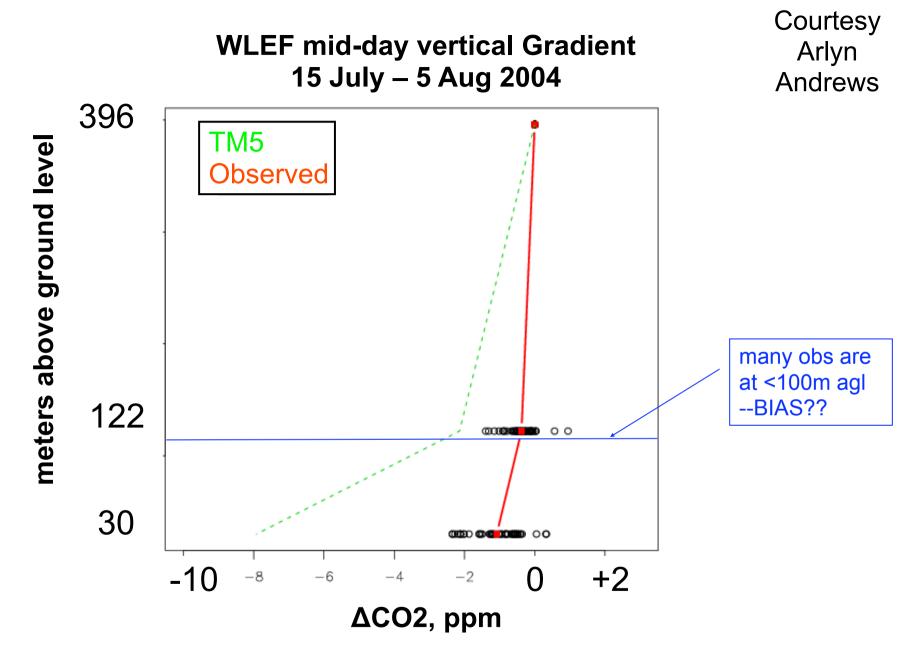
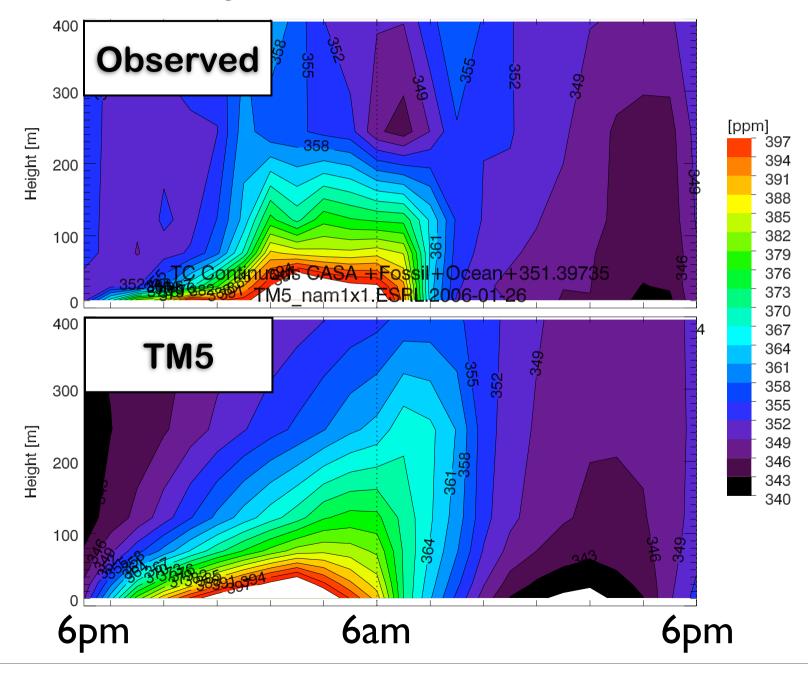
# TM5 Sampling Issue

What's wrong with the simulated surface layer?

