# **TROPOMI CH4 inversions**

Jacob van Peet 29<sup>th</sup> International TM5 meeting Wageningen, 21-22 November 2019



## Outline

- CH4 inversions using TROPOMI data
- Vertical interpolation
- Horizontal merging
- Initial concentration
- Model runs
- Results
- Conclusions and Outlook

### **CH4 inversions using TROPOMI data**

- CH4 measurements from TANSO on board GOSAT can be assimilated into TM5 using 4DVAR
- Using the existing code as a template, try to assimilate TROPOMI CH4 data into TM5
- 1<sup>st</sup> step: try to read and plot the data
- Two formats: "SRON R&D" and the official product, available from https://scihub.copernicus.eu/
  - The R&D format is "1D" (i.e. pixels are included sequentially), while the official format is "2D" (i.e. pixels are ordered in orbits on a lat/lon grid).



R&D

#### official

4

## **Vertical interpolation**

- old: "squeeze" model profile onto measurement grid, pressure levels are changed
- new: ignore model profile outside measurement grid, pressure levels are not changed





# **Horizontal merging**

- Merge TROPOMI data on a regular grid
  - -180 <= lon < 180 and -90 <= lat < 90
- Take the weighted average of the relevant variables.  $\checkmark$ 
  - how to interpret the "mean AK"
- Select the median observation in the gridcell
  - Aki Tsuruta & Ella Kivimäki; FMI



### Ratio between XCH4 and a-priori

#### Ratio between XCH4 and a-priori

file = S5P OFFL L2 CH4 20190314T110140 20190314T124310 07336 01 010202 20190320T125906.nc

file = S5P OFFL L2 CH4 20190314T110140 20190314T124310 07336 01 010202 20190320T125906.nc d lon, d lat = 6.0, 4.0



(xch4-xch4

\_apri)/xch4

apri

(%)

(xch4-xch4

apri)/xch4

apri

(%)

### **Initial concentration**

- Updated by default in 4DVAR. To disable:
  - set the error to 0
  - remove initial concentration from the state vector  $\checkmark$
- Use a mean mix file from the CAMS dataset and convert it to an initial concentration file
  - v18r1-c73-S1-A-model-mmix\_glb600x400\_20180101.hdf
  - provided by Arjo Segers



## **Model runs**

- With surface measurements only
  - data provided by Arjo Segers
  - c73-S1-A-years-2018-INPUT-point\_input.nc4
- With TROPOMI CH4 measurements only
  - new vertical interpolation
  - horizontal merging on 6 x 4 degrees
- 4 sources: biomass burning, rice, wetlands, and other
- Maximum of 10 iterations due to time constraints for this meeting
- Model runs from 1-1-2018 till 1-5-2019
- Analyse from 1-3-2019 till 1-3-2019



run id = 20191115-tm5\_sf-1y\_sat\_only-merge\_6x4

date = 201806, iter = iter-0001



run id = 20191115-tm5\_sf-1y\_sat\_only-merge\_6x4

date = 201806, iter = iter-0010



### CH4 emission for different sources

date = 201806; run id = 20191114-tm5\_sf-1y\_points\_only



date = 201806; i\_iter = iter-0001; run id = 20191114-tm5\_sf-1y\_points\_only



date = 201806; i\_iter = iter-0001; run id = 20191115-tm5\_sf-1y\_sat\_only-merge\_6x4



date = 201806; i\_iter = iter-0010; run id = 20191114-tm5\_sf-1y\_points\_only



date = 201806; i\_iter = iter-0010; run id = 20191115-tm5\_sf-1y\_sat\_only-merge\_6x4



### **Conclusions and outlook**

- The TM5 4DVAR model has been run succesfully on a global scale for a year:
  - with point observations
  - with TROPOMI observations
- After 10 iterations, emission updates show different patterns
- To do:
  - validate with independent measurements (TCCON, stations)
  - check model and measurement uncertainties
  - combined point and satellite assimilation
  - increase maximum number of iterations
  - increase resolution to 3x2 and 1x1 (lon x lat)