# *WinDOAS 2.1* Software User Manual

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WinDOAS is distributed without any restrictions. In case of publications making use of WinDOAS, please mention the following authors in the acknowledgements.

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#### **Table of Contents**

1	11	NTRODUCTION	7
	1.1	Purpose of the Document	7
	1.2	References	7
	1.3	Overview of the Document	8
2	11	NSTALLING WINDOAS	9
	2.1	System and Hardware Requirements	9
	2.2	Installation	9
	2.3	Associate wds File Extension with WinDOAS	9
3	D	ESCRIPTION OF THE USER INTERFACE	11
	3.1	The User Interface Components	11
	3.2	The Four Main Windows in WinDOAS	12
		The Environment Workspace Window	12
		The <b>Projects</b> Window	12
		The <b>Data</b> Window	12
		The <b>Graphs</b> Window	12
	3.3	The Main Menu Bar	12
	3.4	The Toolbar	13
	3.5	The Status Bar	14
	3.6	The Shortcut Menus	14
4	D	ESCRIPTION OF THE ALGORITHMS	15
	4.1	Main WinDOAS features	15
	4.2	Block-diagram structure of the programme	16
	4.3	Slant column fitting	21
		The Beer-Lambert law	21
		DOAS fitting	21
		Intensity fitting	22
	4.4	Fit Parameters	22
		Absorption cross sections	22
		Polynomial	24
		Shift and Stretch	24
		Offset Correction	24
		Ring Effect	25
		Undersampling Correction	25
		Standard Deviations of The Fitted Parameters	25
		AMF	26
		Graphical Display Of Spectral Fit Results	26
	4.5	Wavelength calibration and slit function characterisation	27
	4.6	Ott-line tools	27
		Convolution/Filtering 1 ool	2/
			30
5	~	Undersampung 1001	21
Ð	E 1		
	5.1	Creating a Project	33
		Greating & INEW Project	22
		Inserving Tues	94
		Drowstng Spectra	94
		Tunuung Grupps	)) 36
	52	Saving Town work	90
	5.4	Configuration of a Floject for Dro Reffleval	30
		Congregating 22 1 Topen 1 of Spectra 22nacysis	)0 27
		Analysis Solut For Br() Retrienal	)/ 38
		2 Innejsis Scorp I of DO Ixertevan	20
		Degining Symood Projects Protectios	Jo 28
		1 rojens 1 rojenses	90 42
		2 inaujsis w indow 1 roporaces Runnino Analysis	+2 44
			17

6	Т	HE WORKSPACE ENVIRONMENT TREE	49
	6.1	Creating a New Item in the Workspace Environment Tree	49
	6.2	Observation Sites Properties	50
	6.3	User-Defined Symbols Properties	50
	6.4	Handling Items in the WorkSpace Environment Tree	50
7	Т	HE PROJECTS TREE	51
	7.1	Description of the Projects Tree	
	7.2	Directories and Files Management	
		Spectra Files	
		Ûser-defined Folders	
		Directories structures	
		Handling Spectra	53
	7.3	Handling Items In The Projects Tree	53
	7.4	Project Properties	
		The Spectra Tab Page	
		The Analysis Tab Page	
		The Filtering Tab Page	
		The Calibration Tab Page	
		The Undersampling Tab Page	
		The Instrumental Tab Page	
		The Slit Function Tab Page	
		The ASCII Results Tab Page	
		The NASA-AMES Results Tab Page	
	7.5	Analysis Windows Properties	
	7.6	Configuration of the Fitting Parameters	67
		The Molecules Pages	
		The Continuous Functions Page	
		The Predefined Parameters Pages	
		The Shift and Stretch Pages	
		The Gaps Page	
		The Outputs Pages	
	7.7	Configuration of the Wavelength Calibration procedure	
		The Molecules pages	
		The Continuous Functions Page	
		The Predefined Parameters Page	
		The Shift and Stretch Pages	
		The Gaps Page	
		The Outputs Page	
8	т	HE WINDOAS TOOLS	81
-	8.1	Supported line shapes	
	8.2	About Files Formats	
	8.3	The Convolution/Filtering Tool	
		The General Tab Page	
		The Slit Function Tab Page	
		The Filtering Tab Page	
	8.4	The Ring Tool	
	8.5	The Undersampling Tool	86
9	Δ	NNEXES	
•	91	Files format	87
	<i></i>	Calibration	87
		Cross Sections	
		Solar Spectrum	
		Reference spectrum	
		AMF	
		Slit functions	88 88
	92	Troubleshooting	80
	/ . 4	- 10 a c c c c c c c c c c c c c c c c c c	