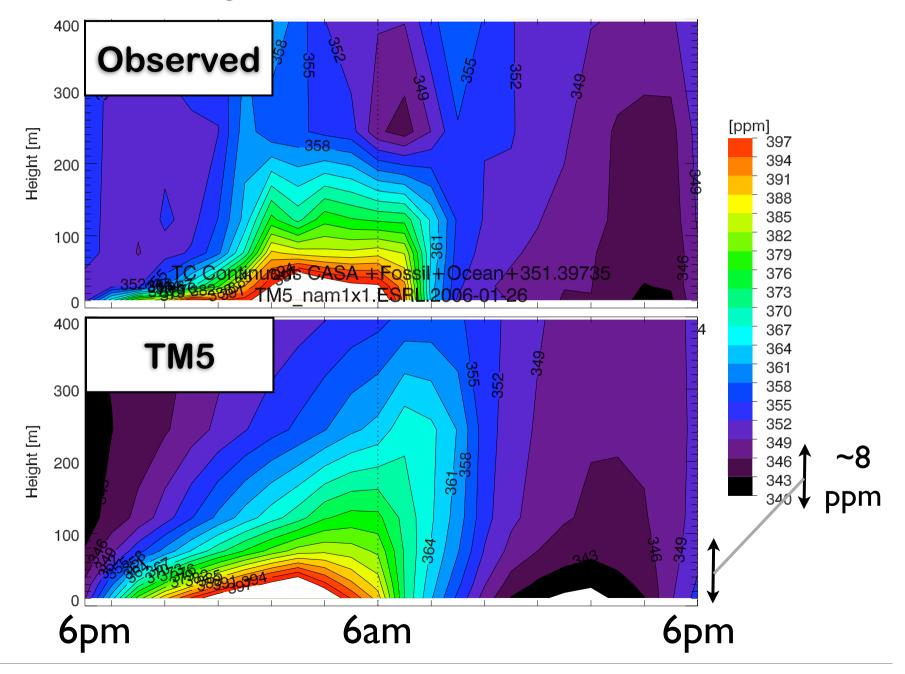


TM5 overestimates vertical gradient from 30 – 400m by factor of 4 near the surface

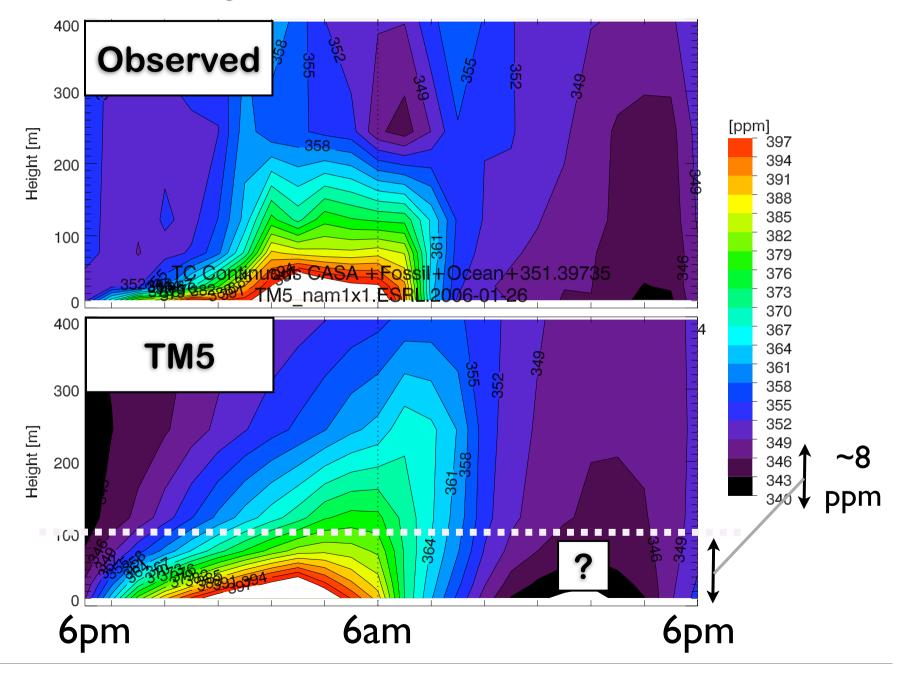
Diurnal cycle of CO<sub>2</sub>, June-10th-2002



Diurnal cycle of CO<sub>2</sub>, June-10th-2002



Diurnal cycle of CO<sub>2</sub>, June-10th-2002



## Hypotheses

A) The vertical diffusion is not fast enough in TM5 to wipe out the near-surface gradient

- B) The way in which emissions are added (including setting the slope) leaves too much CO2 near the surface and not enough at higher levels.
- C) Interpolation of the CO2 profile to the surface is done incorrectly because rzm slopes are used (linear and large)
- D) The operator splitting and the way we sample within it causes biases from including one sample directly after sources.

