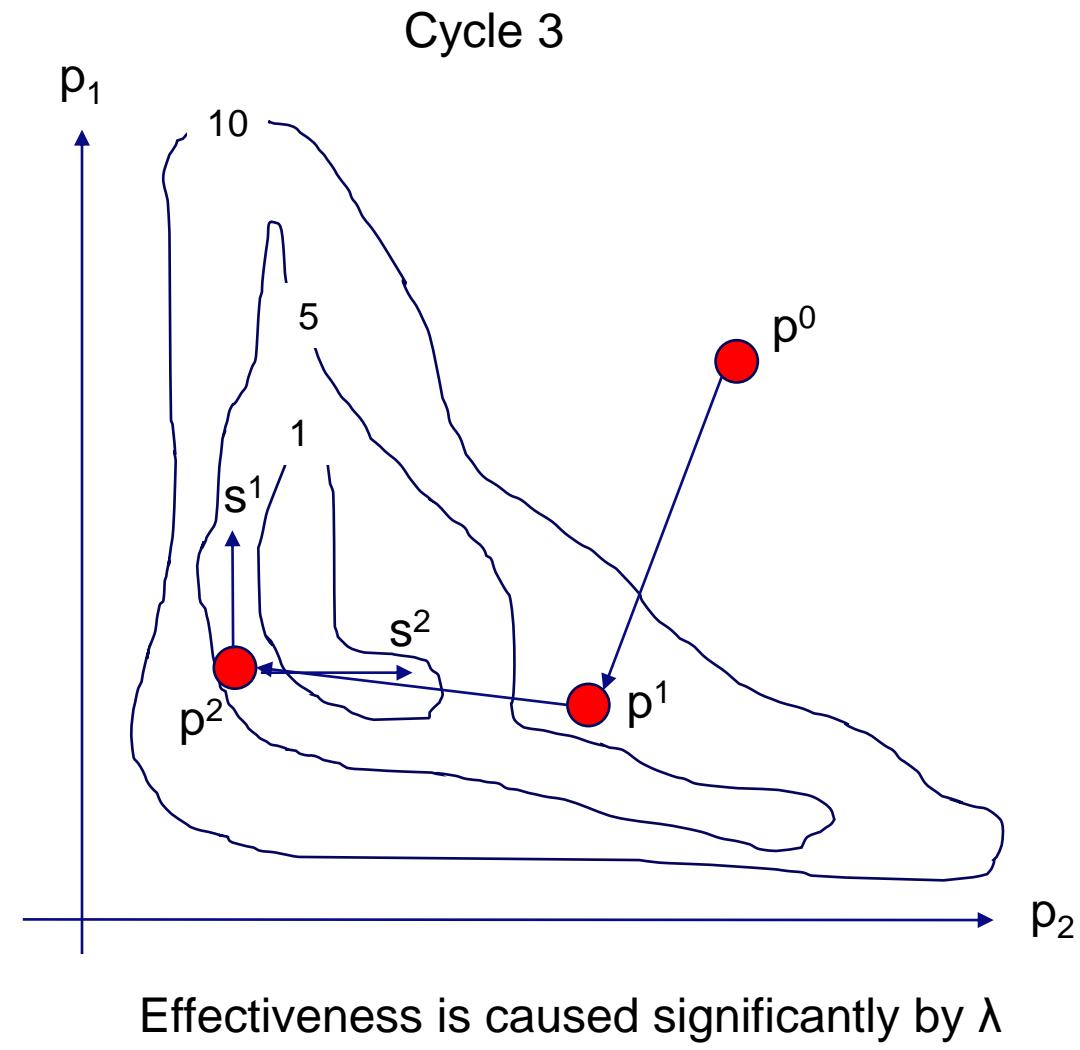
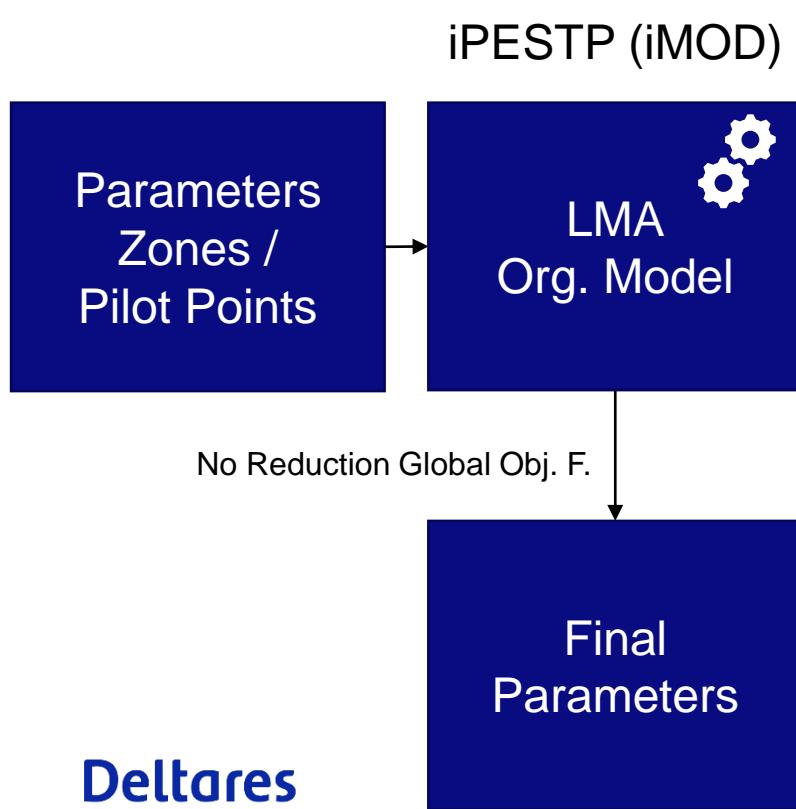
LMA Optimization Method



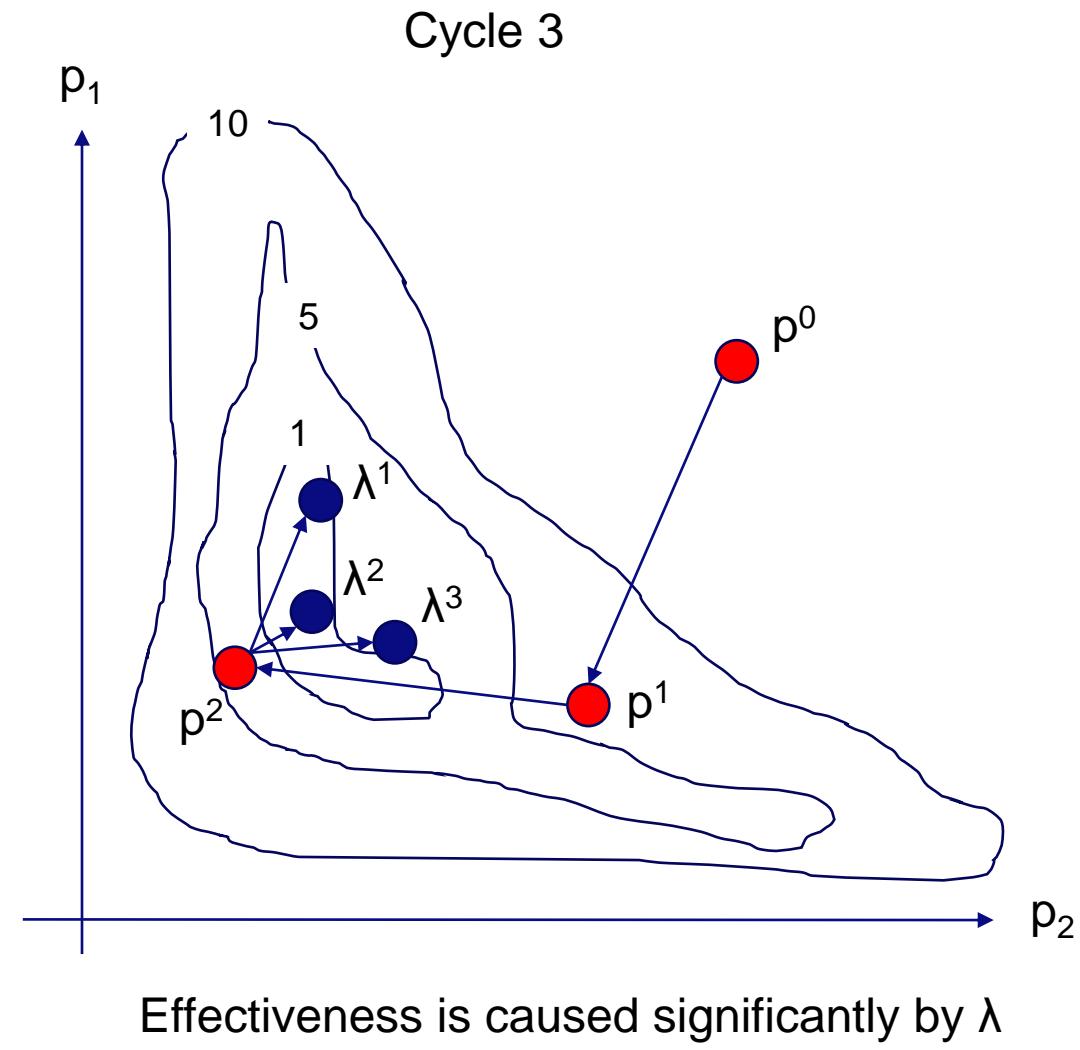
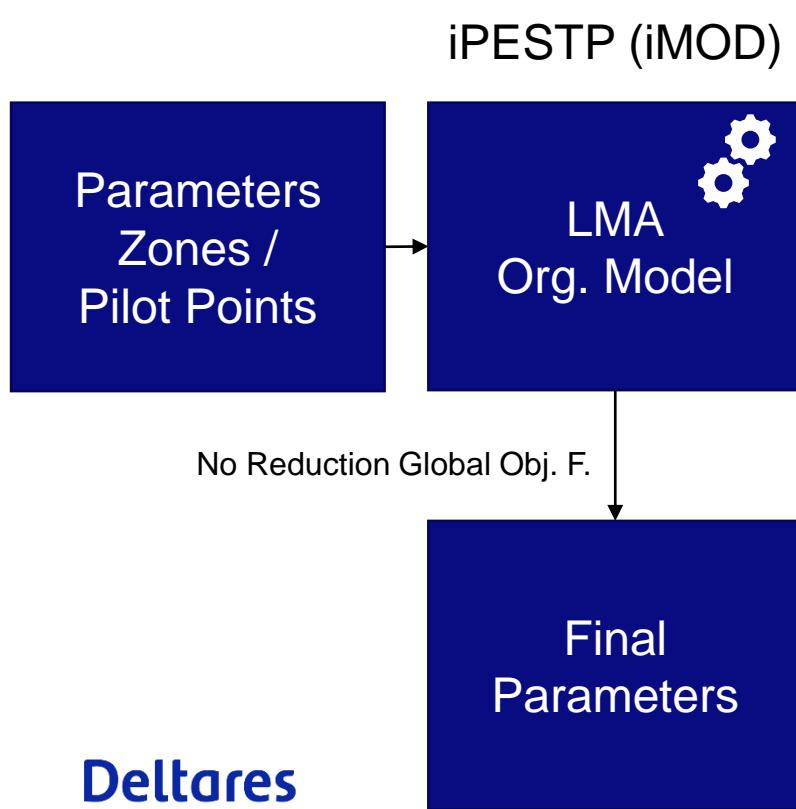
LMA Optimization Method



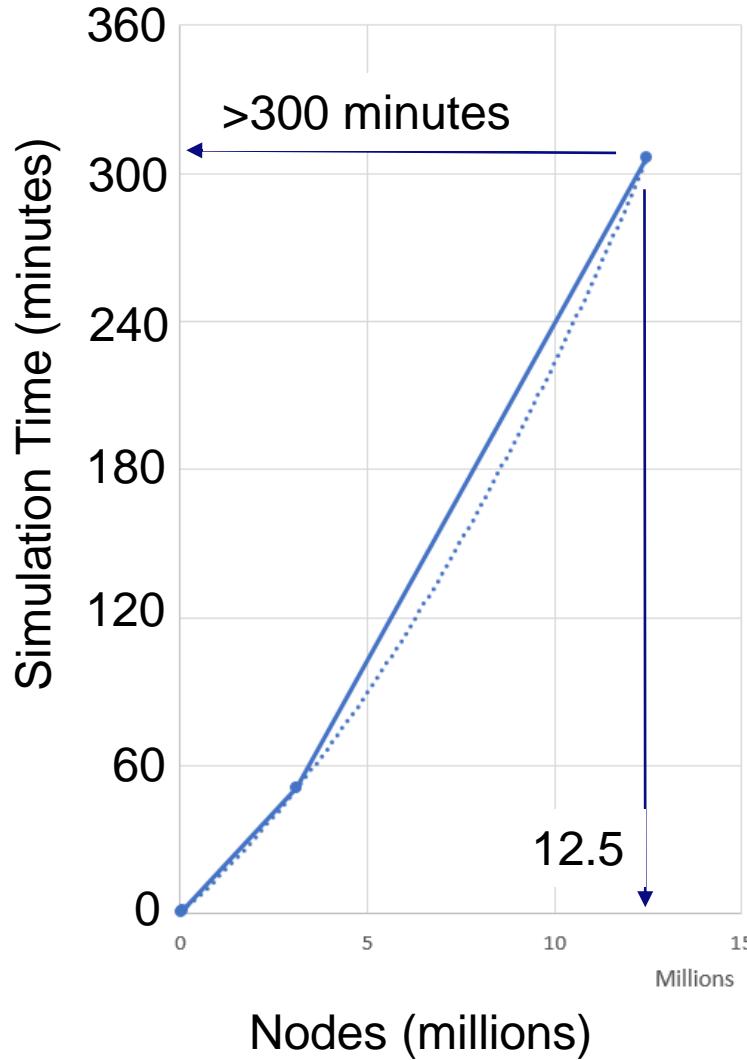
LMA Optimization Method



LMA Optimization Method



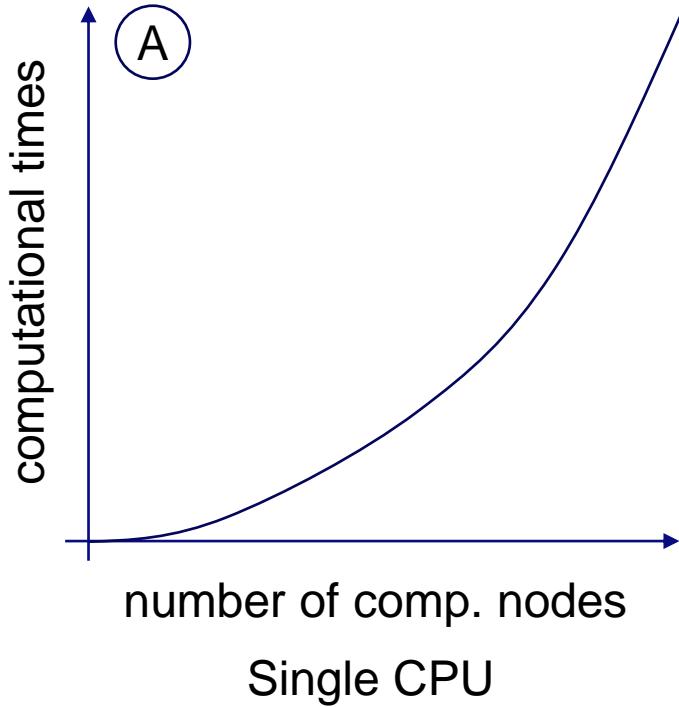
Efficiency of LMA Optimization Method



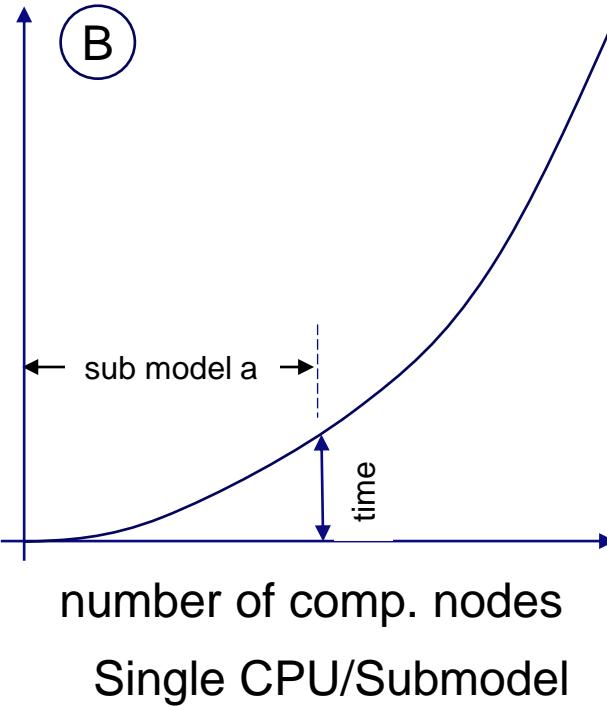
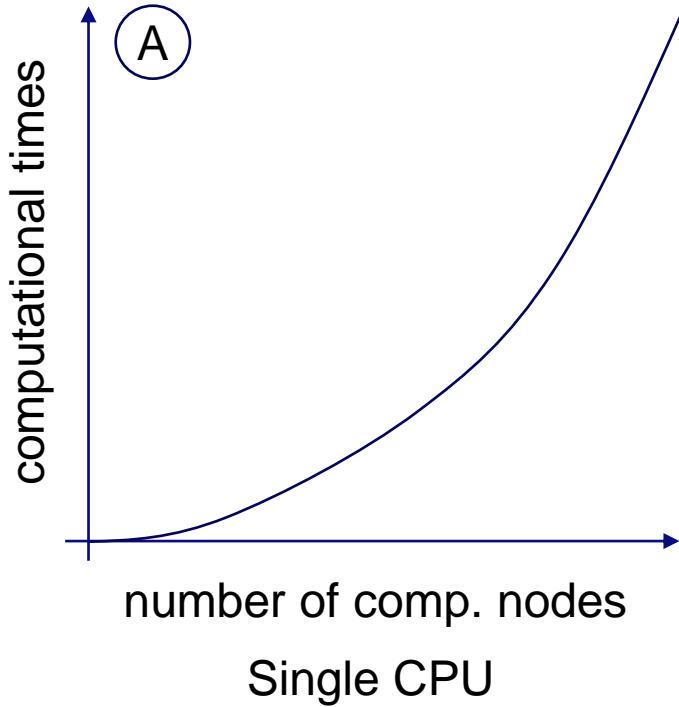
	Single CPU	20 CPU
Sensitivity	2050	427 days
Lambda	3	5 hours
Totally	10 cycles	~0.5 year

It never goes right the first, second, third time ...