## Acronyms and abbreviations

AMF	Air Mass Factor
ASCII	American Standard Code for Information Interchange
BrO	Bromine monoxide
Cic	Colour Index
DOAS	Differential Optical Absorption Spectroscopy
ERS	European Remote Sensing Satellite
FT	Fourier Transform
FWHM	Full Width at Half Maximum
GOME	Global Ozone Monitoring Experiment
IASB-BIRA	Belgian Institute for Space Aeronomy
MDI	Multiple Document Interface
nm	nanometers
NDSC	Network for the Detection of Stratospheric Change
NLLS	Non Linear Least Squares
OHP	Observatoire de Haute Provence, France
RAM	Random Access Memory
RMS	Root Mean Square
RRS	Rotational Raman Scattering
SCD	Slant Column Densities
SFP	Slit Function Parameter
SVD	Singular Value Decomposition
SZA	Solar Zenith Angle
VCD	Vertical Column Densities

#### 1 Introduction

#### 1.1 Purpose of the Document

WinDOAS is a software developed at IASB/BIRA for analysis of UV-visible spectra by the method of Differential Optical Absorption Spectroscopy (DOAS) [12]. It takes the advantage of the experiment acquired by IASB/BIRA from early nineties in DOAS measurements and analysis. Several European groups of the DOAS community contribute also to the improvement and the evolution of the software by their helpful suggestions, applications and inter-comparison exercise.

A separate version, GWinDOAS, has been developed for GOME, (Global Ozone Monitoring Experiment), a nadir-viewing UV-visible spectrometer aboard on the European satellite ERS-2. GOME measures the solar radiation back scattered from the atmosphere in the spectral range 230-800 nm with a high spectral resolution allowing the observation of minor trace gases implied in the ozone depletion, like BrO.

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#### 1.3 Overview of the Document

Chapter 2 : Installing WinDOAS	This chapter covers system and hardware requirements and guide you in the installation of the software.	
Chapter 3 : Description of The User Interface	<ul> <li>This chapter covers the main components of the WinDOAS user interface :</li> <li>the four child windows;</li> <li>the menu bar;</li> <li>the toolbar;</li> <li>the status bar;</li> <li>the right-click shortcut menus.</li> </ul>	
Chapter 4 : Description of The Algorithms	This chapter summarises the main features of WinDOAS and describes the structure of the programme and the nature of the coded algorithms.	
Chapter 5 : Quickstart	This chapter describes how to create a project, insert files and display spectra. Using a predefined configuration provided with the package (set up for BrO retrieval), it guides you step by step towards complete configuration of a project.	
Chapter 6 : The Workspace Environment Tree	This chapter describes how to manipulate and configure <b>Observation Sites</b> and <b>User-defined symbols</b> items in the <b>Workspace Environment tree</b> .	
Chapter 7 : The Projects Tree	<ul> <li>This chapter describes the organisation, the handling and the configuration of the different kinds of items in the <b>Projects</b> tree. It covers :</li> <li>&gt; the management of files and directories structures in the <b>Projects</b> tree;</li> <li>&gt; the configuration of projects;</li> <li>&gt; the parameterisation of the analysis;</li> <li>&gt; the parameterisation of the wavelength calibration.</li> </ul>	
Chapter 8 : The WinDOAS Tools	<ul> <li>This chapter covers the main features supported by WinDOAS about convolution and describes the three off-line tools included in the software :</li> <li>&gt; the Convolution/Filtering tool;</li> <li>&gt; the Ring tool for creating Ring effect cross sections;</li> <li>&gt; the Undersampling tool for creating undersampling cross sections.</li> </ul>	

Chapter 9: Annexes Description of the format of input files and troubleshooting.