# Hypotheses

The vertical diffusion is not fast enough in TM5 to wipe out the near-surface gradient

- B) The way in which emissions are added (including setting the slope) leaves too much CO2 near the surface and not enough at higher levels.
- C) Interpolation of the CO2 profile to the surface is done (1) incorrectly because rzm slopes are used (linear and large)

(2)

(6)

D) The operator splitting and the way we sample within it causes biases from including one sample directly after sources.

### (B) Emissions added incorrectly?

One could distribute emissions through levels
1-3 instead of only having them affect level-1

Currently, we set the slope of the tracers in the do\_add\_2d routine to have maximum mixing ratios near the surface after emissions, even further skewing the profile (C) Interpolation during sampling introduces biases?

Results from TransCom Continuous intercomparisons demonstrated large sensitivity to sampling method

 Not only horizontal interpolation, but also vertical interpolation is important contributor

Many observations are at surface, hence extrapolation is used from few points in profile

### (D) Operator splitting and sampling

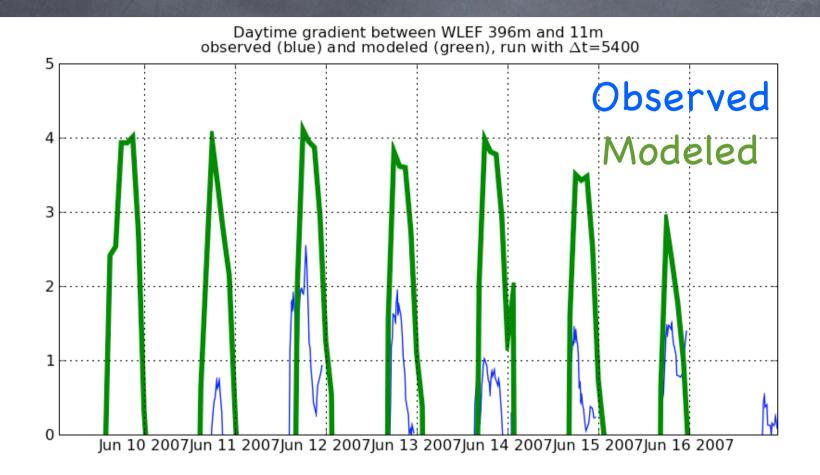
We currently sample (\*) twice per sequence:

\*XYZVSC - \*CSVZYX

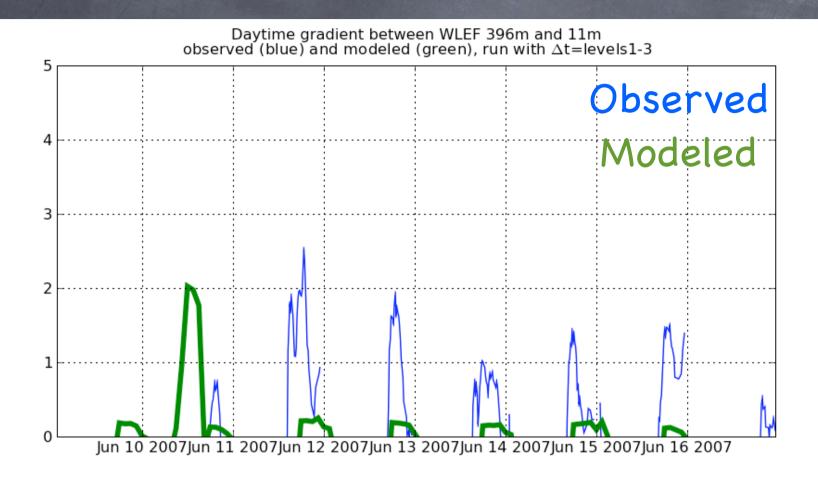
and average the result. One sample is after tracer updates by S and C, what if these tendencies are really large?

Small time step avoids numerical inaccuracies

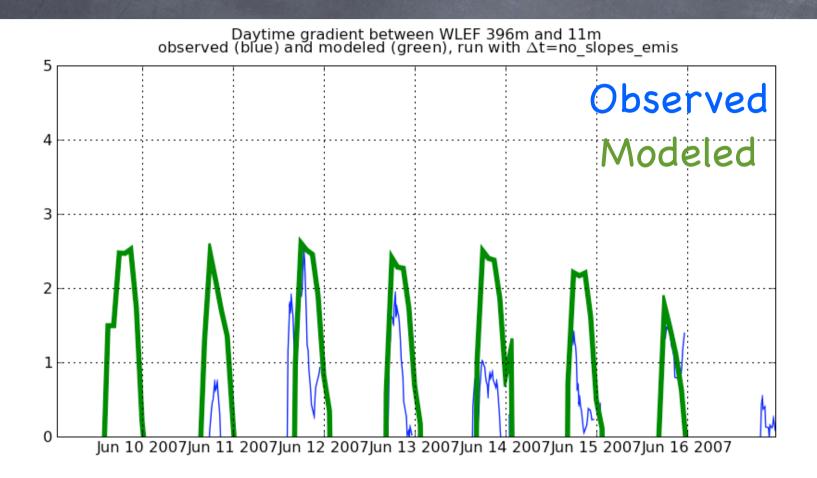
Smart sampling too?