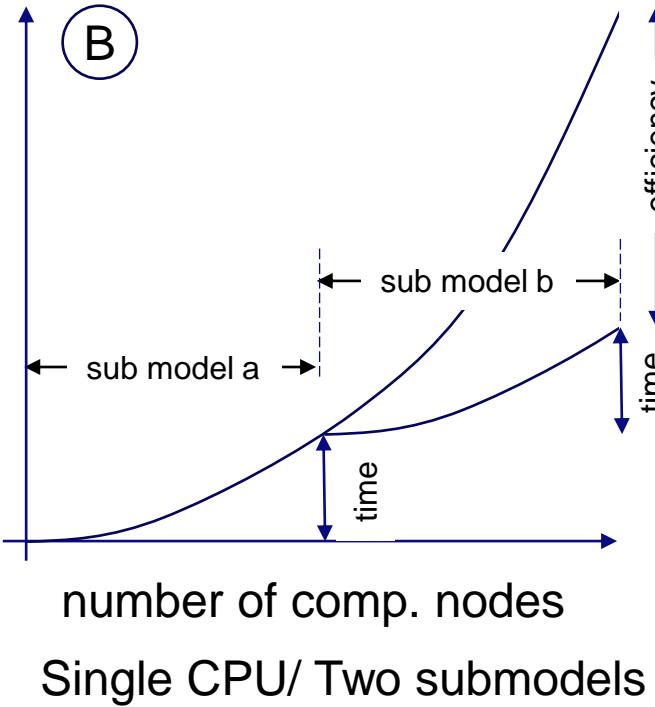
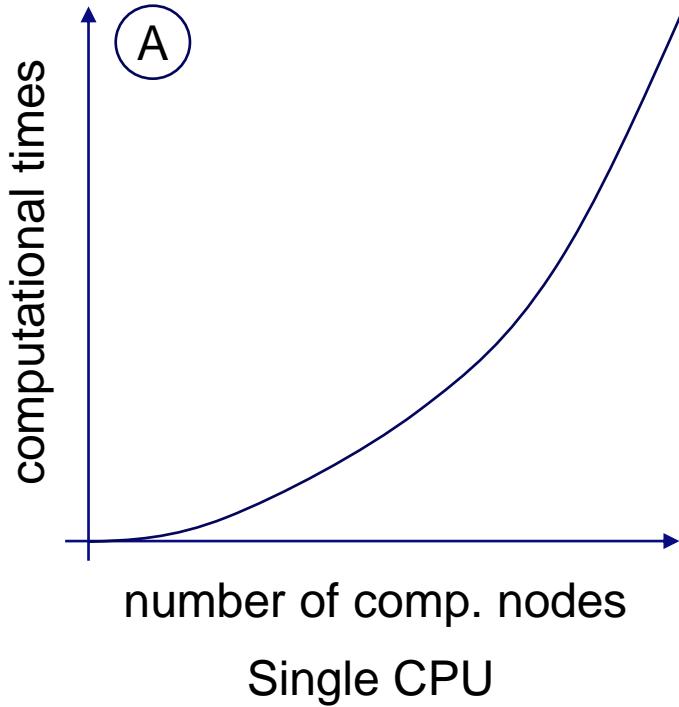
Runtimes of Models



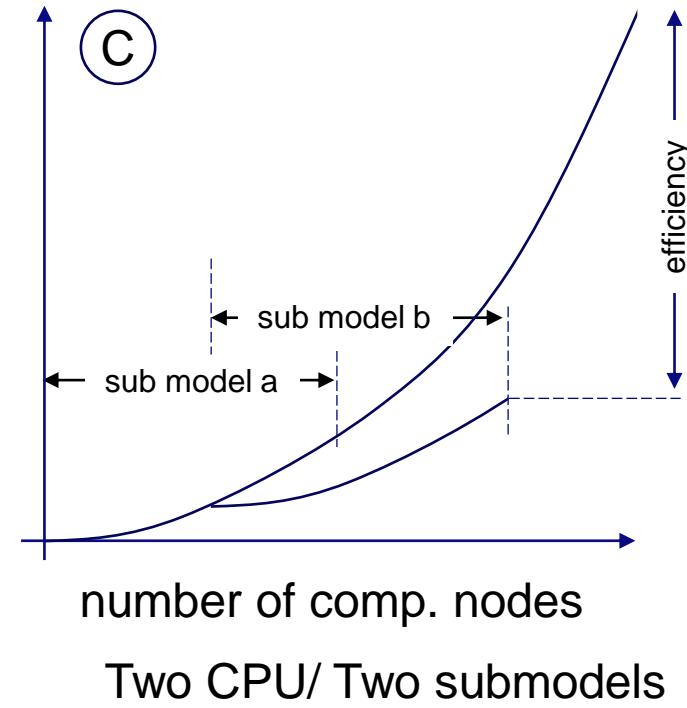
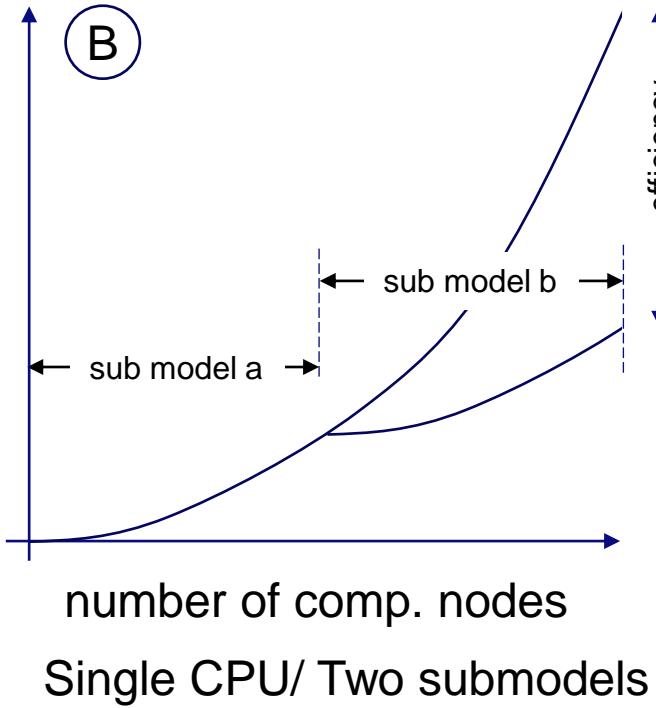
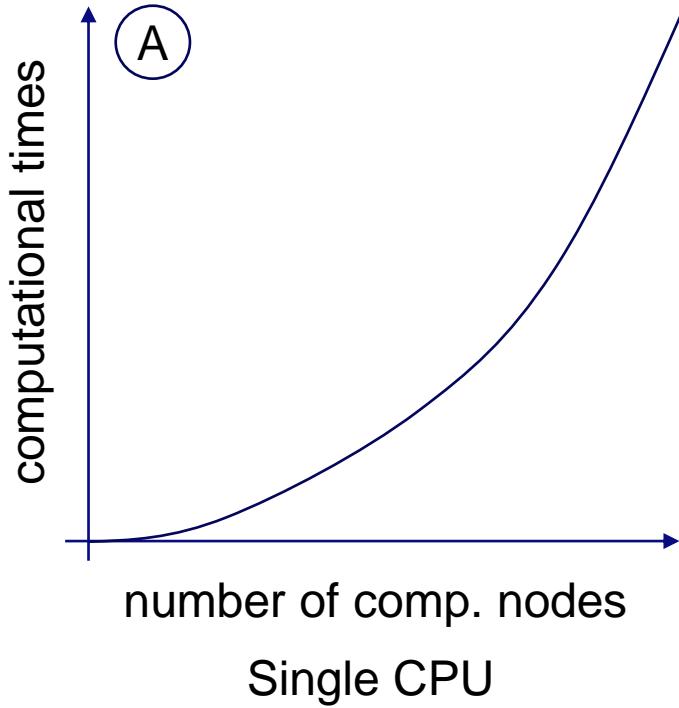
Runtimes of Models



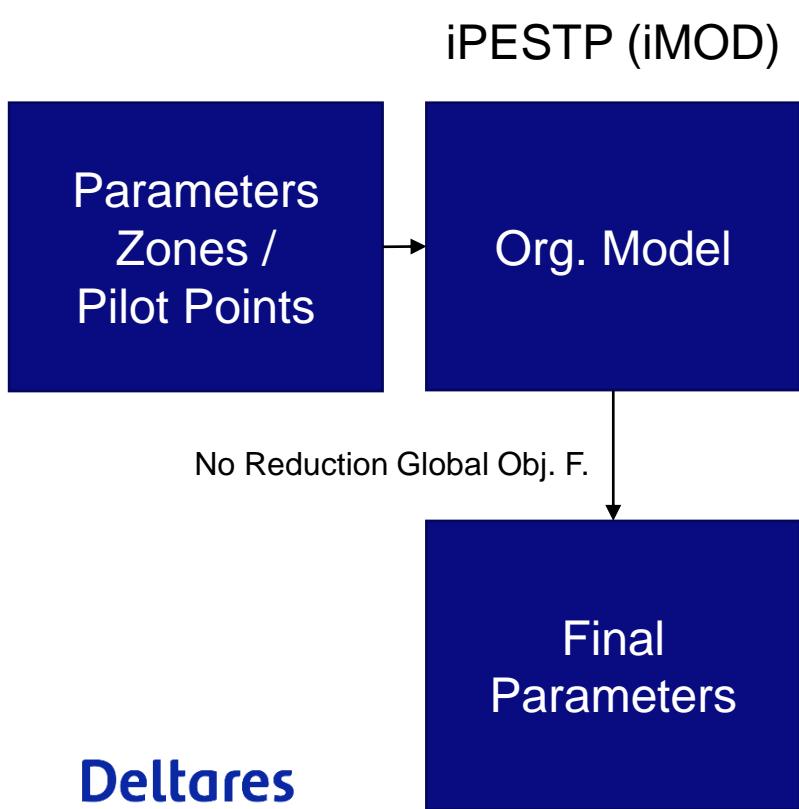
Runtimes of Models



Runtimes of Models

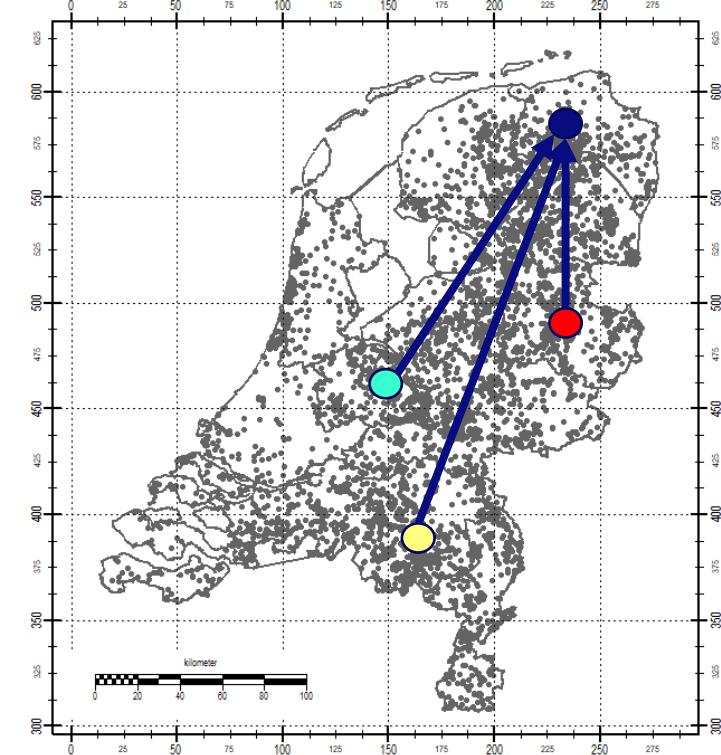
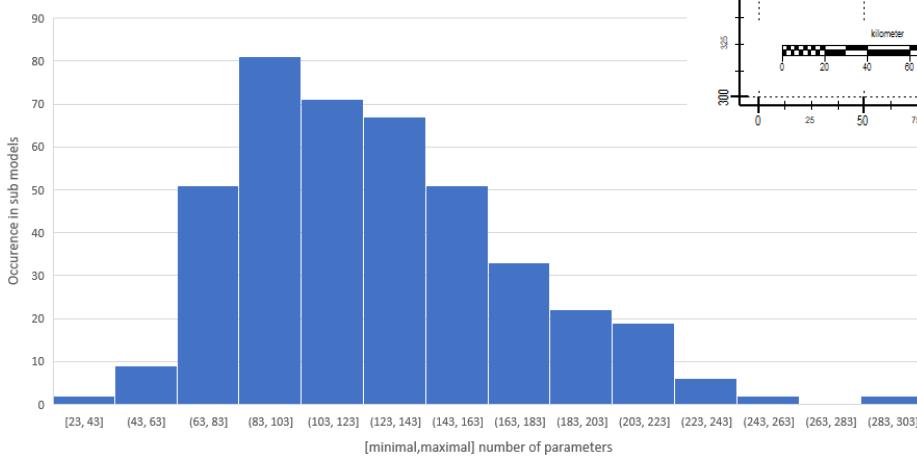


Submodel Model Optimization Method (SMOM)



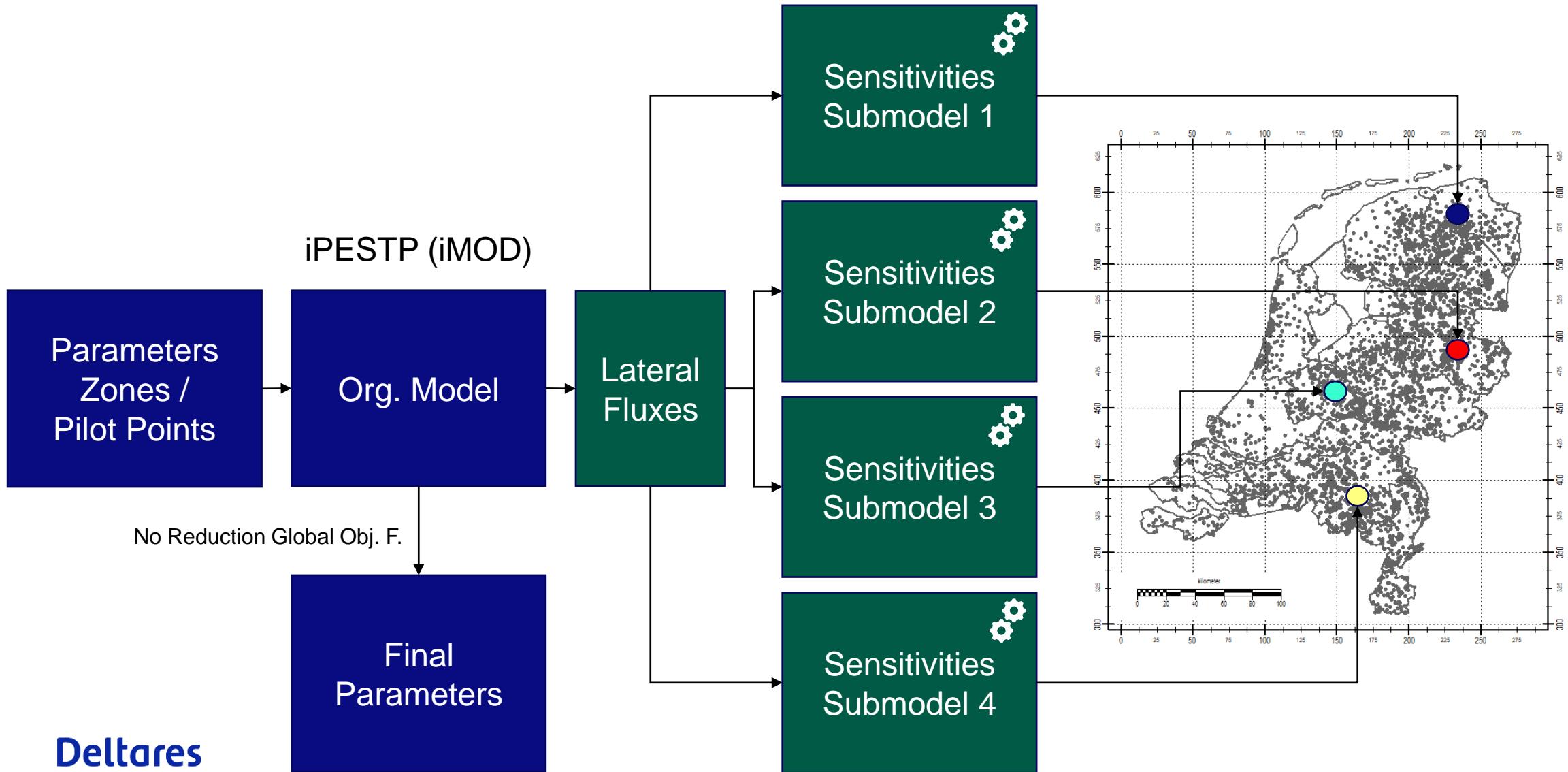
Questions:

- Effects groundwater in Groningen that of Twente, Utrecht or Brabant?
- Can sensitivities be approximated by smaller models?
- Why compute Groningen for parameters that do not extent in that region?

