

# SFIT4 – Retrieval parameters

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# Retrieval parameters



# Retrieval parameters

## Overview

- rt** The main parameter `rt` can be used to switch off and on the retrieval altogether.
- rt.lm** Switches on or off the Levenberg-Marquardt iteration scheme.
- rt.convergence** the iteration is considered converged when  $\text{rt.convergence} > D\_CHI = (CHI\_2\_MAX_{i-1} - CHI\_2\_MAX_i)$
- rt.max\_iteration** maximum number of iterations



# Retrieval parameters

## Overview

| ITER | FIT_RMS | GAMMA    | CHI^2_X | CHI^2_Y | CHI^2   | CHI^2_OLD | DCHI^2     |
|------|---------|----------|---------|---------|---------|-----------|------------|
| 1    | 25.5294 | 1.00E+05 | 0.000   | 886.616 |         |           |            |
| 2    | 15.7601 | 1.00E+05 | 0.000   | 332.528 | 332.528 | 886.616   | 554.088226 |
| 3    | 9.8038  | 1.00E+04 | 0.004   | 134.162 | 134.166 | 332.528   | 198.361710 |
| 4    | 5.1080  | 1.00E+03 | 0.038   | 32.608  | 32.646  | 134.166   | 101.519910 |
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| 7    | 1.8797  | 1.00E+00 | 0.281   | 2.594   | 2.875   | 3.048     | 0.172872   |
| 8    | 1.8844  | 1.00E-01 | 0.300   | 2.573   | 2.873   | 2.875     | 0.002286   |

FINAL: MEAN\_SNR= 86.6267 MEAN\_FIT\_RMS(%)= 1.88443 NVAR= 185 NFIT= 3477

| BAND | SCAN | RMSSNR (CALCULATED) | (EFFECTIVE) | (RETRIEVED) | CHI^2 |
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NEGATIVE MIXING RATIO VALUES FOUND FOR : 03668

NEGATIVE MIXING RATIO VALUES FOUND FOR : 03686



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**rt.max\_iteration** maximum number of iterations

For all retrieval parameters a priori is given and a standard deviation  $\sigma$  in the form

**rt.x.apriori** the apriori of a given value. It is actually applied in the forward calculation. Meaning it can also be used in forward calculations.

**rt.x.sigma** the entry in the  $S_A$  matrix corresponding to this parameter.



# Retrieval parameters

## Wave number scaling and shifting

`rt.wshift` wave number shift.

- ▶ Shift works on the internal grid  
**band.X.calc\_point\_space**
- ▶ This is only useful for microwindows (small) because the mismatch is a wavenumber dependent polynomial. For small wave number regions, this can be approximated by a shift.
- ▶ This is on top of `rt.wave_factor`, which is a scaling.
- ▶ The artificial grid needs to be more dense than the measured grid

`rt.dwshift` wave number shift for each retrieved gas separately, except the first retrieved one.

- ▶ all lines of each gas are shifted by the same amount (!!!)



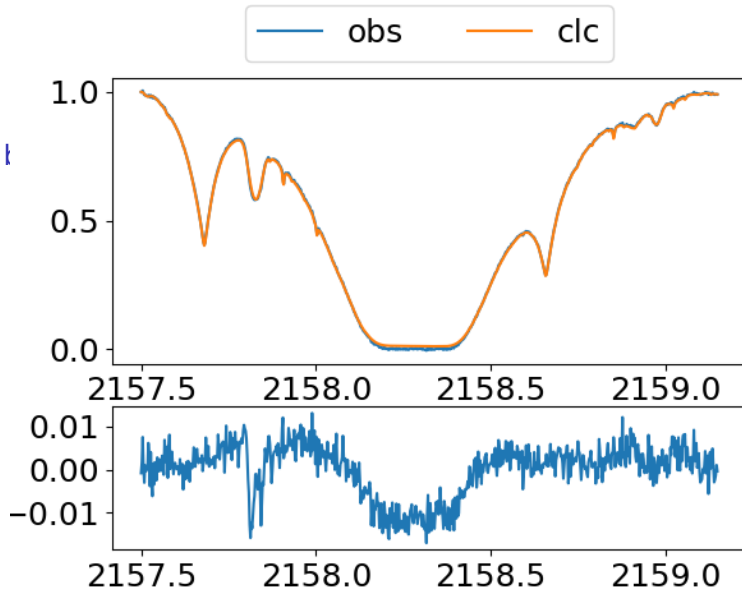
## offset in window

`band.zshift` calculates and retrieves an offset in the microwindow. Two types:

`.type = 1` retrieves the offset in this MW **ONLY ONE!!!**



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- `.type = 2` uses the offset which is retrieved in another microwindow. THIS MUST BE LATER THAN THE MW USED FOR **ZSHIFT.TYPE=1**

`band.zshift.apriori` is also an **FW parameter**



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`band.zshift.apriori` is also an **FW parameter RETRIEVAL ONLY POSSIBLE IF THERE IS AN SATURATED PART IN THE MW.**



# channeling in window

The channeling in a MW is calculated via

`band.x.beam` = 1,2 The beams with the numbers 1 and 2 are calculated

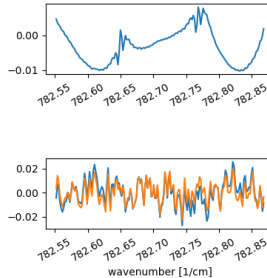
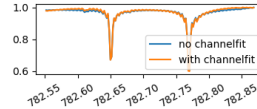
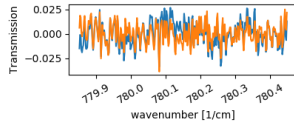
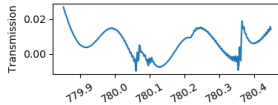
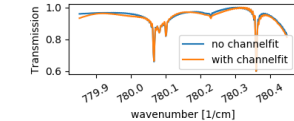
`band.x.beam.model` = IP, PS which one is better has to be checked.