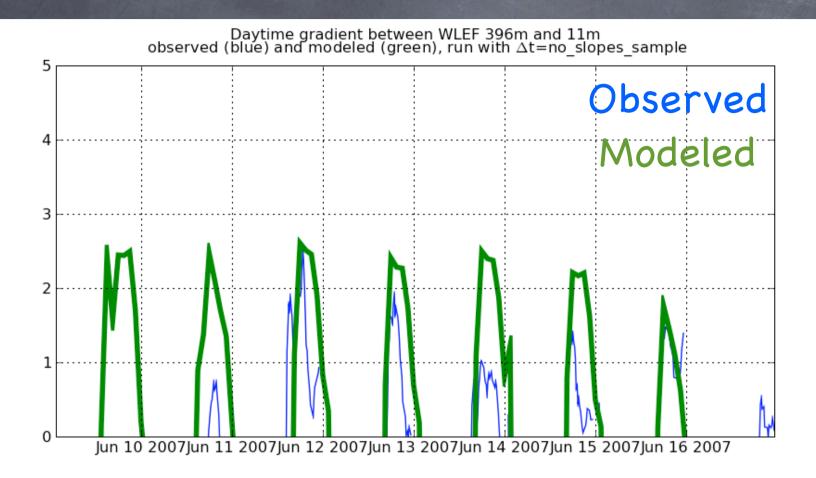
#### $\Delta t = 5400$ seconds



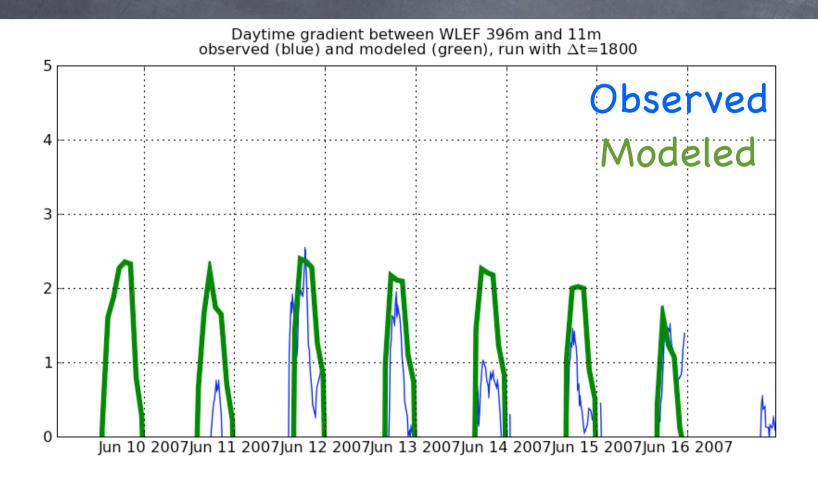
#### emissions in levels 1-3 (mass-weighted)



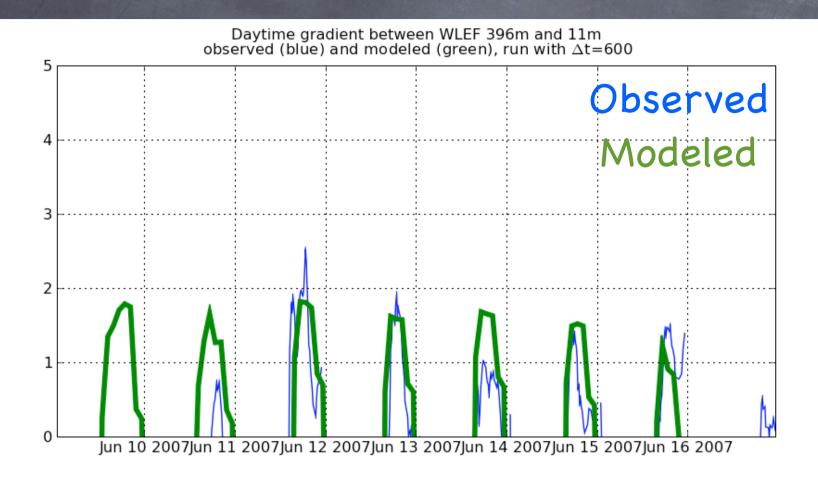
no slopes setting in emissions (do\_add\_2d)