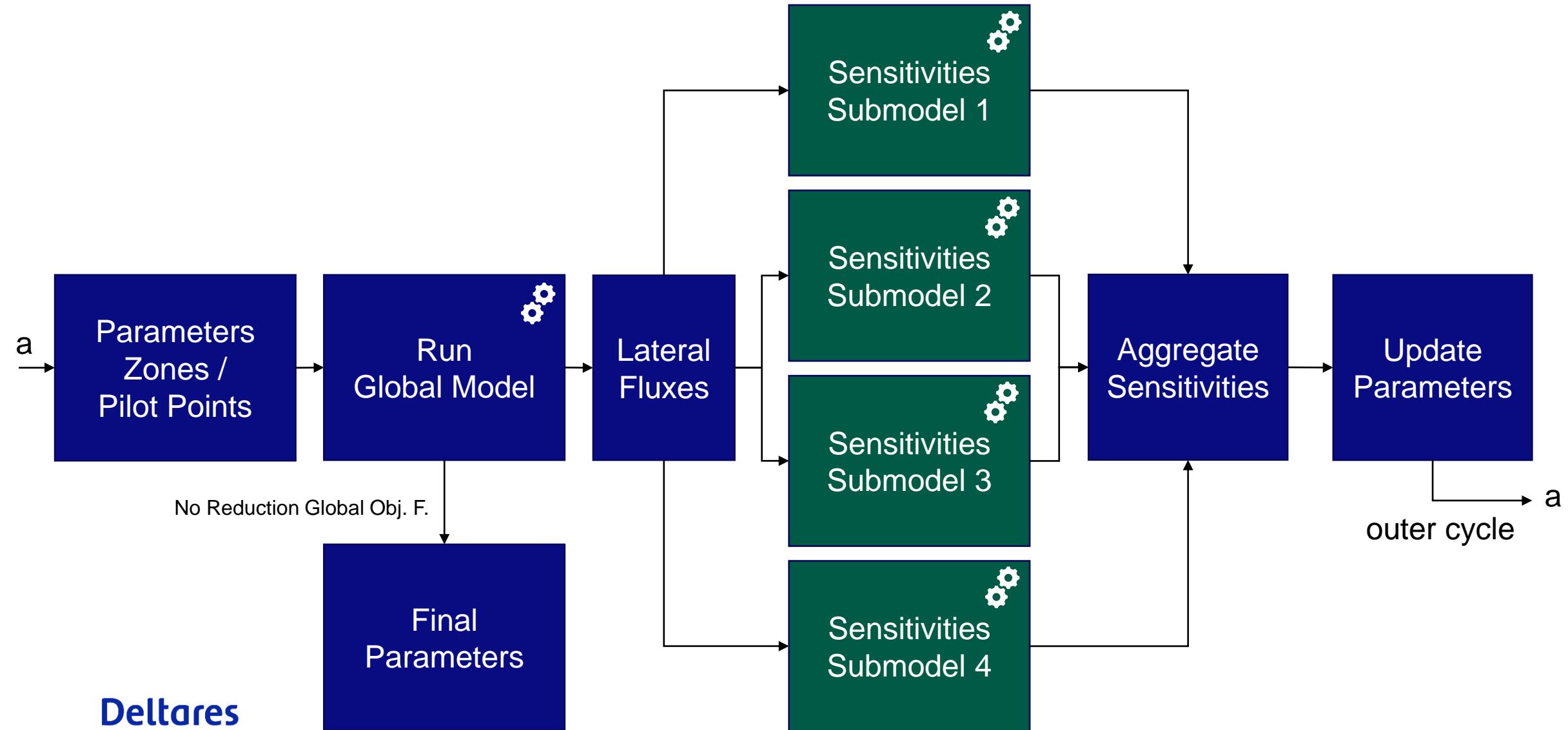


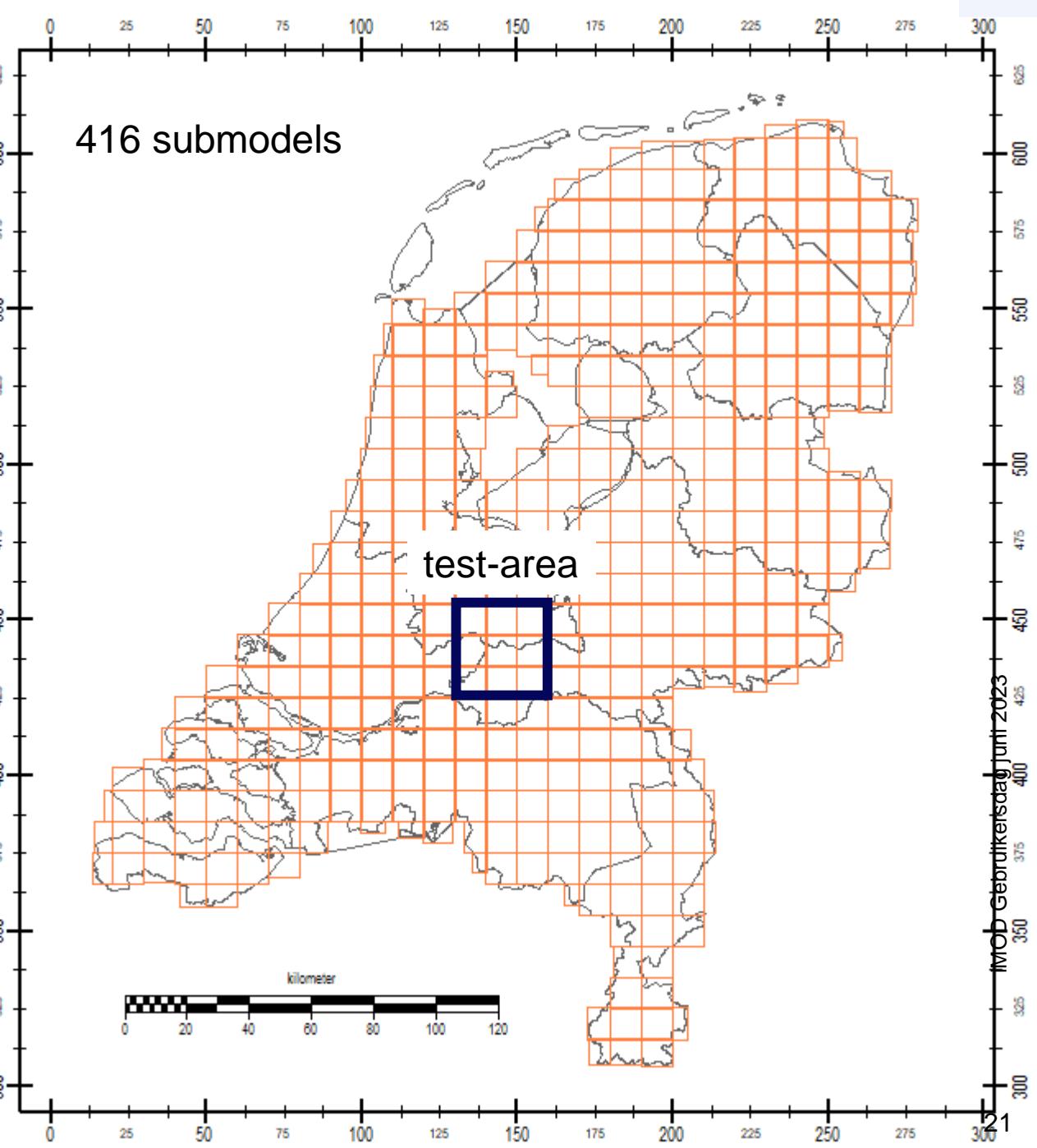
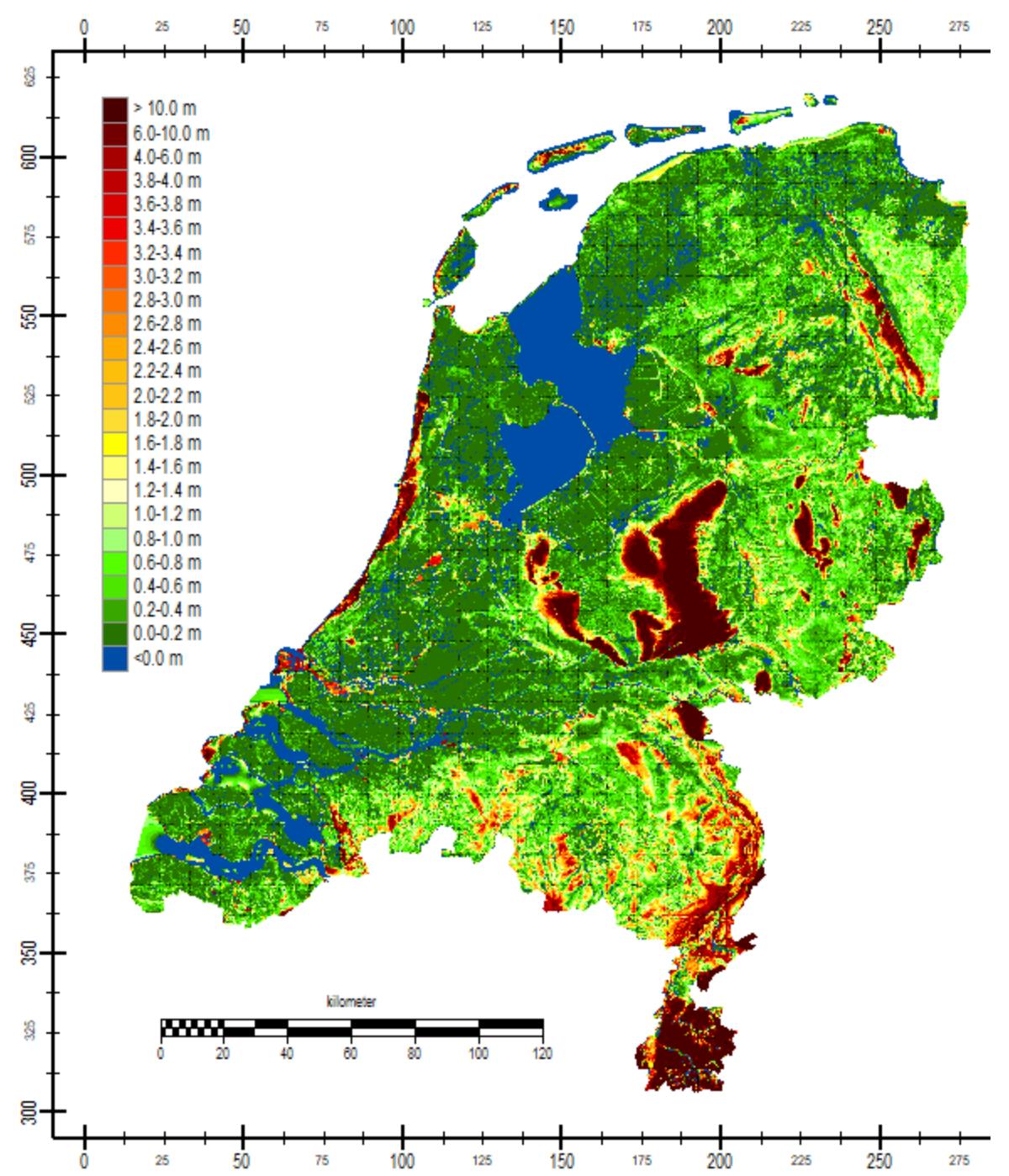
number of
parameter per
submodel

Submodel Model Optimization Method (SMOM)

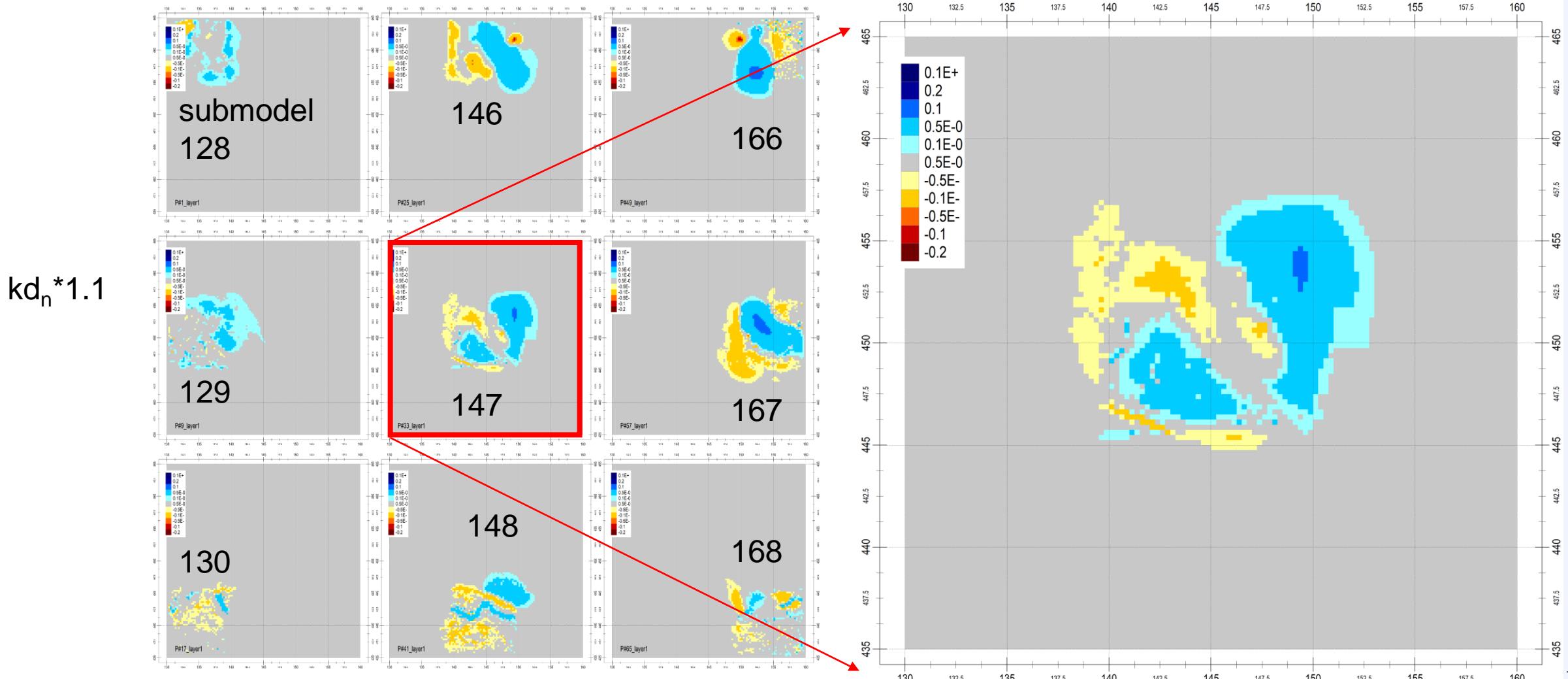


Submodel Model Optimization Method (SMOM)

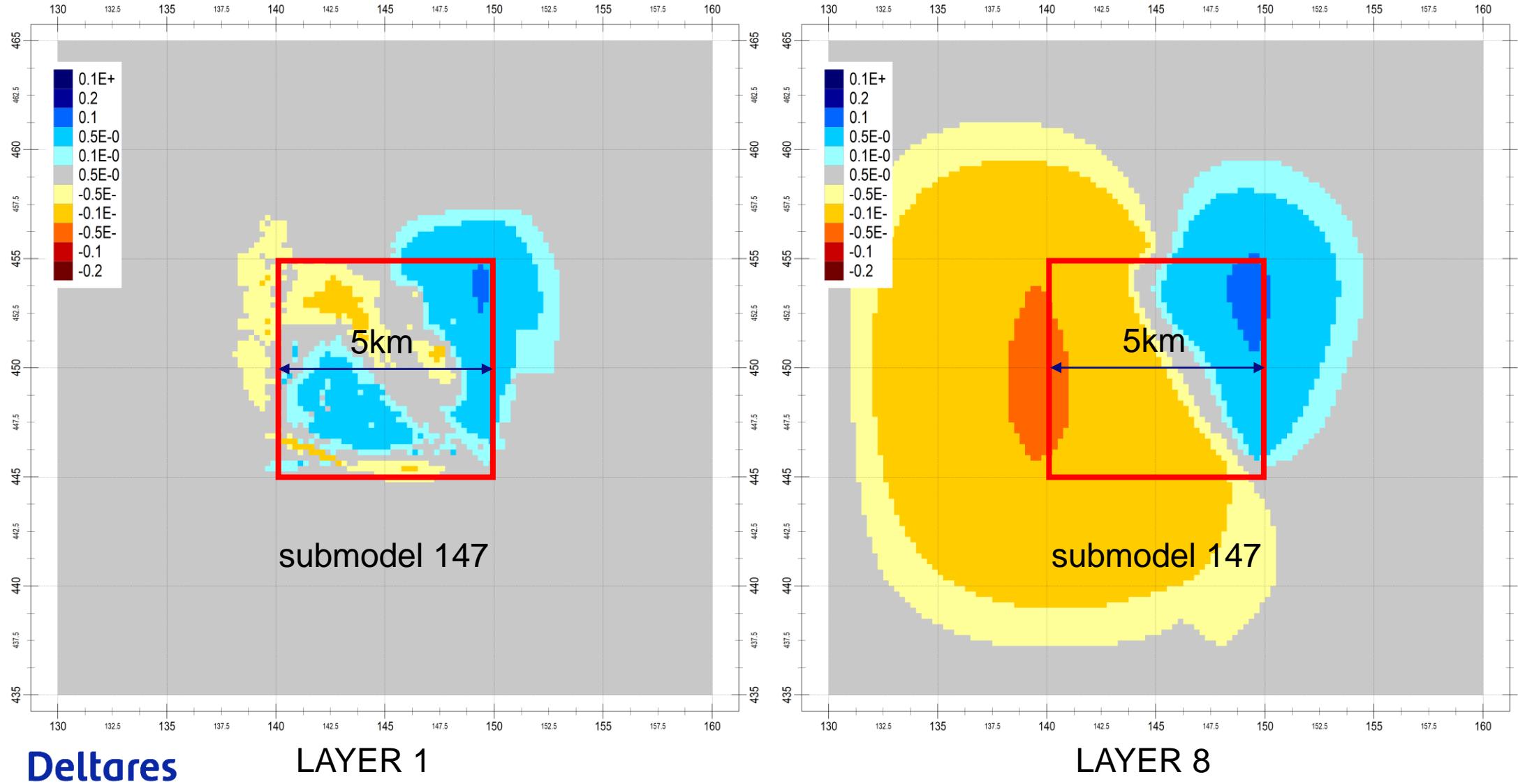




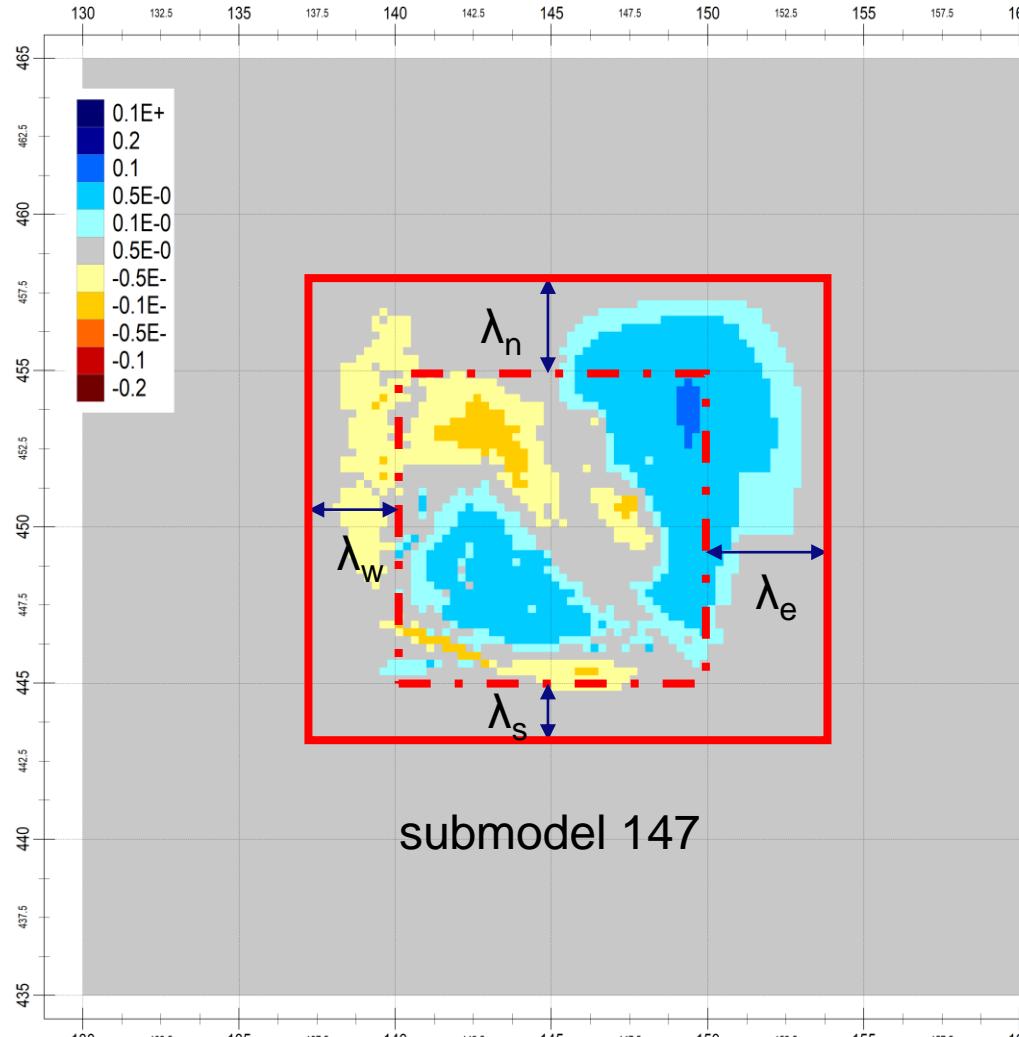
Submodel Model Optimization Method (SMOM)



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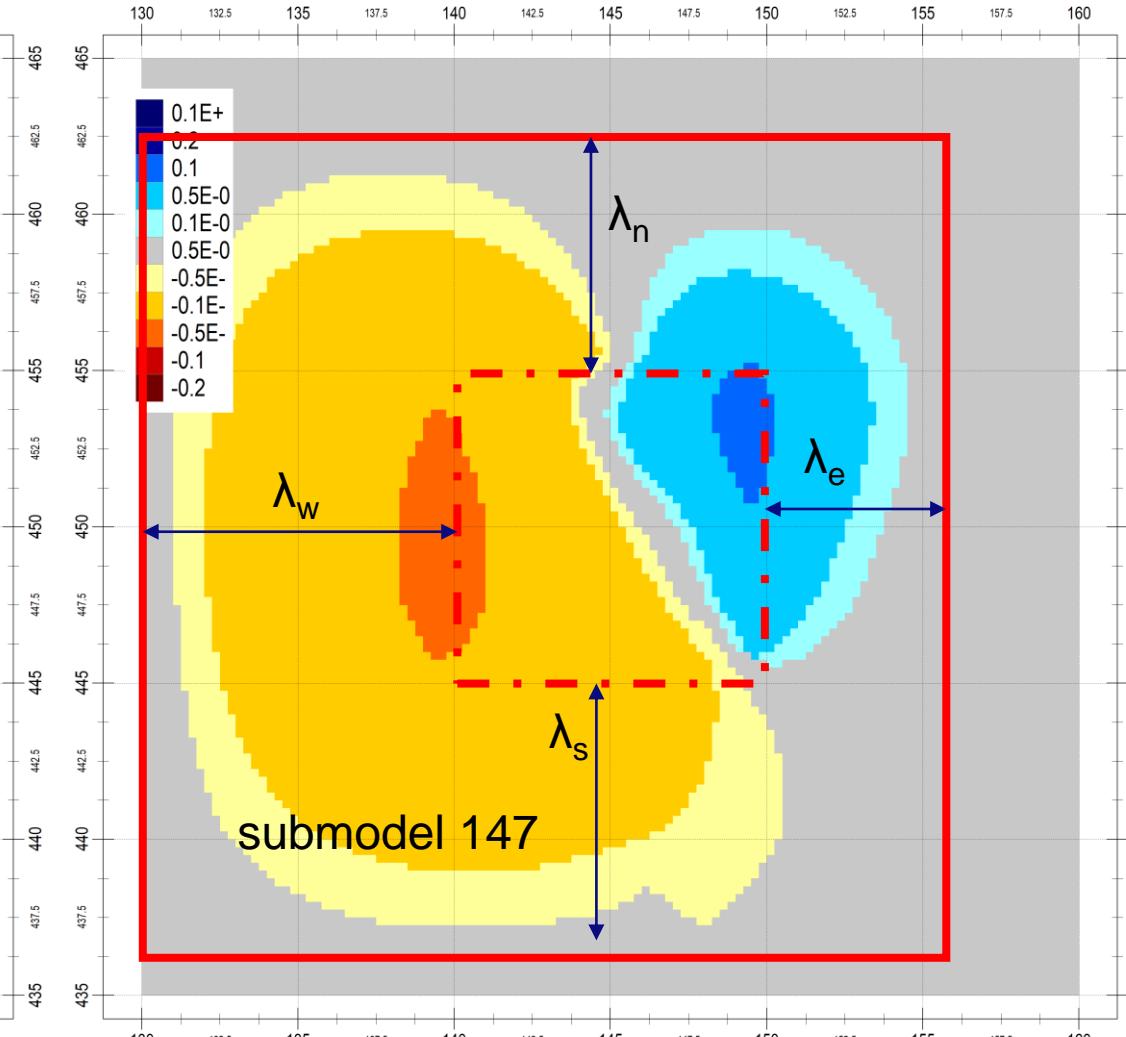