### 2 Installing WinDOAS

#### 2.1 System and Hardware Requirements

WinDOAS runs under Windows 95/NT. A Pentium computer with at least 64 megabytes of RAM is recommended.

#### 2.2 Installation

The program with the "Software User Manual" (this document) and on-line help is usually distributed by E-mail. To install the package, just copy the sent files in a dedicated directory.

The example application described in section 5.2 is available on request.

#### 2.3 Associate wds File Extension with WinDOAS

The configuration of applications developed with WinDOAS is saved in ASCII files. The default extension for WinDOAS configuration files is *wds*. You can open an existing configuration file using the options sequence **Files-Open** from the menu bar or the icon **Open** from the toolbar. The same result will be obtained easier by double clicking on the name of any file with extension *wds*. For that, apply the following instructions :

- in My Computer or Windows Explorer window, apply the command View-Options from the menu bar;
- select the **File Types** tab page;
- click **New Type** pushbutton;
- enter WinDOAS as Description of type;
- enter \*.wds as Associated extension;
- click New pushbutton to open a New Action popup window;
- enter WinDOAS as Action;
- use Browse pushbutton to search for WinDOAS program in your path tree;
- click **OK**;
- click **Change Icon** pushbutton and then **Browse** pushbutton to search for **WinDOAS** program;
- select WinDOAS icon;
- Click ok.

### 3 Description of The User Interface

#### 3.1 The User Interface Components



The WinDOAS user interface is a Multiple Document Interface (MDI) including the following components :

➢ four main child windows;

1. the Workspace Environment window;

- 2. the Projects window;
- 3. the **Data** window;
- 4. the Graphs window.
- ➤ a menu bar;
- $\triangleright$  a toolbar;
- $\triangleright$  a status bar;
- right-click shortcut menus.

These components are further detailed in the following sections.

### 3.2 The Four Main Windows in WinDOAS

The Environment Workspace Window	In this window, the user should list all relevant symbols that will be used in the configuration of his projects and analysis windows. The <b>Observation Sites</b> coordinates are used to (re-)calculate zenith angles. For balloons and satellite measurements, they allow operating a selection on spectra to analyse, according to specified geolocation coordinates (the sites).
The <b>Projects</b> Window	WinDOAS is based on the notion of projects. A project can be defined as a set of files sharing the same configuration of analysis.
	WinDOAS allows defining several projects in a session, giving users the possibility to handle several analysis configurations. A project includes the definition of spectral windows and the list of files to analyse with this configuration.
The <b>Data</b> Window	This window displays the available information on the current spectrum and analysis results.
The <b>Graphs</b> Window	Spectra, fitting results and in general, any kind of graphical information, are displayed in the Graphs window.

### 3.3 The Main Menu Bar

Application Control	Files	usual Windows optic	ns for applications control :
		New	starts a new WinDOAS application;
		Open	opens an existing WinDOAS application;
		Save	saves the current WinDOAS application;
		Save as	renames the current WinDOAS application;
		Print	prints the content of the active <b>Graphs</b> or <b>Data</b> window;
		Print Setup	selects and set up the printer;
		Exit	exits WinDOAS
	Edit	for the moment, thi giving direct access to	s option only includes the <b>Goto</b> sub-option of any record in the current open spectra file;
	Projects	global operations on the <b>Projects</b> tree :	
		New Remove All Run Analysis Run Calibration	creates a new project in the <b>Projects</b> tree removes all projects from the <b>Projects</b> tree; applies analysis on all files of all projects; applies wavelength calibration on all files of all projects;
	Tools	gives access to WinD	OAS tools;

Windows Organization	View	calls back the selected child window;	
	Windows	usual Windows options for organising child windows in the current application or activating the selected child window;	
Online Help	Help	gives access to online help.	