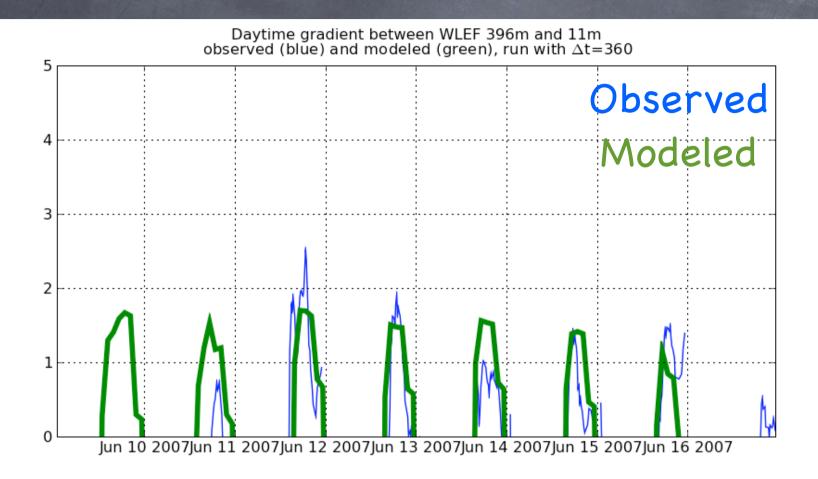
#### no vertical slopes in sampling



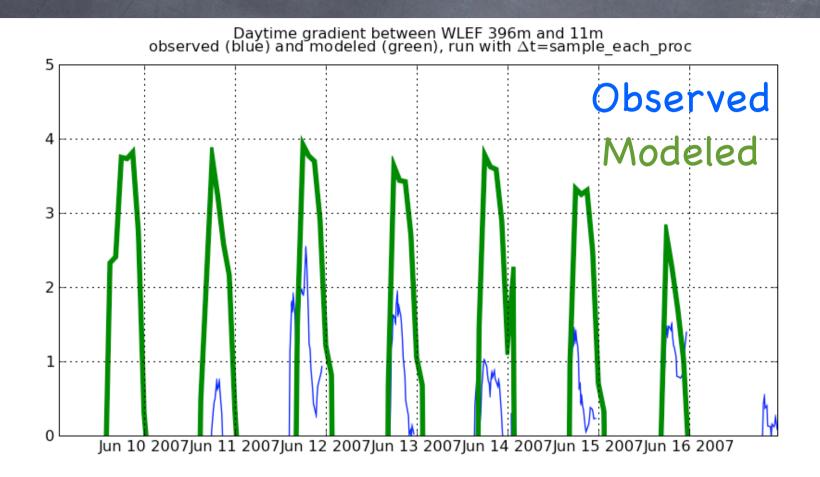
 $\Delta t = 1800$  seconds



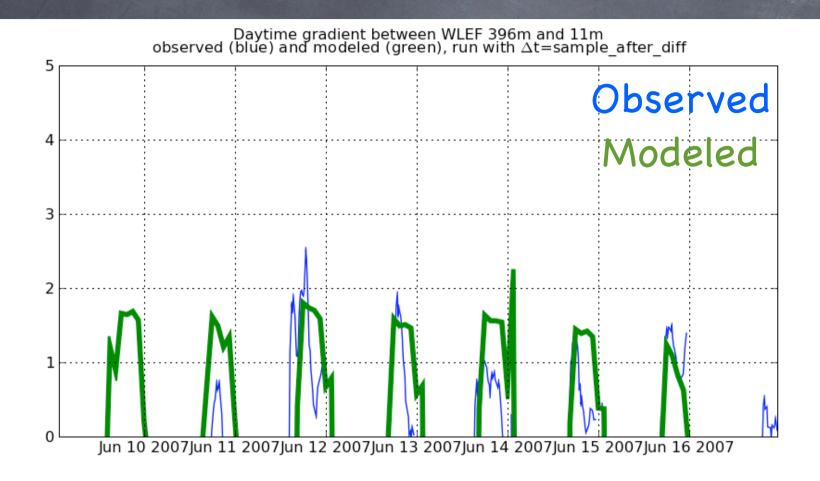
 $\Delta t = 600$  seconds



 $\Delta t = 600$  seconds

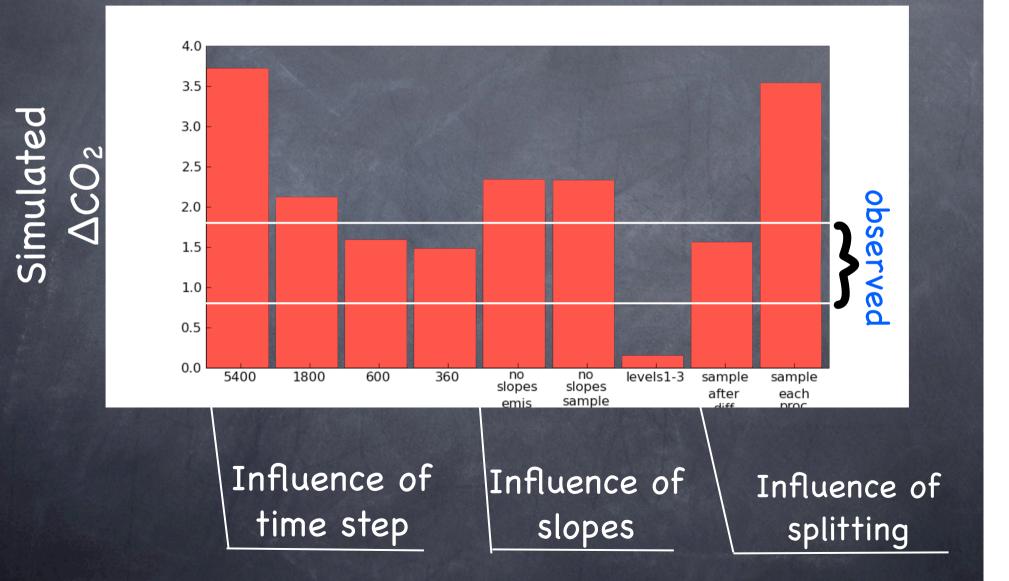


Sample after each process (X\*Y\*Z\*V\*S\*C\*)

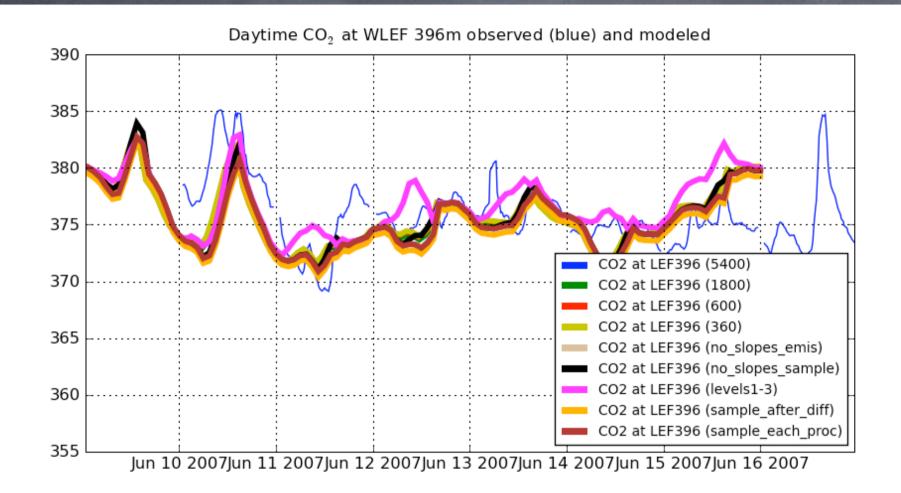


Sample after diffusion only (XYZV\*SC-CSV\*ZYX)

### Summary



## And note:



Mixing ratios at 396m are hardly affected by any of the changes attempted!

## Conclusions

- A dangerous time step dependence of the numerical results of TM5 exists, manifesting itself strongest near the surface
- Careful sampling is the best way to minimize these errors
- For now, it seems prudent to \*stop\* using vertical slopes in the emissions and sampling routines, at least near the surface
- Smarter sampling can help, but 'smart' is application specific and thus not easy to implement
- A more generic solution is under development (WUR: Maarten)
- This problem is not CO2 specific: it exists for all tracers, and especially for those with large tendencies in S or C !!!