Submodel Model Optimization Method (SMOM)



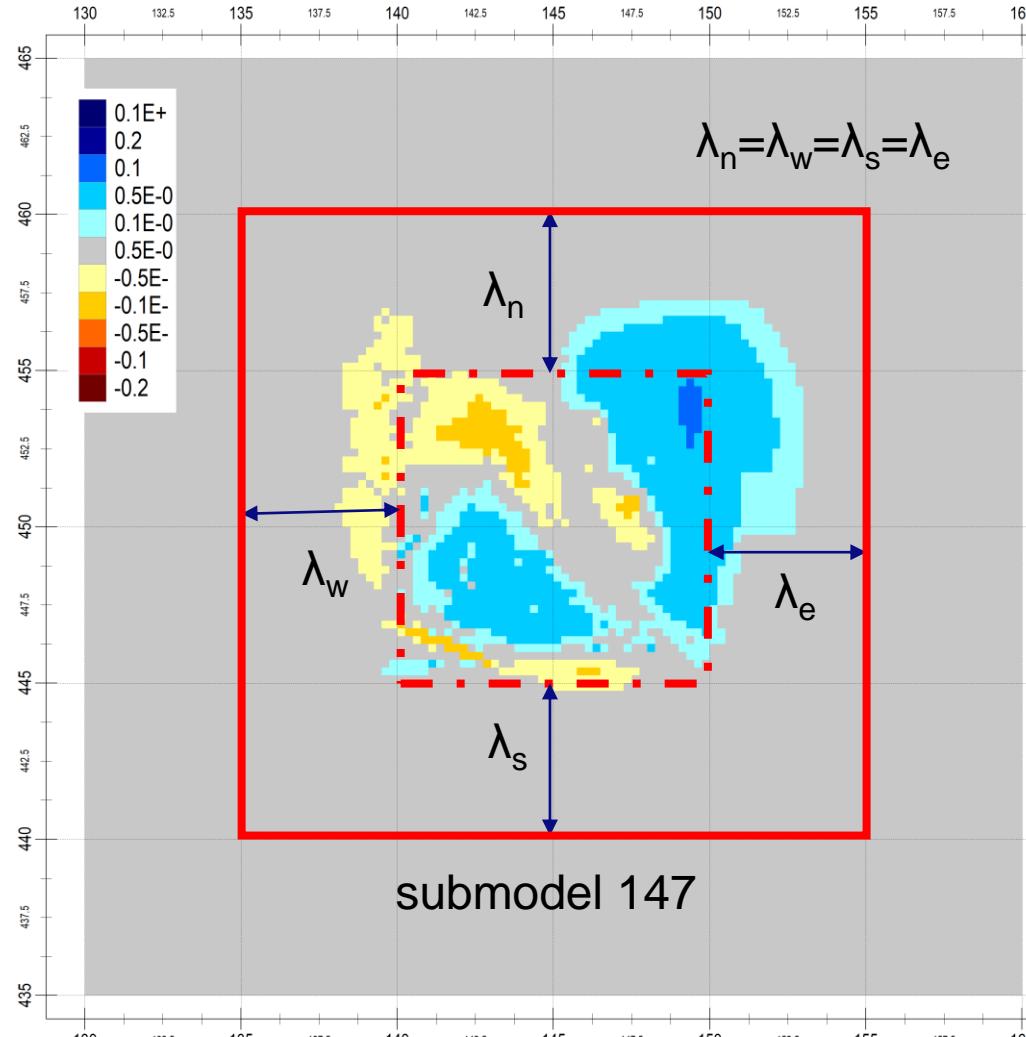
Deltas

LAYER 1



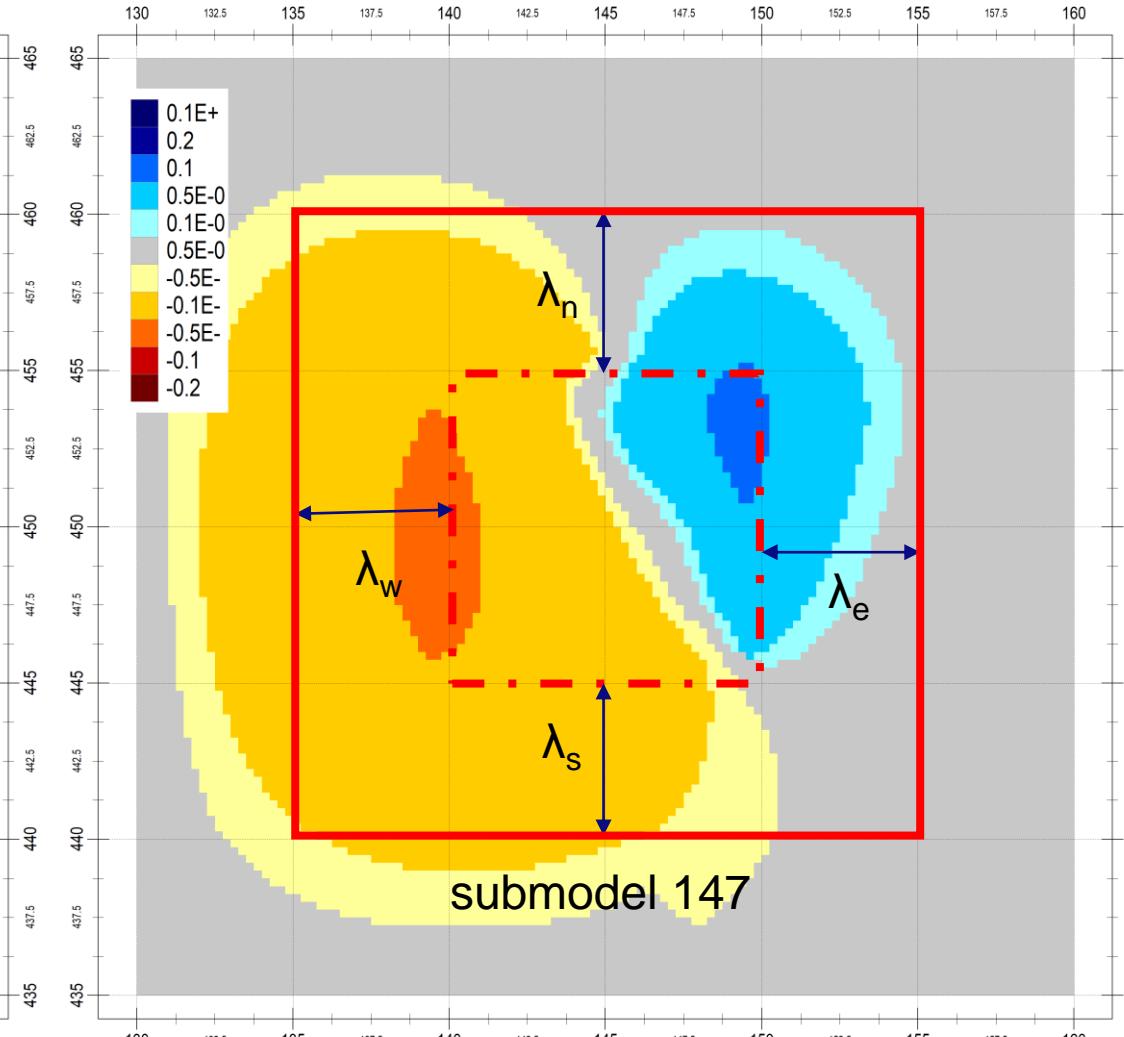
LAYER 8

Submodel Model Optimization Method (SMOM)



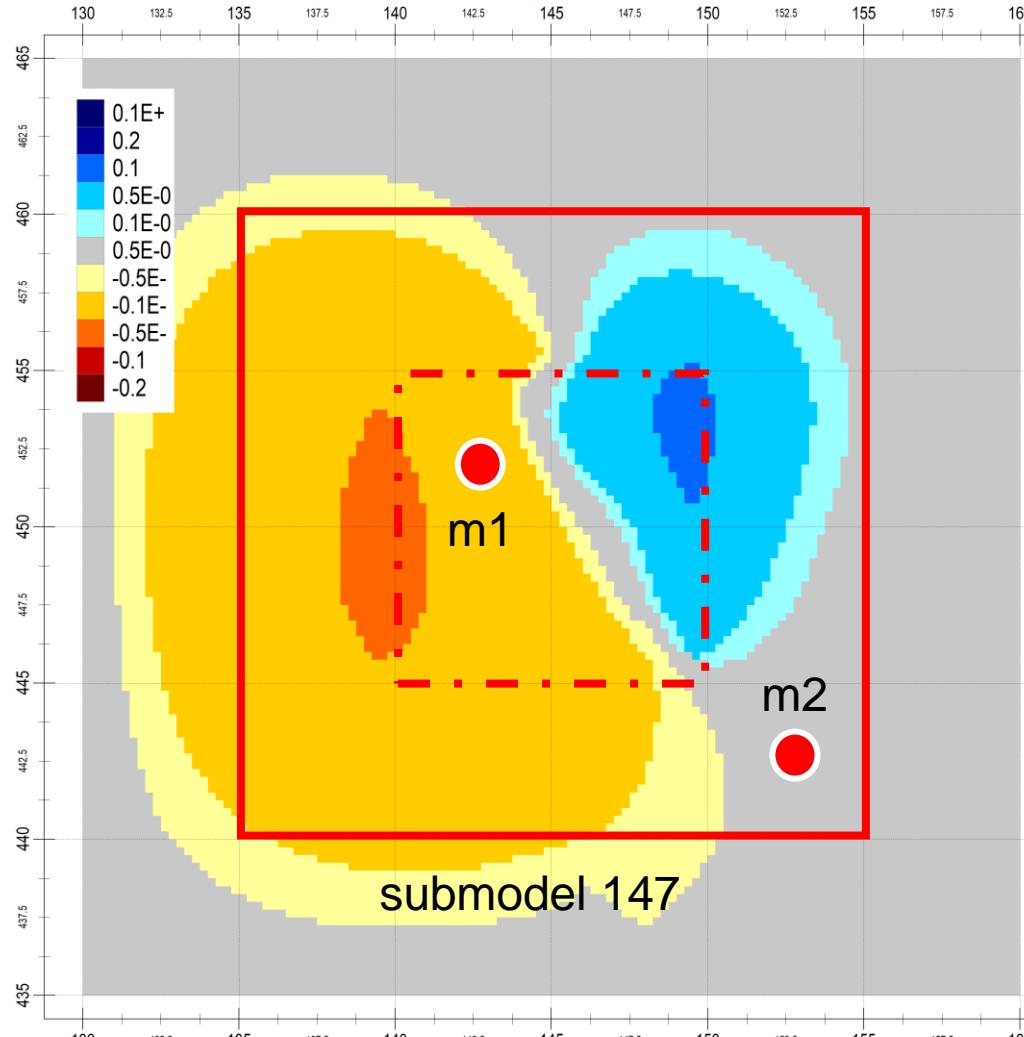
Deltas

LAYER 1



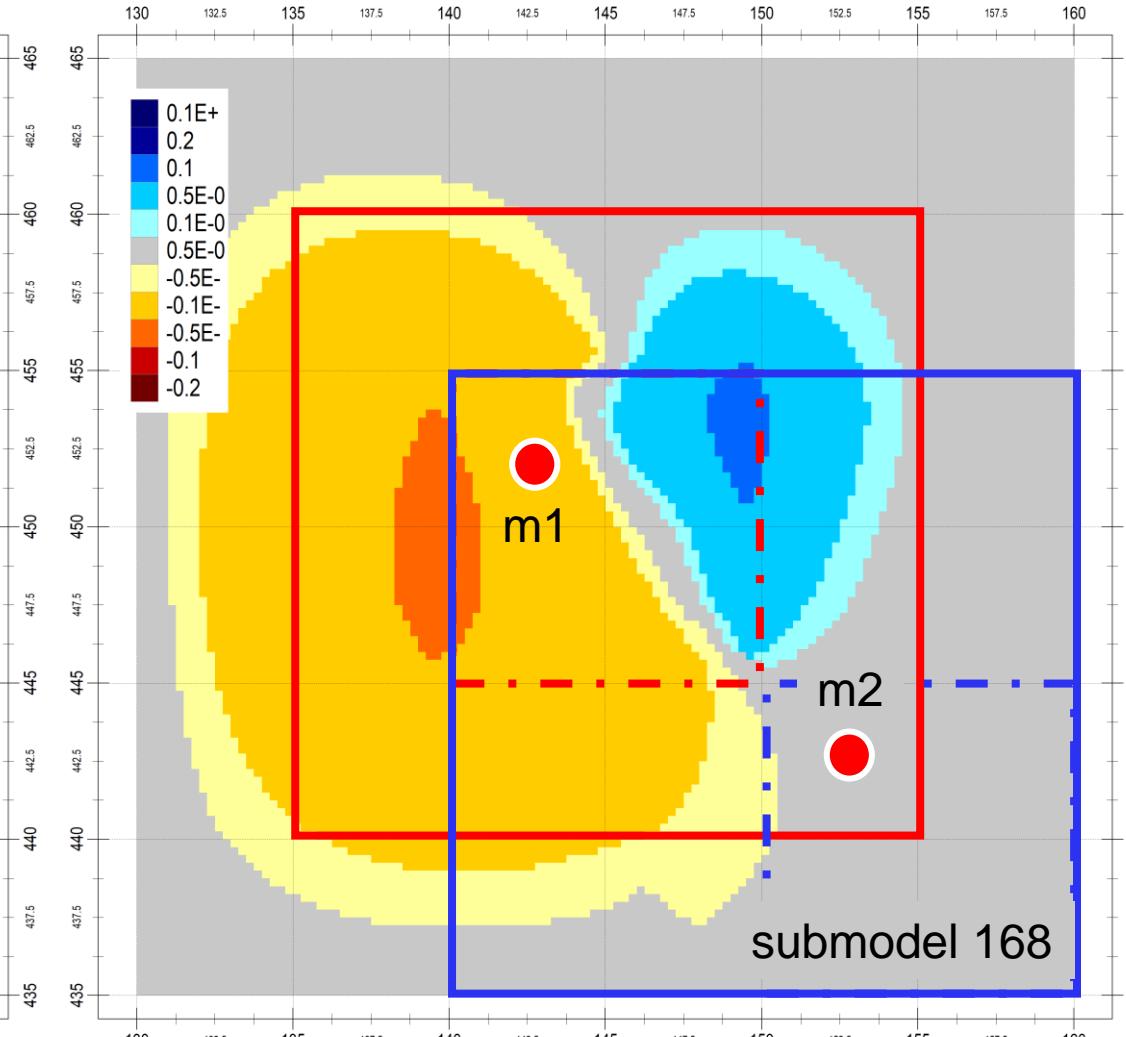
LAYER 8

Submodel Model Optimization Method (SMOM)



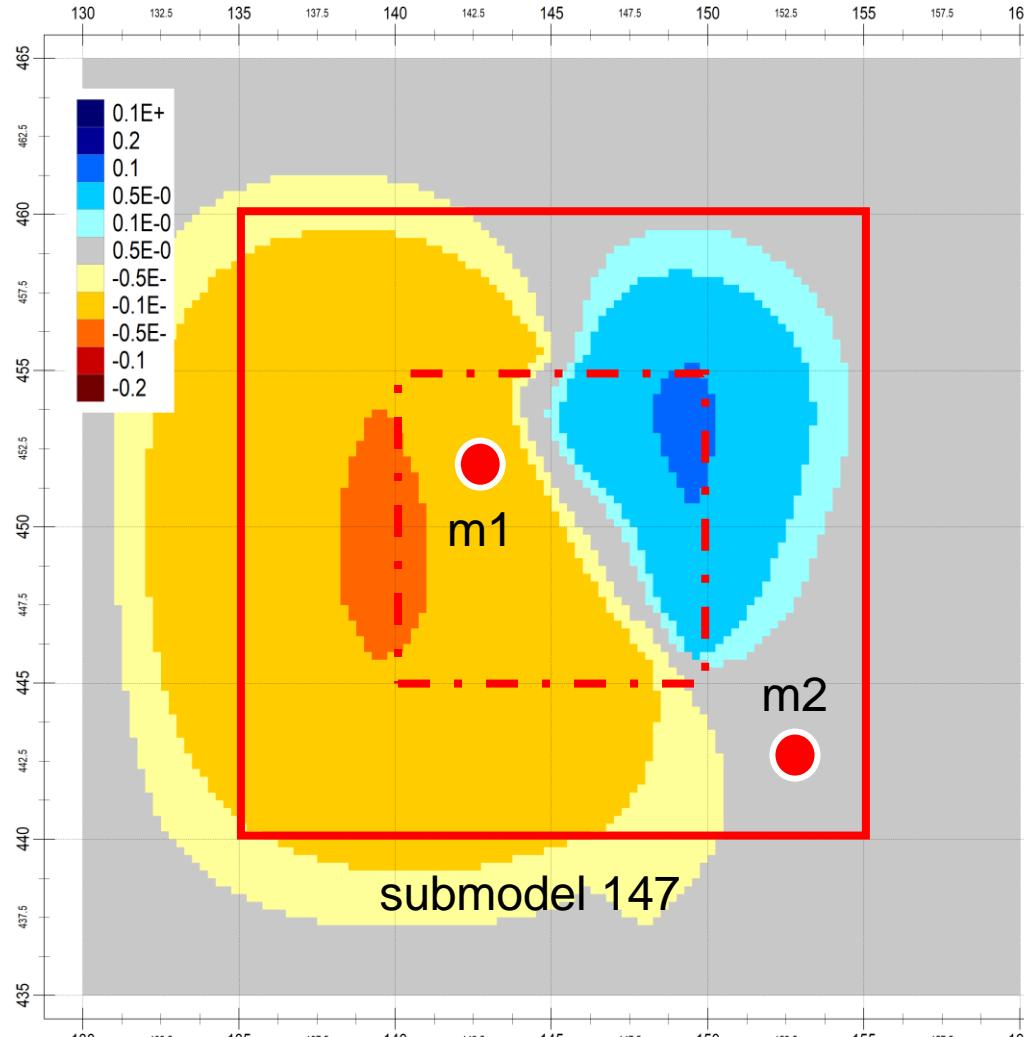
Deltares

LAYER 8



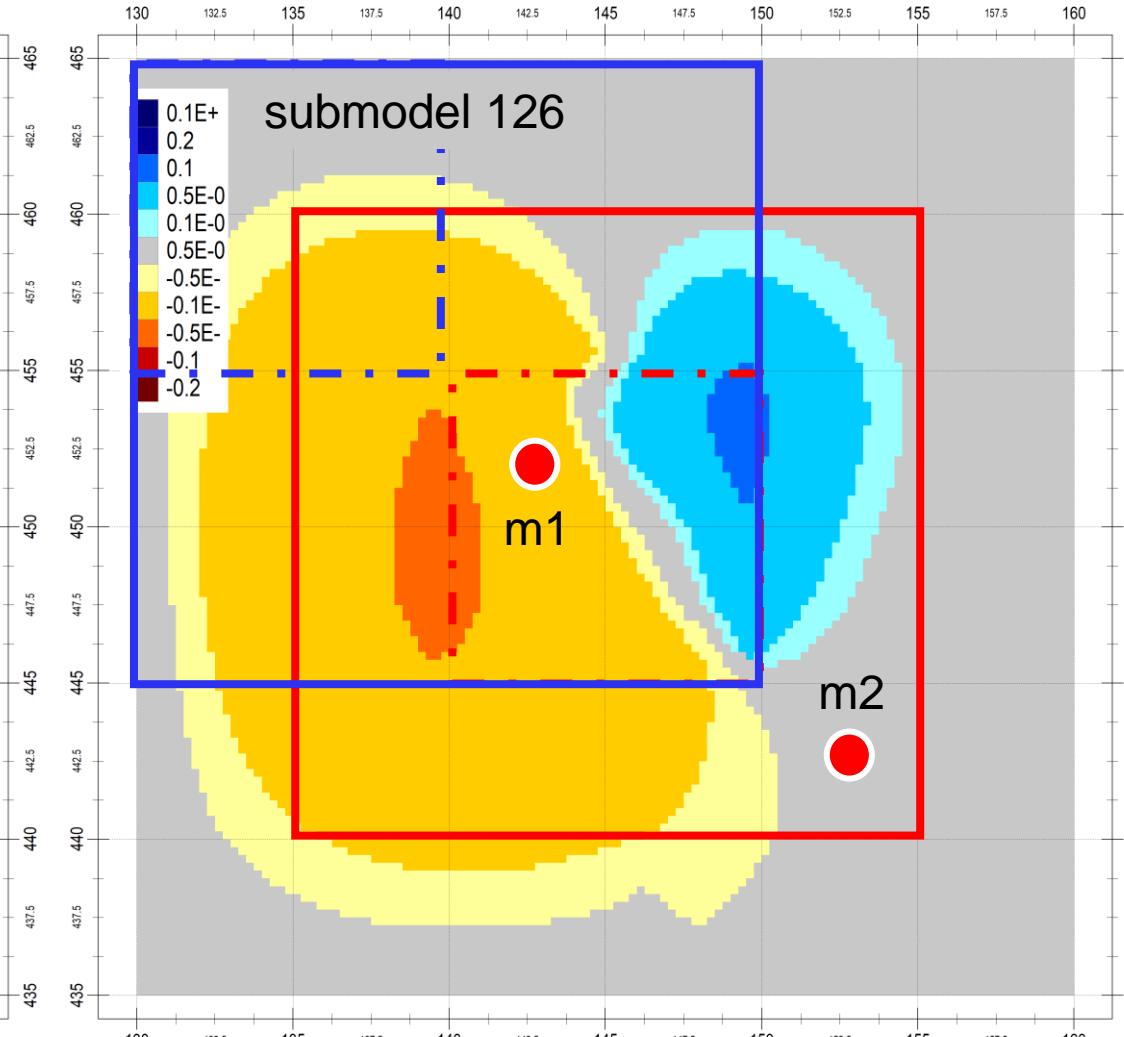
LAYER 8

Submodel Model Optimization Method (SMOM)