`band.x.beam.1.apriori` = AMP, FREQ, PHAS, SLOPE defines the apriori values of the beam 1

`band.x.beam.1.sigma` standard deviations for all parameters, i  
sigma = 0, parameter is not retrieved,



# channeling in window



and 2 are

is to be

defines

rameters, i

## rt.slope and rt.background

rt.slope and rt.background can be used to model a sensitivity function of the instrument, caused by a filter or the wave number dependent sensitivity of the instrument itself.

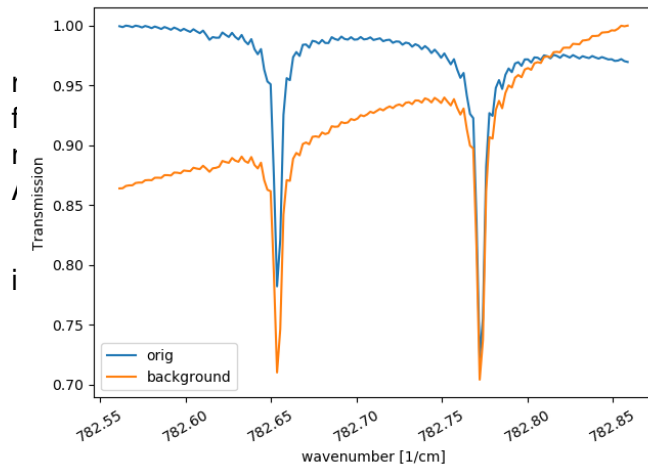
A function

$$\text{rt.slope} * \nu + \text{rt.background} * \nu^2 \quad (1)$$

is multiplied to the calculated spectrum.



# rt.slope and rt.background



sensitivity  
wave  
f.

(1)

# Retrieval parameters

## Construction of the $S_A$ matrix.

The  $S_A$  matrix is constructed from the sigma values given. How this is actually done, depends on the parameters. In principle the  $S_A$  is constructed as a diagonal matrix and inverted in the code to yield the  $S_A^{-1}$  matrix. Some caveats:

`gas.profile.x.correlation` off diagonals using the sigma values as maxima

`.type = 1` gaussian with FWHM = `.width`

`.type = 2` exponentially with FWHM = `.width`

`.type = 3` not used

`.type = 4` the  $S_A$  matrix is read in from `file.sa_matrix`

`.type = 5` the  $S_A^{-1}$  matrix is read in from `file.sa_matrix`



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`gas.profile.x.regmethod` lets you chose between OE optimization and the Thikonov-Phillips regularization with L1 constraint (smoothness constraint)

- `.type = 'OE'` optimal estimation (Rodgers, 2000)
- `.type = 'TP'` Thikonov-Phillips with smoothness constraint
- `.lambda` strength of the regularization in TP. The higher the value the less is the regularization.

The smoothness constraint is calculated from `file.stalayers` in order to adapt for non-unique altitude layering. The matrix is scaled using the `gas.profile.x.sigma` values.





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**FIT\_RMS** mean variance of the residuum

**GAMMA** the Levenberg Marquardt Parameter

**CHI\_2\_X** A measure of the deviation of the retrieved state from the A PRIORI



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**CHI\_2\_Y** A measure for the retrieval quality

$$\chi_Y^2 = \frac{(y_M - y_C)^T S_\epsilon (y_M - y_C)}{m}$$

$\chi_Y^2 = 1$  if the residuum is reduced to the noise as specified in  $S_\epsilon$



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- ▶ In the lower half, the retrieval diagnostics for the last calculation are shown for each MW.
- ▶ A warning if retrieved profiles have negative parts



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Iteration stopped if either  $DCHI < rt.convergence$  or  $ITER > rt.max\_iteration$ , what ever comes first.



# The finish

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|------|------|---------------------|-------------|-------------|-------|
| 1    | 1    | 138.44              | 138.44      | 113.72      | 1.48  |
| 2    | 1    | 146.53              | 146.53      | 143.52      | 1.04  |
| 3    | 1    | 327.50              | 327.50      | 185.55      | 3.12  |
| 4    | 1    | 319.57              | 146.69      | 81.72       | 2.40  |

NEGATIVE MIXING RATIO VALUES FOUND FOR : H2O

NEGATIVE MIXING RATIO VALUES FOUND FOR : C2H4

|            |     |            |     |            |     |            |     |            |     |             |     |
|------------|-----|------------|-----|------------|-----|------------|-----|------------|-----|-------------|-----|
| 03         | : T | H2O        | : T | CO2        | : F | 03668      | : F | 03686      | : F | C2H4        | : F |
| 9.5734E+18 |     | 5.2139E+22 |     | 7.9696E+21 |     | 1.6077E+19 |     | 1.6077E+19 |     | 9.8413E+14  |     |
| 1.0921E+19 |     | 4.1944E+22 |     | 1.4369E+22 |     | 1.6255E+19 |     | 1.5224E+19 |     | -9.1080E+14 |     |

Iter/Mx:05/15 %RMS=1.125 FitPrm=117 CVRG:T DIVW:F DOFS=4.755 SNR= 162. CHI\_2\_Y= 2.2875

RDRV: DONE. ELAPSED TIME = 200.01916800000001