Deltares

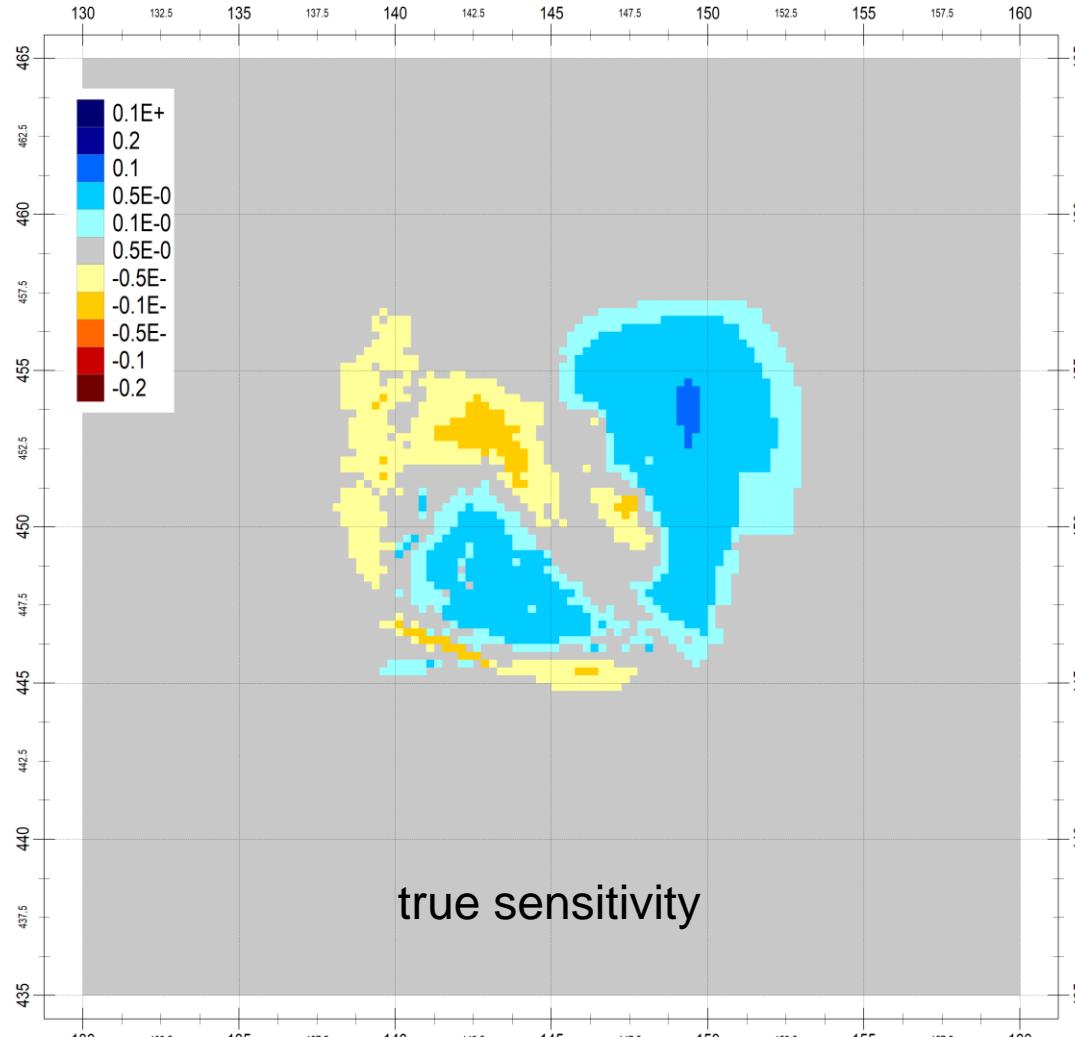
LAYER 8



LAYER 8

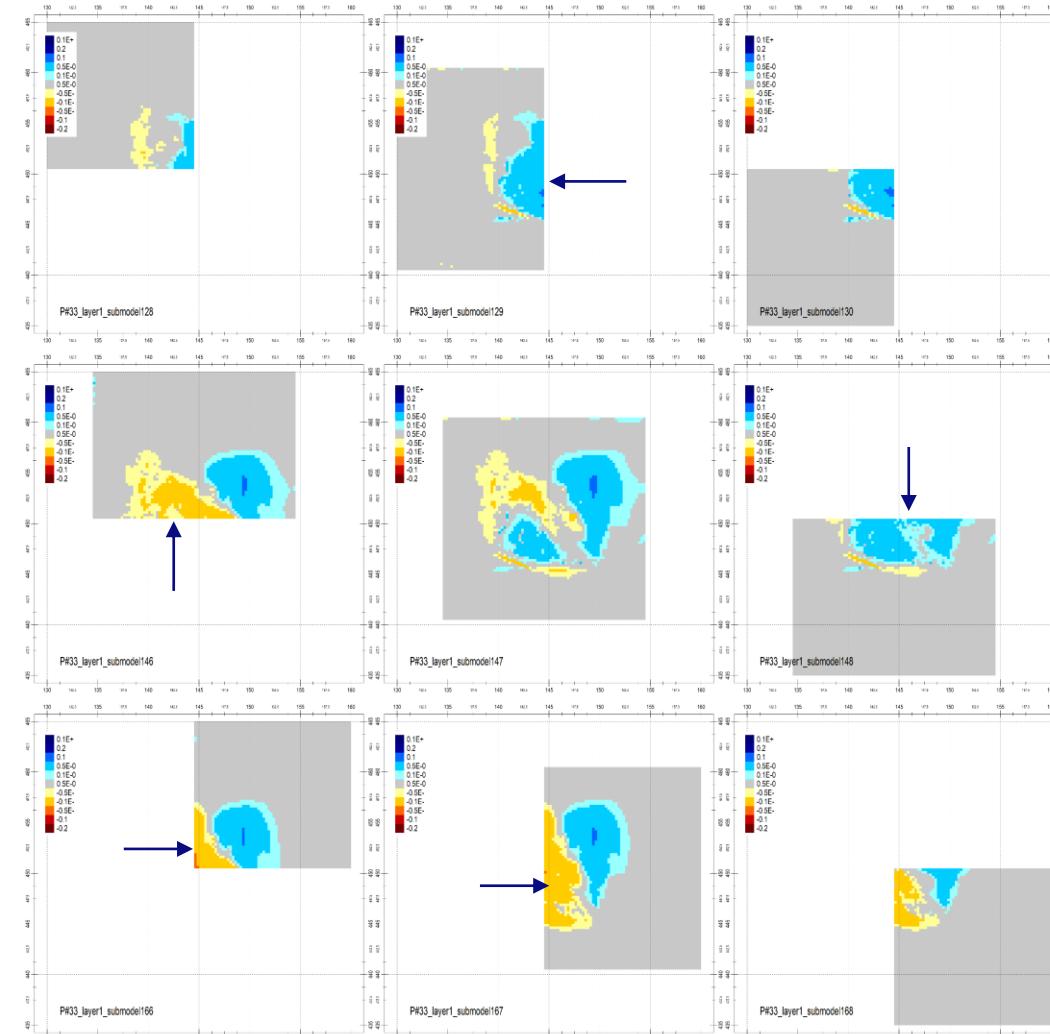
SMOM – Q-EDGE

sensitivities per submodel

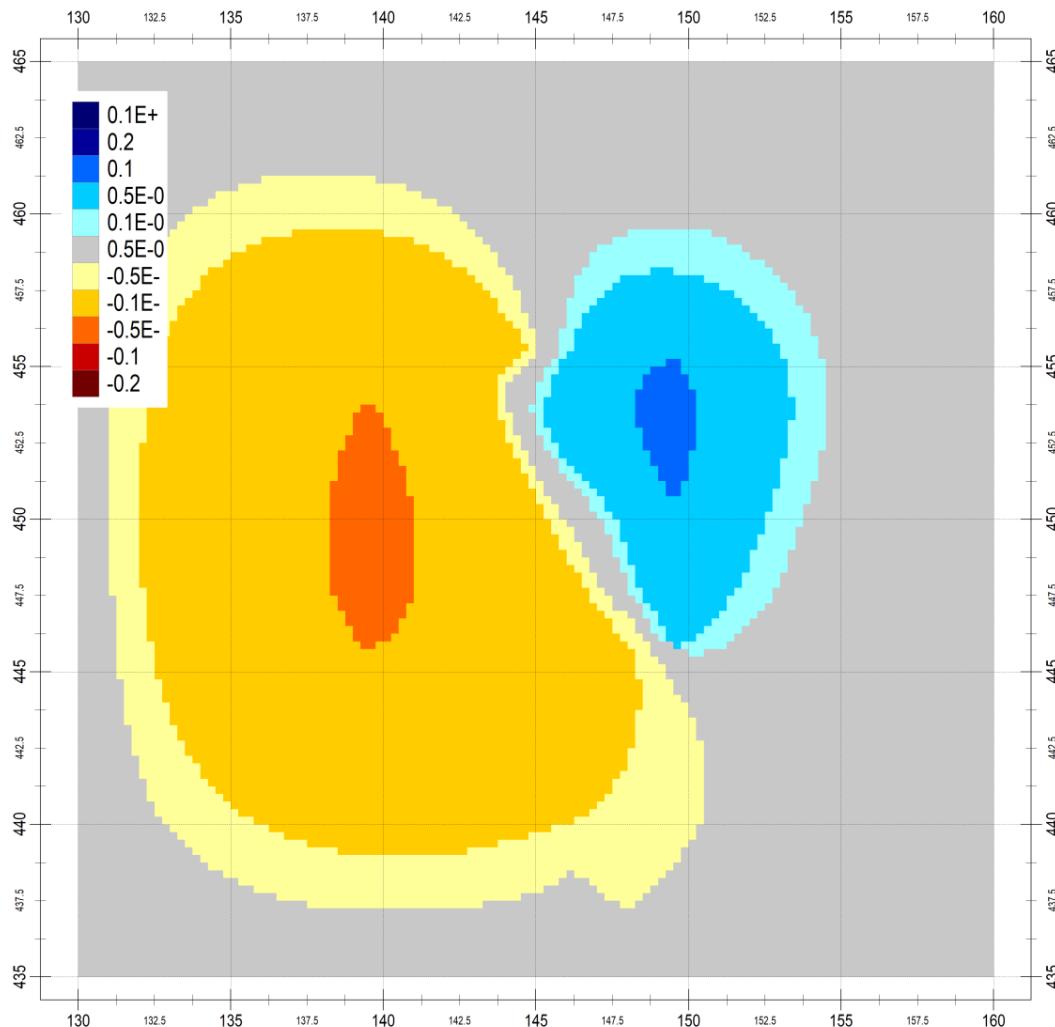


Deltares

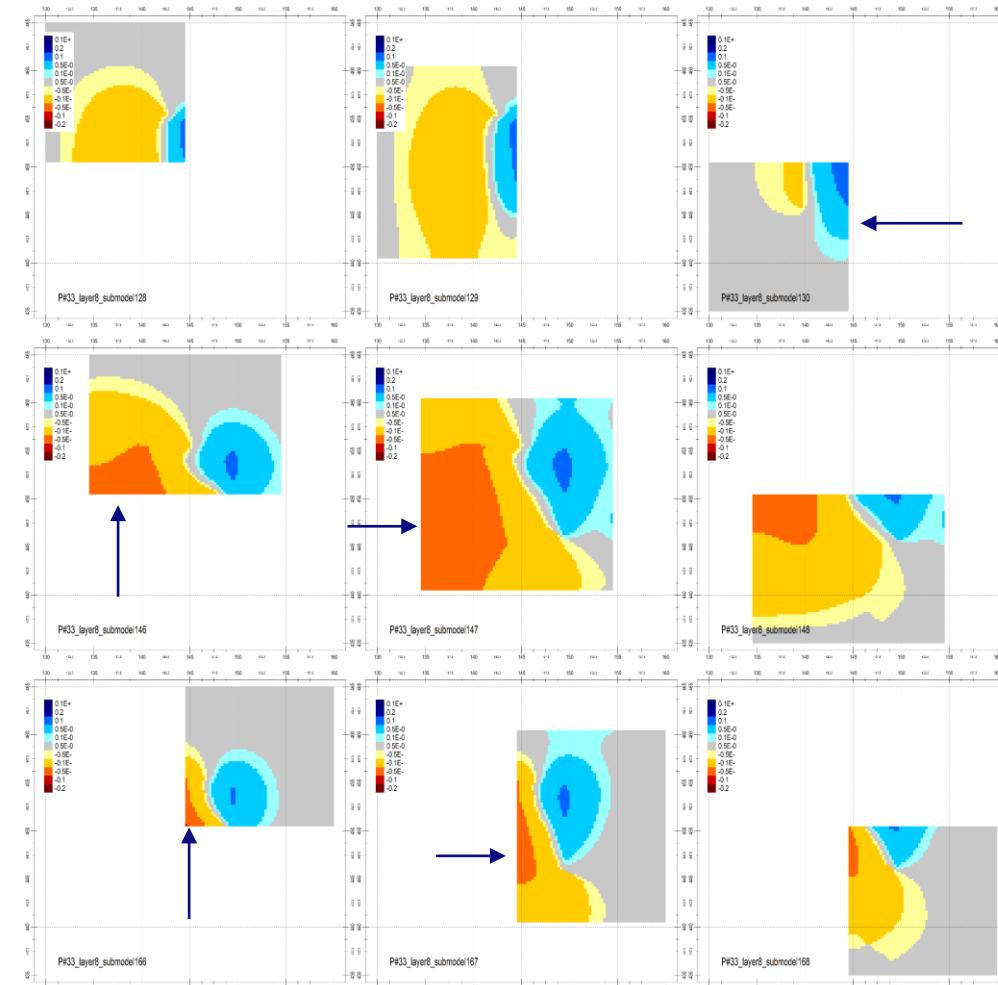
LAYER 1



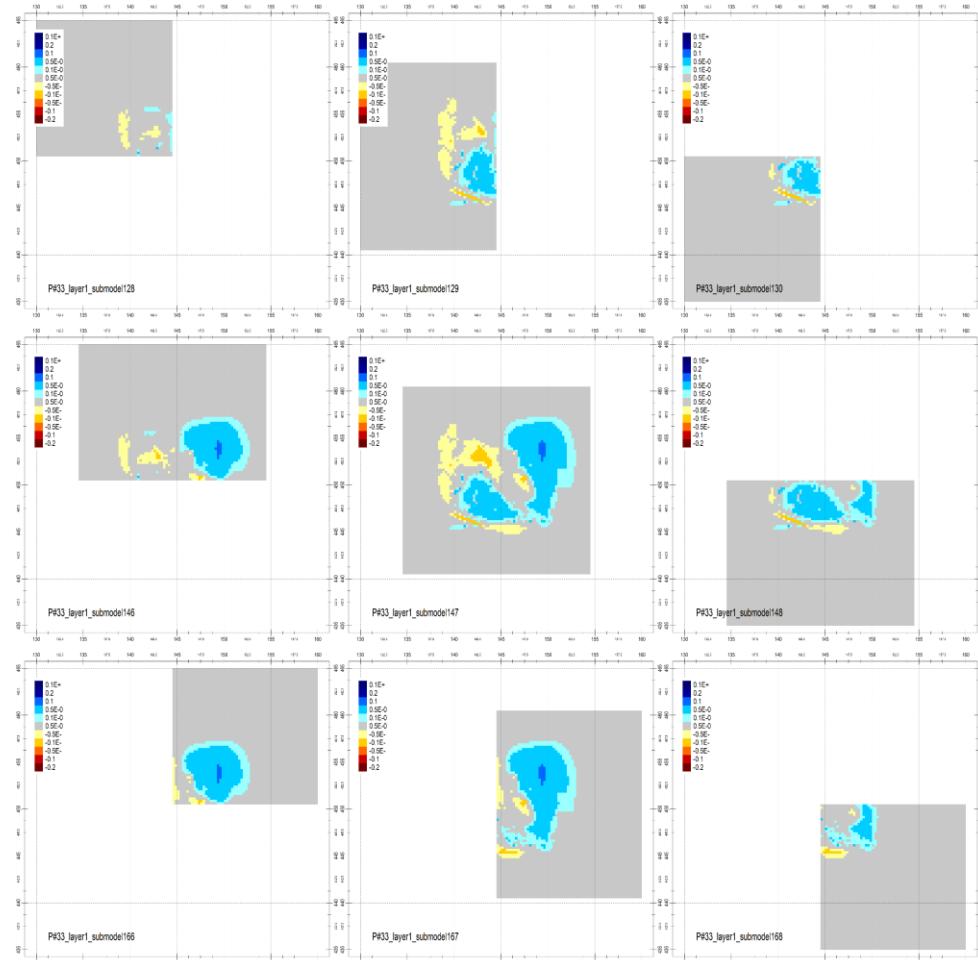
SMOM – Q-EDGE



Deltas

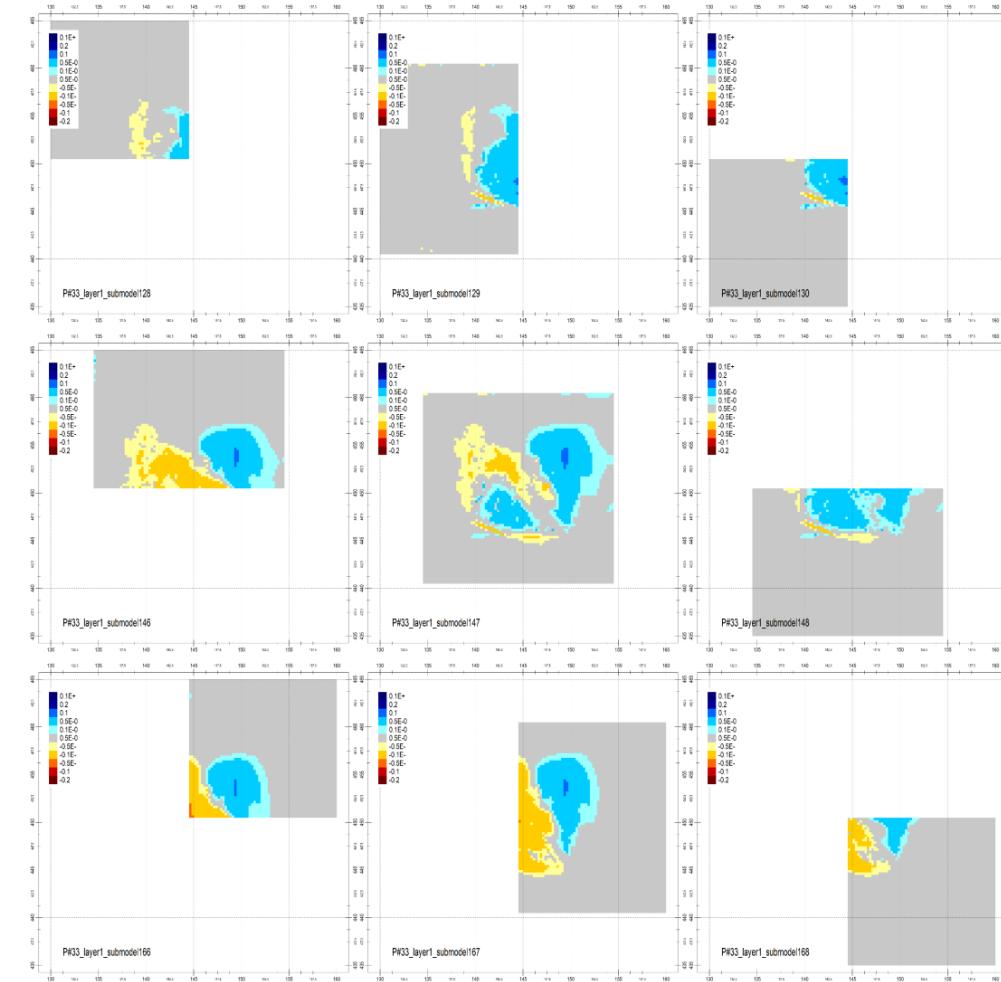


SMOM – GHB-EDGE $\text{cond} = q/dh$



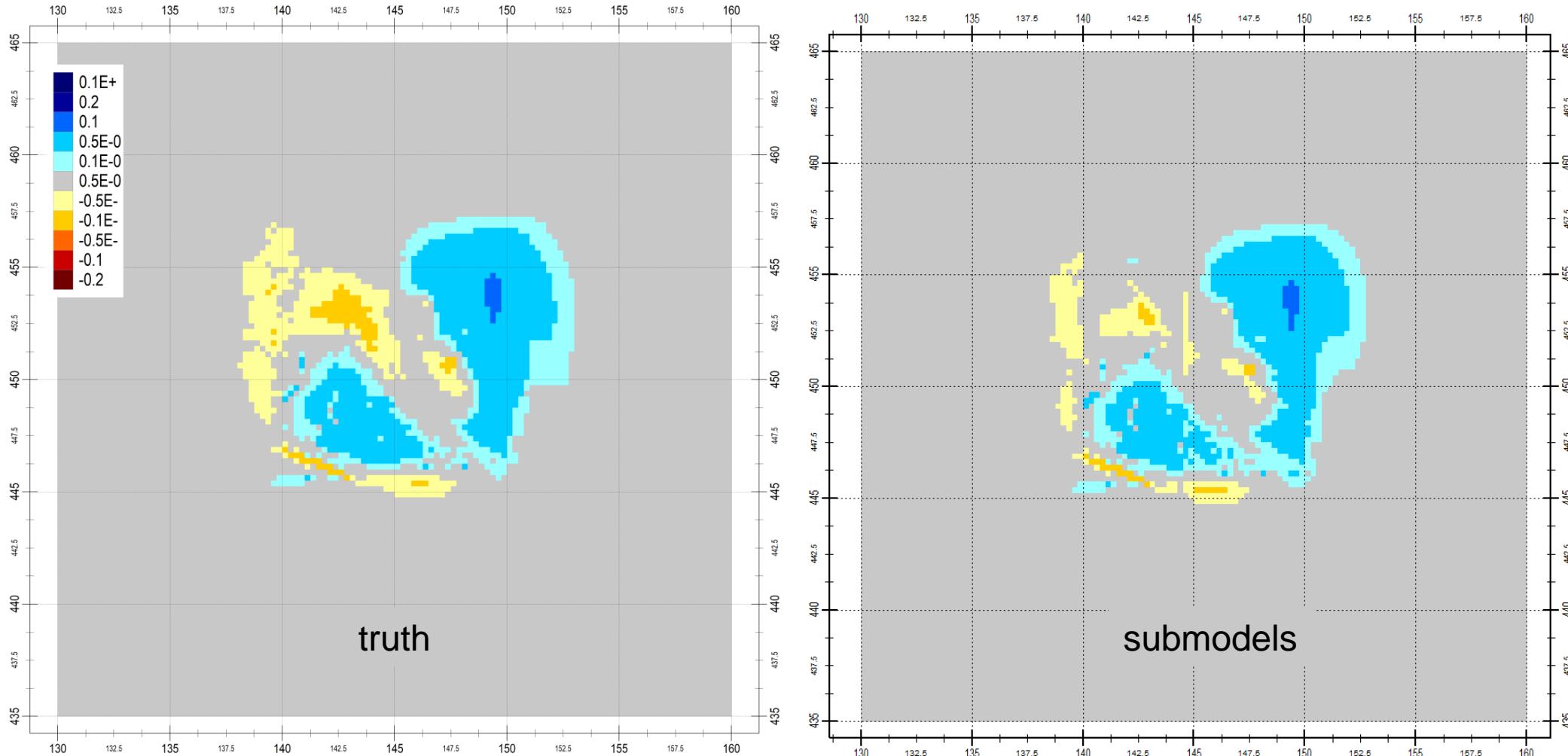
GHB-edge

Deltares

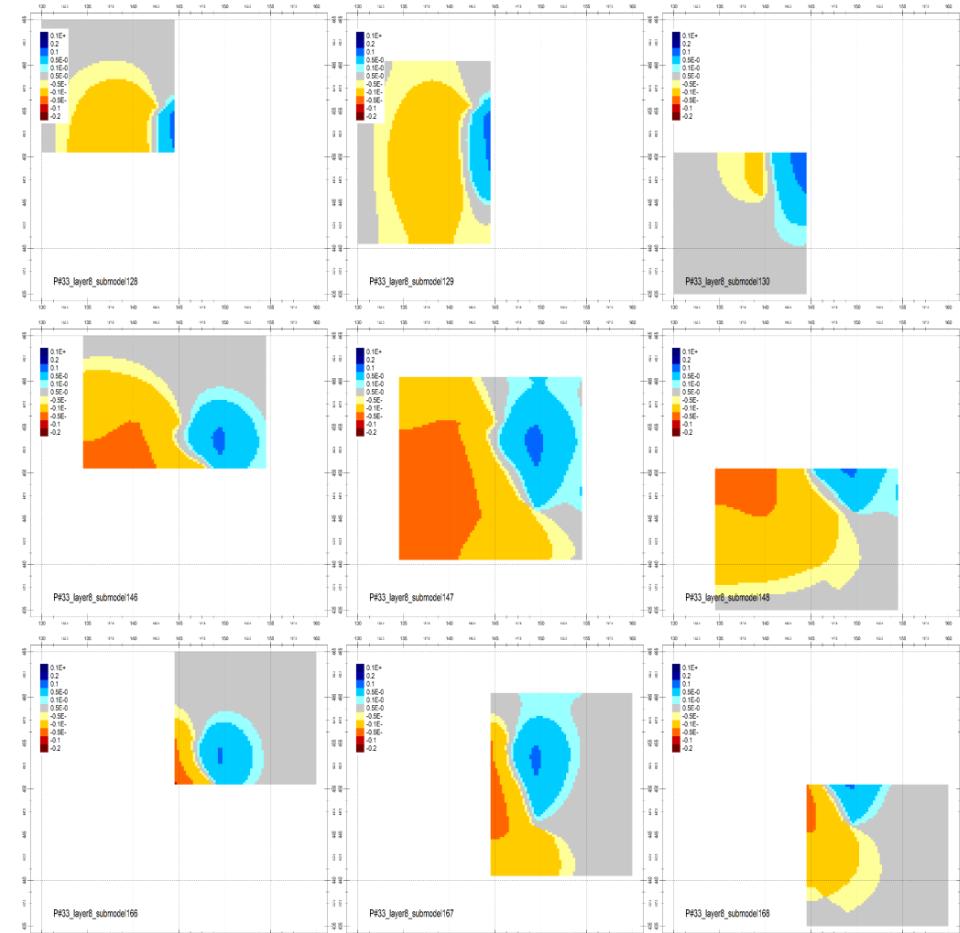
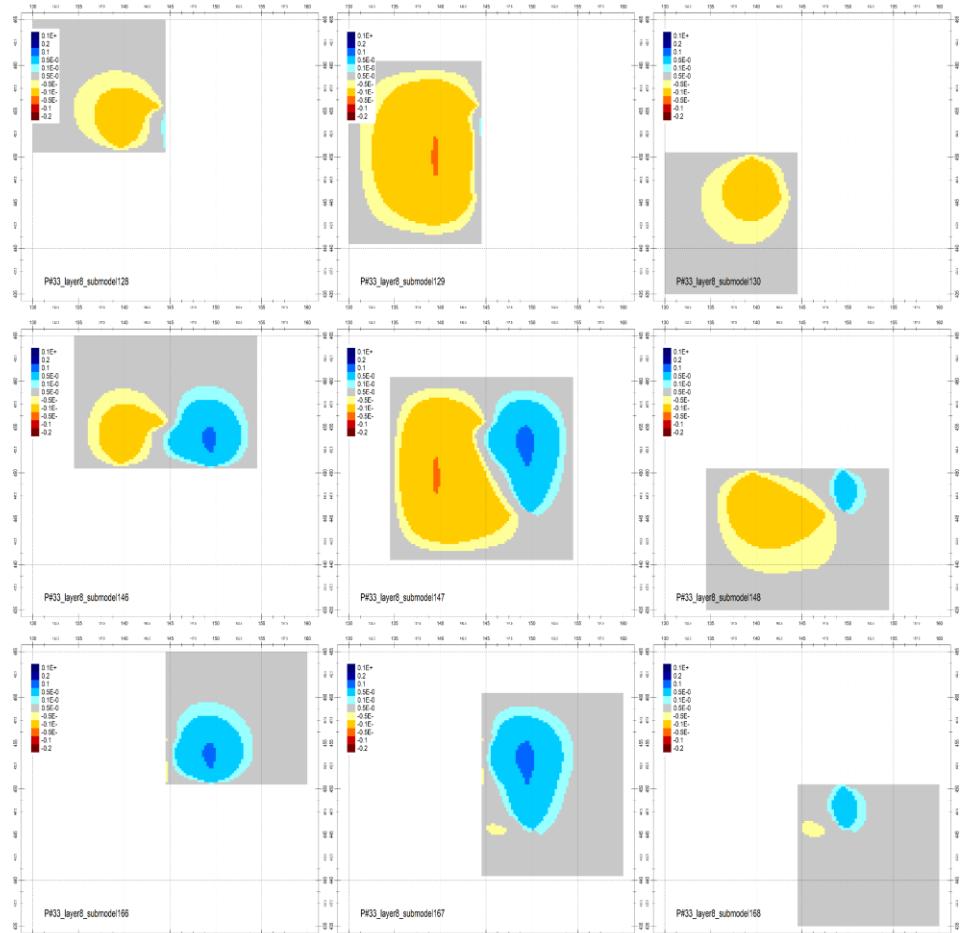


Q-edge

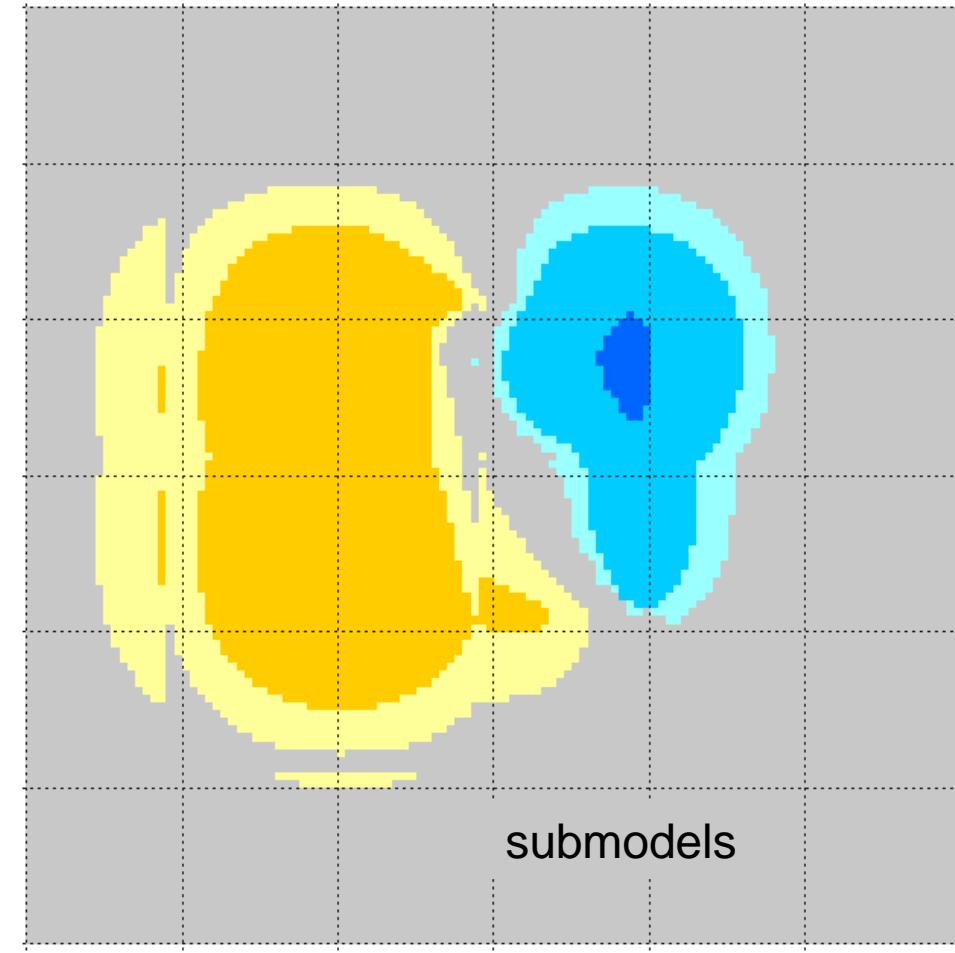
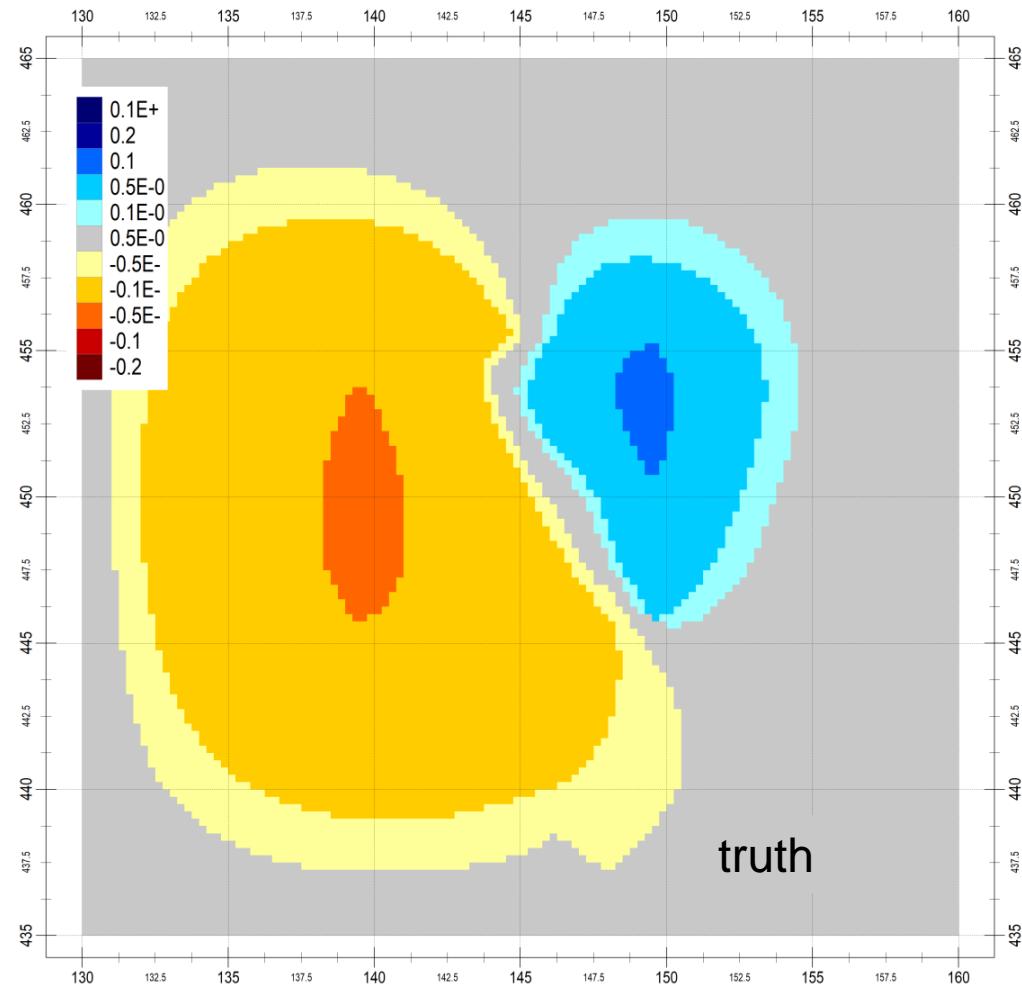
Submodel Model Optimization Method (SMOM)



SMOM – GHB-EDGE $\text{cond} = q/dh$



SMOM – GHB-EDGE



SMOM – RESULTS

full overlap

half overlap

quarter overlap

GHB-BOUNDARY

	TRUTH	10000	5000	2500	1000	0	10000	5000	2500	1000	0	
iterates	0	4	4	4	4	4						
doelf.	199.687	162.829	160.951	162.757	169.232	181.881	228.278	0.98847	0.99956	1.03932	1.11701	1.40195
kd_z128	1	0.4894	0.538	0.56	0.595	0.759	1.467	1.09929	1.14425	1.21576	1.55086	2.99752
kd_z129	1	3.65549	3.025	4.289	8.427	10	0.171	0.82752	1.1733	2.3053	2.73561	0.04678
kd_z130	1	9.7792	10	5.769	9.707	10	1	1.02258	0.58993	0.99262	1.02258	0.10226
kd_z146	1	0.67596	0.678	0.686	0.671	0.68	0.648	1.00302	1.01486	0.99267	1.00598	0.95864
kd_z147	1	3.46153	3.625	3.688	3.542	2.638	0.591	1.04723	1.06543	1.02325	0.76209	0.17073
kd_z148	1	4.66887	7.408	6.155	2.238	1.13	0.457	1.58668	1.31831	0.47934	0.24203	0.09788
kd_z166	1	0.33774	0.343	0.33	0.331	0.338	0.29	1.01557	0.97708	0.98004	1.00077	0.85865
kd_z167	1	4.75738	5.612	5.17	4.337	6.615	7.479	1.17964	1.08673	0.91164	1.39047	1.57208
kd_z168	1	0.58062	0.582	0.591	0.398	0.535	0.26	1.00238	1.01788	0.68548	0.92143	0.4478
							sum	1.0871	1.04308	1.06512	1.18131	0.80582
iterates	0	4	4	4	4	4	4					

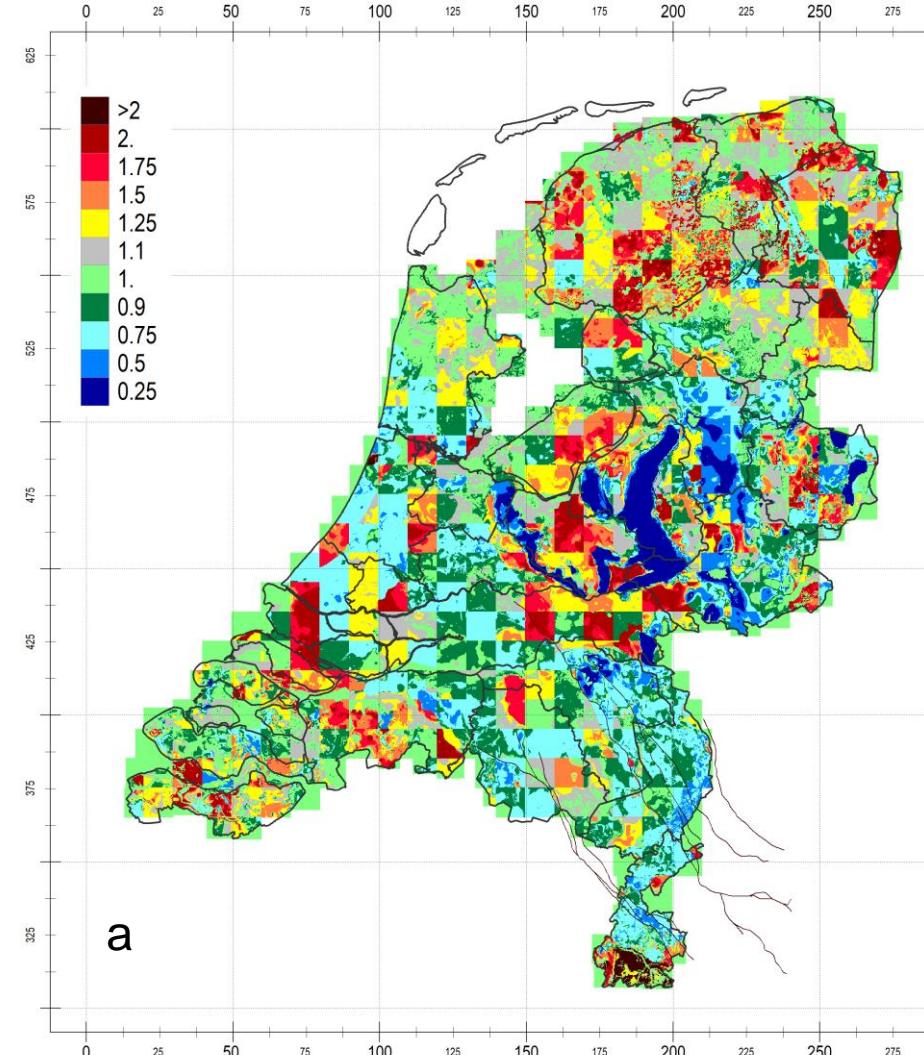
Q-BOUNDARY

	TRUTH	10000	5000	2500	1000	0	10000	5000	2500	1000	0	
iterates	0	4	4	4	4	4						
doelf.	199.687	162.829	164.933	164.939	178.618	175.524	183.853	1.01292	1.01296	1.09697	1.07797	1.12912
kd_z128	1	0.4894	0.629	0.823	0.61	0.492	0.456	1.28523	1.68163	1.24641	1.0053	0.93174
kd_z129	1	3.65549	4.524	0.165	2.224	1.585	0.923	1.23759	0.04514	0.6084	0.43359	0.2525
kd_z130	1	9.7792	10	10	0.922	2.169	1	1.02258	1.02258	0.09428	0.2218	0.10226
kd_z146	1	0.67596	0.672	0.683	0.692	0.675	0.646	0.99414	1.01042	1.02373	0.99858	0.95568
kd_z147	1	3.46153	3.631	3.48	2.122	2.517	1.942	1.04896	1.00534	0.61302	0.72714	0.56102
kd_z148	1	4.66887	9.067	10	1.038	2.545	1.128	1.94201	2.14184	0.22232	0.5451	0.2416
kd_z166	1	0.33774	0.413	0.382	0.338	0.325	0.422	1.22283	1.13105	1.00077	0.96228	1.24948
kd_z167	1	4.75738	7.103	4.686	4.828	5.667	4.131	1.49305	0.985	1.01484	1.1912	0.86833
kd_z168	1	0.58062	0.177	1.673	0.229	0.182	0.1	0.30485	2.88142	0.39441	0.31346	0.17223
							sum	1.17236	1.32271	0.69091	0.71094	0.59276
iterates	0	4	4	4	4	4	4					

fractional errors

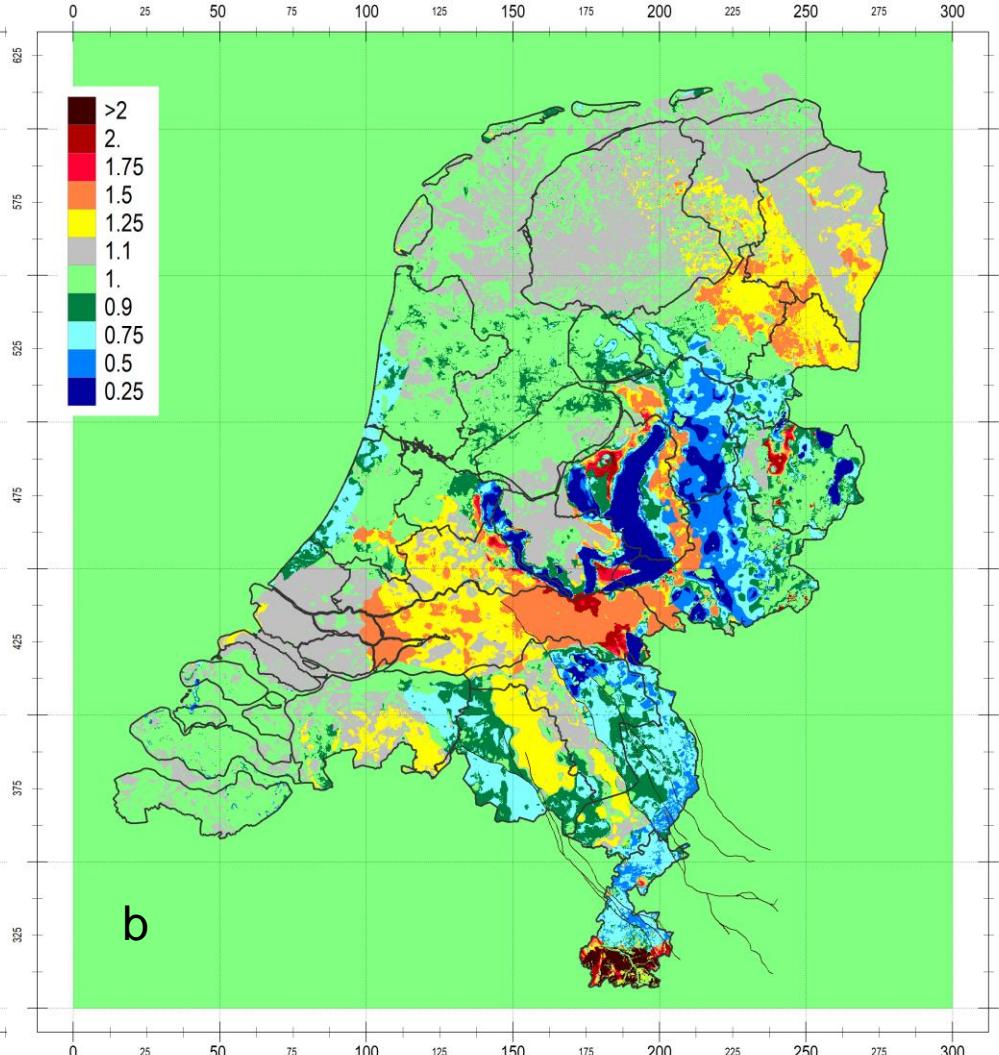
RESULTS

kd_2 submodel



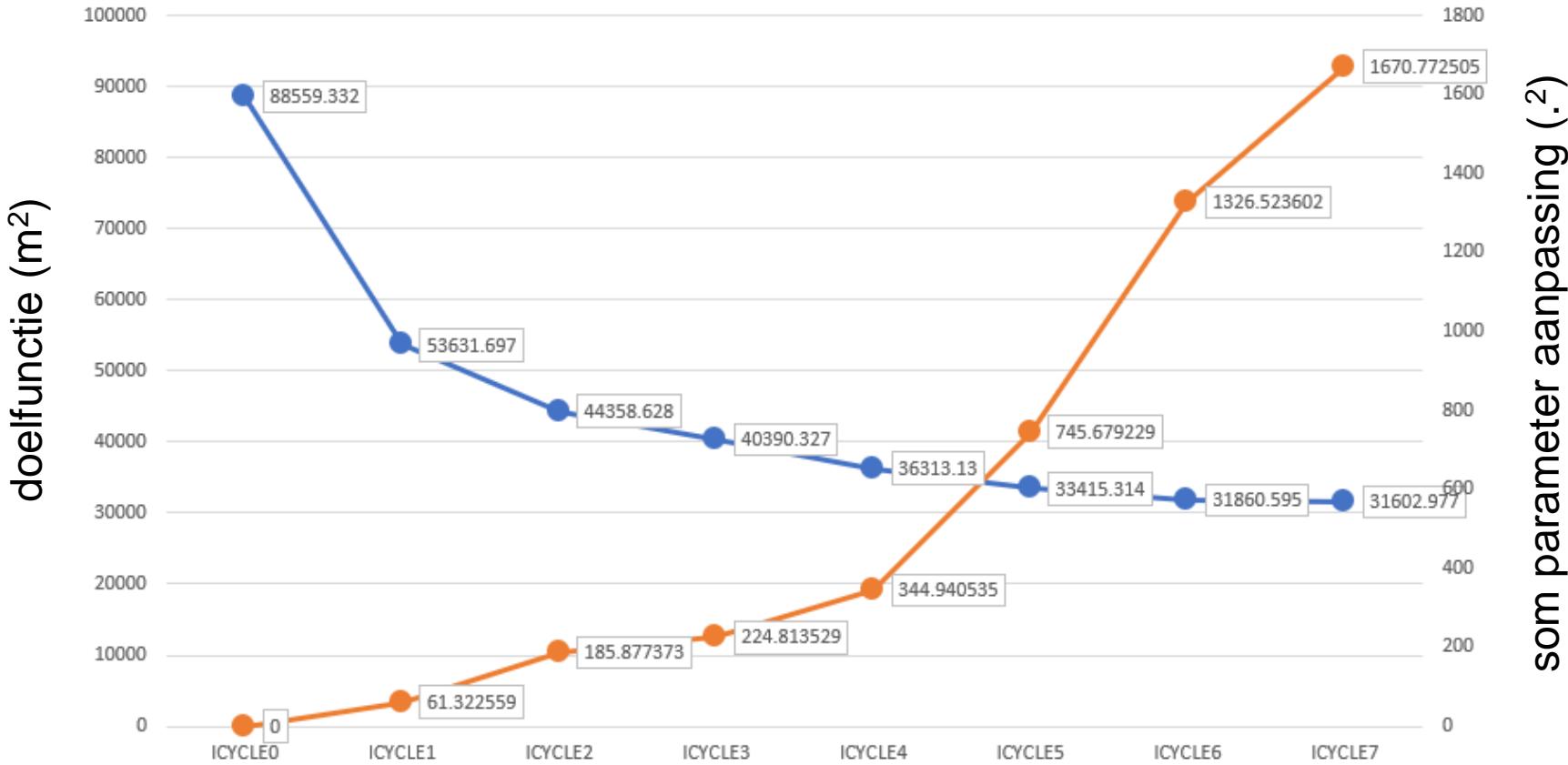
a

kd_2 globalmodel



b

RESULTS



Synthetic Example

FUNCTION=RUNFILE

```
PRJFILE_IN=D:\MODELS\TEST.PRJ
NAMFILE_OUT=D:\MODELS\TEST.NAM
IPESTP=1
NCPU=6
ISOLVE=1
MODFLOW=D:\IMODFLOW.EXE
```

NCYCLE=10
SEPMODELS=D:\MODELS\SBM.IDF
BNDSUBMODEL=3

iMOD. User Manual

EXAMINE= (optional) Use this keyword to examine on a particular location what entries exists in the model. The information is listed in the file EXAMINE.TXT in the OUT-PUT_FOLDER. EXAMINE can have the following values:

- EXAMINE>0: enter the X- and Y-coordinate of the location, e.g. EXAMINE=1200025,0,750000,0 to denote an X-coordinate of 1200025.0 and an Y-coordinate of 750000.0. For that location the correct row- and column number will be computed and all model entry is listed in the output file.
- EXAMINE<0: enter the row- and column number of the model cell to be listed in the output file

NCYCLE= (optional) Specify the number of outer cycles to be used to update boundary conditions for "separate"-modelling.

OUTER-UPDATE= (optional) Specify OUTERUPDATE=1 to update parameters found from the separate SEPMODELS using Levenberg-Marquardt rather than aggregate the individual parameters (OUTERUPDATE=0). The parameter SVD_EIGV is used to truncate the eigenvalues from the combined Jacobian matrices which are read from the file LOG_PEST_JACOBIAN.TXT.

SYNPARAM= (optional) Use this keyword to specify how parameters are adjusted.

1. Specify SYNCPARAM=1 to synchronize parameters by averaging sensitivities and residuals among the separate models and create a global adjustment factor for all separate models. This is the default value.
2. Specify SYNCPARAM=2 to aggregate all individual sensitivities in a global jacobian matrix to compute adjustment factor for each separate model individual;
3. Specify SYNCPARAM=3 to aggregate all individual sensitivities in a global jacobian matrix and compute a single adjustment factor for each separate model based on explained variance.

NMODELS= (optional) Specify the number of separate models, e.g. NMODELS=4 to specify four separate models which are all equally distributed of the total modeled area. This area can be given with the keywords WINDOW or NETWORKID, and in case these are all absent, the area is derived from the first active IDF from the PRJ file.

BUFMODELS= (optional) Specify the size of the overlapping buffer in between separate models, e.g. BUFMODELS=250.0 to denote a buffer size of 250 meter. BUFMODELS is read only in case NMODFLOW>0 and by default BUFMODELS=0.0 meter.

INTPARAM= (optional) Specify INTPARAM=0 to specify that parameters are modified within the entire model area, which includes the area of the buffer (in case BUFMODELS>0 meter). In case INTPARAM=1 the parameters are optimized for the model area outside of the buffer which avoid optimizing parameters for adjacent separate models that might overlap. The default setting is INTPARAM=0.

SEPMODELS= (optional) If NMODELS is absent, specify the IDF files that configures the layout of separate models, e.g. SEPMODELS=D:\DATA\SEPMODELS.IDF. The number need to be unique but can be discontinuous, so values of 1,3,10,54 are valid and denote finally four separate models. The NodataValue in the SEPMODELS are ignored.

SKIPMODEL= (optional) If SKIPMODEL=1, the initial global model is skipped as it is assumed that it is computed beforehand. Thereafter, existing separate models will be skipped for the first cycle to be able to perform a restart. Whether a models exists is determined by the existence of the file LOG_PEST_JACOBIAN.TXT. Initial values are read from the PRJ-files created initially and are called "SEPMODEL(i).PRJ". By default SKIPMODEL=0 and all separate models are optimized initially.

IPKS= (optional) Whenever SIM_TYPE=1 or SIM_TYPE=2 IPKS=1 to apply the PKS package instead of the PCG package. The solver is more robust than the PCG solver, so you might want to use it in case of non-convergence due to huge contrasts in conductivity using the multi-core applications. This option is applicable whenever OUT is specified, whenever RUNFILE_OUT is specified, this keyword has any effect.

NRPROC= (optional) Enter the number of processors to be used for the PKS, e.g. NRPROC=5 to specify 5 processors (PKS runs in parallel mode). By default NRPROC=1 and the PKS is used in serial mode.

PKSMERGE= (optional) Specify whether result files need to be merged after the simulation. PKSMERGE=0 does not merge the files and PKSMERGE=1 can be specified which deletes the original IDF file after merging.

XS= (optional) or **NSEP=** (optional) The method to apply the load of the partial models, these are:
- Bisection Partitioning.

LOAD= (optional) Describes the load distribution (PARTOPT=2, the LOAD-zones) or the in- and active nodes (PARTOPT=3, the active nodes).

PARTOPT= (optional) 1 or SIM_TYPE=2

RUNFILE=D:\PROGRAMS\IMODFLOW.EXE

RUNTYPE= (optional) 1 or SIM_TYPE=4 or SIM_TYPE=5

RUNNAME= (optional) The file that will be created, e.g. RUN-1005 compatible model, all arrays are listed in RUN-1 or IDEBUG=2 to export all model input to

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EXCLUDE= (optional) Specify EXCLUDE=1 (this is the default value) to exclude model area outside the specifies extent of the sepmodel as defined in SEPMODELS. These areas are treated as nodatavalues and are excluded in the modeling and optimization. In this case no overlap of the optimization per sepmode occurs. In case EXCLUDE=0, the area surrounding the specific sepmode is included and causes an influence on the optimized values.

Synthetic Example

BND for Submodels
Original Model

PEST information
after aggregation
Original Model input
Submodels

The screenshot shows a Windows desktop environment. In the center is a Notepad window titled "single_zone_sm500mtr_4msr_2param.TXT - Notepad". The content of the file is as follows:

```
SUMMARY OF SEP-MODELLING
          cycle1   cycle2   cycle3
0       0.270    0.015    0.008    0.004 ← Global Objective Function
1           0.059    0.000    0.000
2           0.008    0.000    0.000
3       0.100    0.000    0.000 ← Local Objective Functions

ICYCLE=      1
IPRM PT      PARAM  F_FINAL     MEAN      STDEV  F_MDL1   F_MDL2   F_MDL3
1 KH        AQUIFER1_KH  0.298    0.254    0.114   0.151    0.236    0.376
2 KH        AQUIFER2_KH  0.979    1.152    0.603   0.599    1.062    1.794

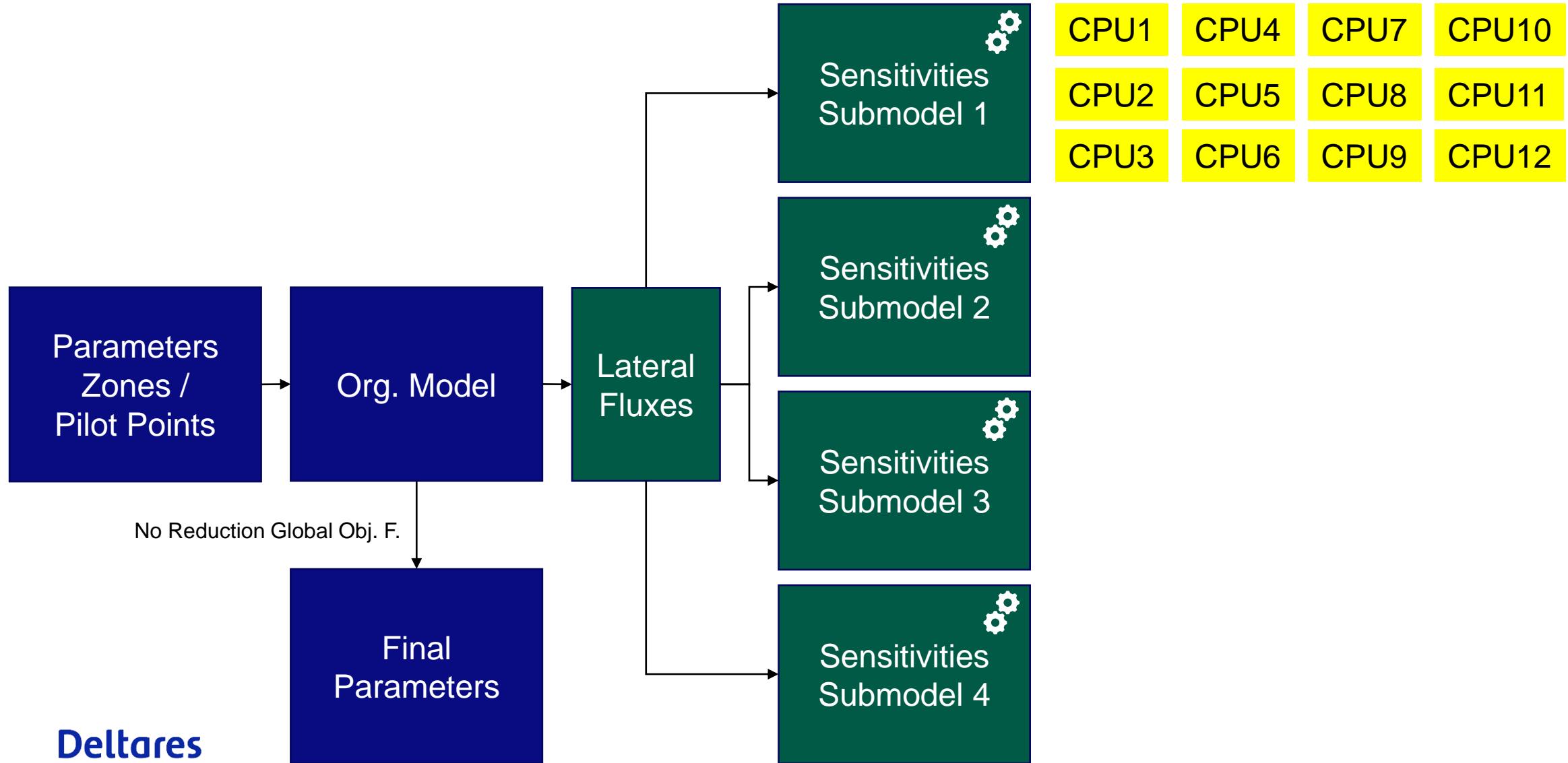
ICYCLE=      2
IPRM PT      PARAM  F_FINAL     MEAN      STDEV  F_MDL1   F_MDL2   F_MDL3
1 KH        AQUIFER1_KH  0.233    0.226    0.049   0.281    0.186    0.211
2 KH        AQUIFER2_KH  1.228    0.800    0.350   0.396    0.996    1.008

ICYCLE=      3
IPRM PT      PARAM  F_FINAL     MEAN      STDEV  F_MDL1   F_MDL2   F_MDL3
1 KH        AQUIFER1_KH  0.265    0.269    0.048   0.292    0.214    0.302
2 KH        AQUIFER2_KH  1.384    0.942    0.361   0.538    1.231    1.058
```

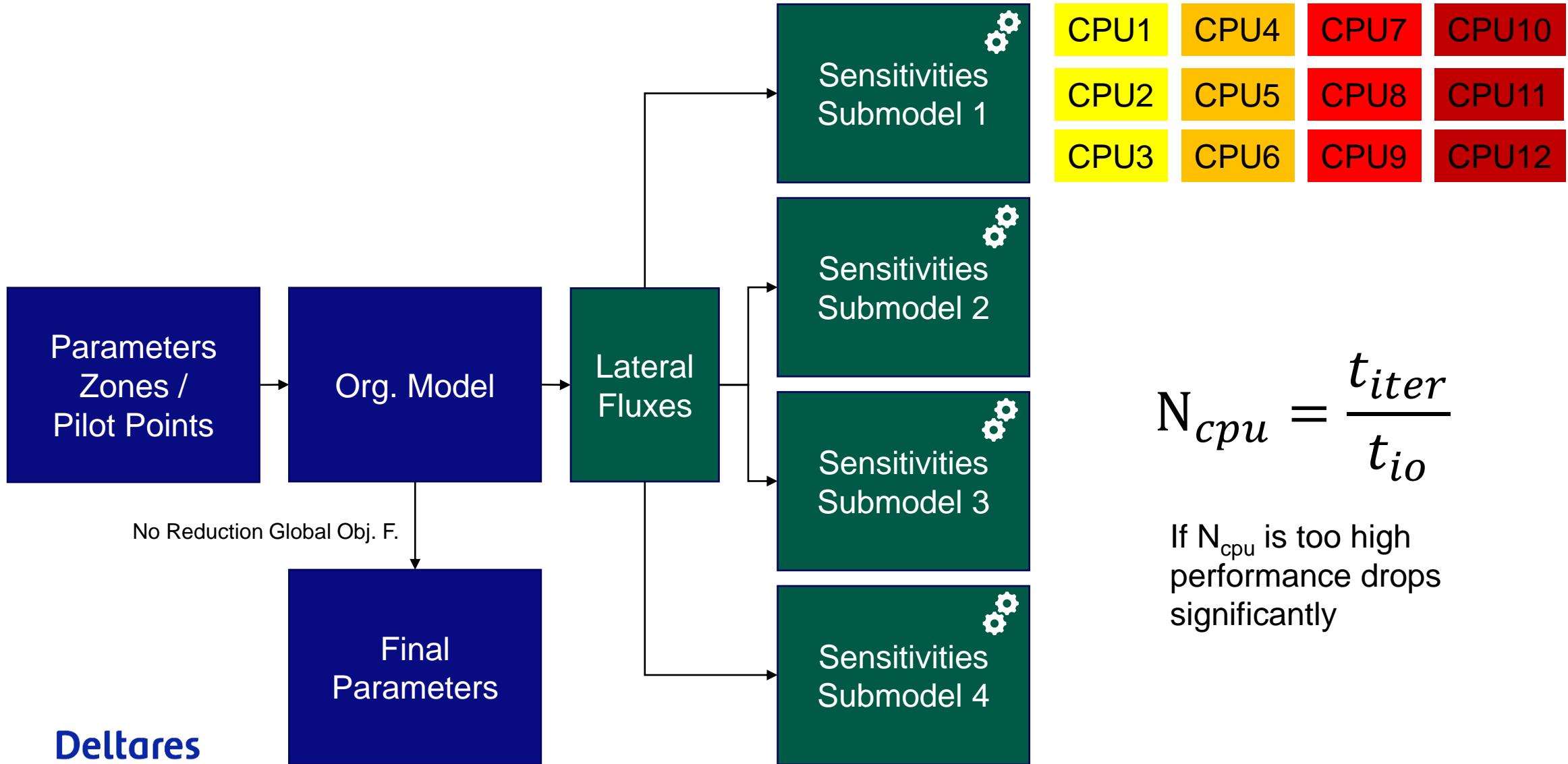
Three arrows point from the text "Global Objective Function" and "Local Objective Functions" to the numerical values in the Notepad file. To the right of the Notepad window is a Total Commander file browser showing a directory tree for "m\SEPMODEL1*.*". At the bottom right is a command prompt window with the text "Alt+ F4 Exit".

Performance

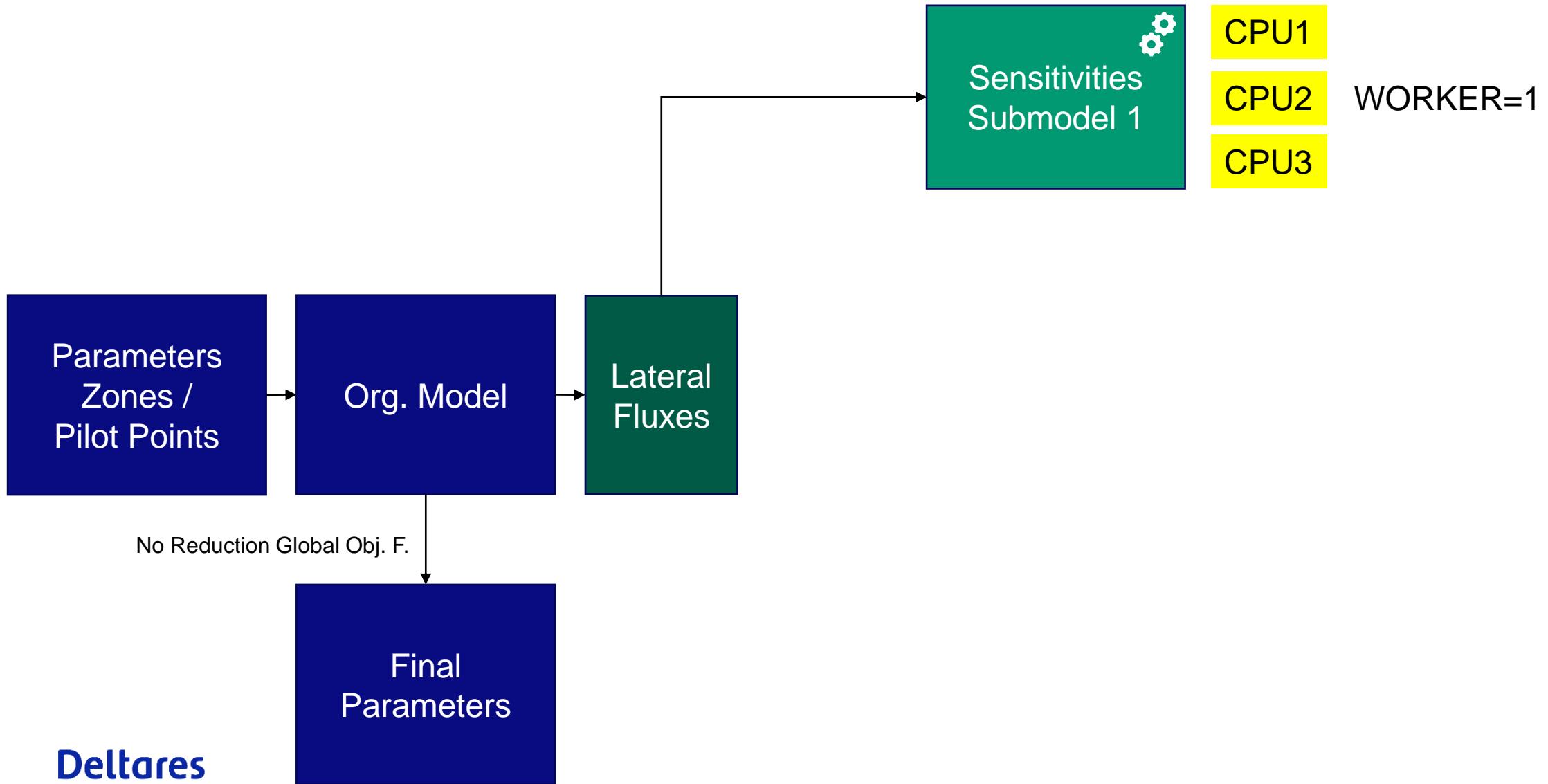
NCPU=12



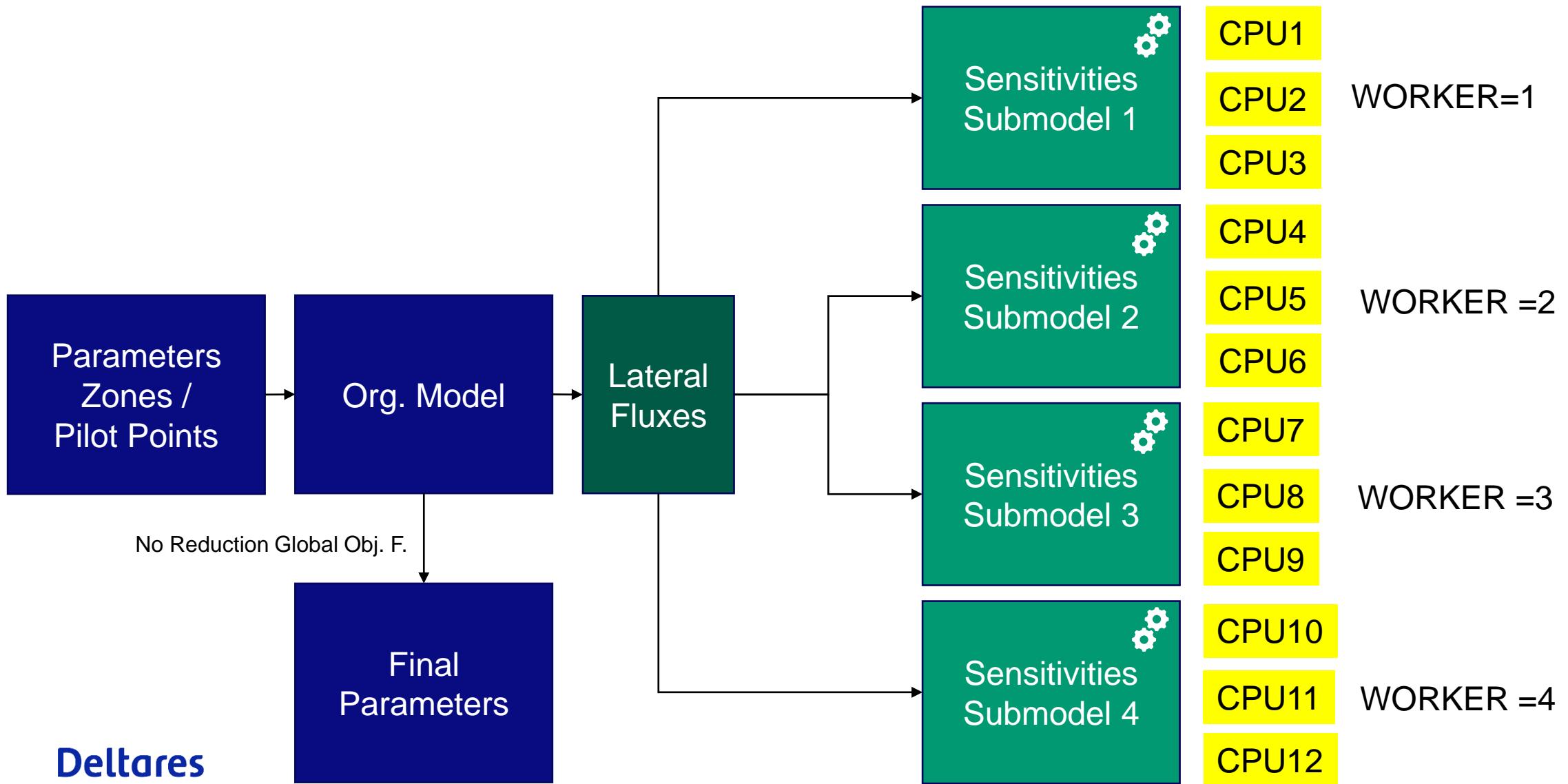
Performance



Performance



Performance



Synthetic Example

FUNCTION=RUNFILE

```
PRJFILE_IN=D:\MODELS\TEST.PRJ
NAMFILE_OUT=D:\MODELS\TEST.NAM
IPESTP=1
NCPU=6
ISOLVE=1
MODFLOW=D:\IMODFLOW.EXE
```

NCYCLE=10
SEPMODELS=D:\MODELS\SBM.IDF
BNDSUBMODEL=3
NWORKERS=3

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EXAMINE= (optional) Use this keyword to examine on a particular location what entries exists in the model. The information is listed in the file EXAMINE.TXT in the OUT-PUT_FOLDER. EXAMINE can have the following values:

- EXAMINE>0: enter the X- and Y-coordinate of the location, e.g. EXAMINE=1200025,0,750000,0 to denote an X-coordinate of 1200025.0 and an Y-coordinate of 750000.0. For that location the correct row- and column number will be computed and all model entry is listed in the output file.
- EXAMINE<0: enter the row- and column number of the model cell to be listed in the output file

NCYCLE= (optional) Specify the number of outer cycles to be used to update boundary conditions for "separate"-modelling.

OUTER-UPDATE= (optional) Specify OUTERUPDATE=1 to update parameters found from the separate SEPMODELS using Levenberg-Marquardt rather than aggregate the individual parameters (OUTERUPDATE=0). The parameter SVD_EIGV is used to truncate the eigenvalues from the combined Jacobian matrices which are read from the file LOG_PEST_JACOBIAN.TXT.

SYNCPARAM= (optional) Use this keyword to specify how parameters are adjusted.

1. Specify SYNCPARAM=1 to synchronize parameters by averaging sensitivities and residuals among the separate models and create a global adjustment factor for all separate models. This is the default value.
2. Specify SYNCPARAM=2 to aggregate all individual sensitivities in a global jacobian matrix to compute adjustment factor for each separate model individual;
3. Specify SYNCPARAM=3 to aggregate all individual sensitivities in a global jacobian matrix and compute a single adjustment factor for each separate model based on explained variance.

NMODELS= (optional) Specify the number of separate models, e.g. NMODELS=4 to specify four separate models which are all equally distributed of the total modeled area. This area can be given with the keywords WINDOW or NETWORKID, and in case these are all absent, the area is derived from the first active IDF from the PRJ file.

BUFMODELS= (optional) Specify the size of the overlapping buffer in between separate models, e.g. BUFMODELS=250.0 to denote a buffer size of 250 meter. BUFMODELS is read only in case NMODFLOW>0 and by default BUFMODELS=0.0 meter.

INTPARAM= (optional) Specify INTPARAM=0 to specify that parameters are modified within the entire model area, which includes the area of the buffer (in case BUFMODELS>0 meter). In case INTPARAM=1 the parameters are optimized for the model area outside of the buffer which avoid optimizing parameters for adjacent separate models that might overlap. The default setting is INTPARAM=0.

SEPMODELS= (optional) If NMODELS is absent, specify the IDF files that configures the layout of separate models, e.g. SEPMODELS=D:\DATA\SEPMODELS.IDF. The number need to be unique but can be discontinuous, so values of 1,3,10,54 are valid and denote finally four separate models. The NodataValue in the SEPMODELS are ignored.

SKIPMODEL= (optional) If SKIPMODEL=1, the initial global model is skipped as it is assumed that it is computed beforehand. Thereafter, existing separate models will be skipped for the first cycle to be able to perform a restart. Whether a models exists is determined by the existence of the file LOG_PEST_JACOBIAN.TXT. Initial values are read from the PRJ-files created initially and are called "SEPMODEL(i).PRJ". By default SKIPMODEL=0 and all separate models are optimized initially.

IPKS= (optional) Whenever SIM_TYPE=1 or SIM_TYPE=2 IPKS=1 to apply the PKS package instead of the PCG package. The solver is more robust than the PCG solver, so you might want to use it in case of non-convergence due to huge contrasts in conductivity using the multi-core applications. This option is applicable whenever OUT is specified, whenever RUNFILE_OUT is specified, this keyword has any effect.

NRPROC= (optional) Enter the number of processors to be used for the PKS, e.g. mode. By default NRPROC=1 and the PKS is used in serial mode.

PKSMERGE= (optional) Specify whether result files need to be merged after the simulation. PKSMERGE=0 does not merge the files and PKSMERGE=1 can be specified which deletes the original IDF file after merging.

XS= (optional) or **NSEP>0** The method to apply the load of the partial models, these are:

- Bisection Partitioning,
- Scribes the load distribution (PARTOPT=2, the LOAD-zones) or the in- and active nodes (PARTOPT=3, the - and active nodes),

PARTOPT= (optional) 1 or SIM_TYPE=2

NMODFLOW=D:\PROGRAMS\IMODFLOW.EXE

RUNFILE= (optional) SIM_TYPE=4 or SIM_TYPE=5

RUNFILE (optional) file that will be created, e.g. RUNFILE=RUN005 compatible model, all arrays are listed in RUNFILE=1 or IDEBUG=2 to export all model input to RUNFILE

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Conclusies

- Met half overlappende submodellen wordt voldoende gevoeligheid per parameter verkregen om parameters op globale schaal aan te passen. Het effect wordt dan, gezien vanuit het meetpunt, door het gehele model gebied door gegeven;
- Het noodzakelijkerwijs opschalen van het model voor de kalibratie wordt hiermee voorkomen;
- Het verspreiden van submodellen over verschillende computers vergroot de efficientie;
- Grottere spreidingslengte dwingen grotere buffers af rondom submodellen maar doordat submodellen elkaar allemaal overlappen wordt dit verholpen;
- Met 15 computers en 416 submodellen is het LHM niet-stationair gekalibreerd op basis van de GxG binnen 1-2 weken.

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