### 3.4 The Toolbar



The WinDOAS toolbar proposes several shortcuts that allow easy access to the following options :

Application Control	New	starts a new WinDOAS application;
	Open	opens an existing WinDOAS application;
	Save	saves the current WinDOAS application;
Printing	Print	prints the content of the active Graphs or Data window;
Display	Zoom out	graph <b>zoom in</b> is performed by clicking and dragging the mouse over the wanted screen area; this shortcut recovers the original size of the active graph;
Spectra Browsing	Goto	direct access to a given record in the current spectra file;
	First spectrum	goes back to the first spectrum in the current file;
	Previous spectrum	goes back to the previous spectrum in the current file; this button speeds up the spectra browsing when held down;
	Next spectrum	goes to the next spectrum in the current file; this button also speeds up the spectra browsing when held down;
	Last spectrum	goes to the last spectrum in the current file;
	Continue	continues the paused spectra browsing or analysis;
	Pause	in non-stop spectra browsing or analysis, marks a pause;
	Stop	stops spectra browsing or analysis.

#### 3.5 The Status Bar

The status bar is divided in three sections :

online help	a short help text is displayed as a menu bar command or a shortcut menu option is pointed out with the cursor;
graphs coordinates	the current coordinates (in user units) are displayed as the cursor is moved on the graph;
time section	the current time is displayed with an accuracy of 1 minute.

#### 3.6 The Shortcut Menus

Menu commands and toolbar buttons are completed in right-click shortcut menus. The options proposed in the shortcut menus depend on the calling object : the **Data** window, a graph or any item in the **Workspace Environment** or **Projects** trees.

## 4 Description of The Algorithms

This chapter summarises the main features of WinDOAS and describes the structure of the programme and the nature of the coded algorithms.

### 4.1 Main WinDOAS features

Analysis	۶	DOAS/intensity fitting modes;
	۶	shift/stretch fully configurable for any spectral item (cross section or spectrum);
	۶	possibility to filter spectra and cross sections before analysis (supported filters include Kaiser, gaussian, boxcar, Savitsky-Golay);
	≻	possibility to define gaps within fitting intervals (e.g. to eliminate bad pixels);
	≻	possibility to fit an instrumental offset;
	$\triangleright$	possibility to work either in pixel or in wavelength (nm) units;
	4	possibility to define several configurations of spectral windows under a project;
Calibration And Slit Function Characterization	$\checkmark$	wavelength calibration and instrumental slit function characterisation using a non-linear least-squares (NLLS) fitting approach where measured intensities are fitted to a high resolution solar spectrum degraded to the resolution of the instrument. The fitting method (DOAS or intensity fitting) can be chosen different from the method used in the analysis;
	≻	possibility to correct for atmospheric absorption and Ring effect;
	۶	supports different analytical line shapes, as described in section 4.6;
Cross Sections Handling	٨	possibility to calculate differential absorption cross sections (by orthogonalisation or high-pass filtering);
	۶	possibility to correct cross sections using wavelength dependent AMF;
	۶	possibility to fix the column density of any selected species;
	$\blacktriangleright$	possibility to convolute cross sections in real-time using a user-defined slit function or the information on calibration and slit function provided by the wavelength calibration procedure;
	$\blacktriangleright$	possibility to handle a difference of resolution between measured and control spectra;
Undersampling	pos froi	sibility to build undersampling cross sections using the information derived m the NLLS wavelength calibration;

Output The output is fully configurable. Analysis results are saved in tabbed columns ASCII files, easily loadable in most spreadsheet programmes.

#### Tools Convolution/Filtering tool:

- standard and  $I_0$ -corrected convolutions are supported;
- possibility to create an effective slit function taking into account the (finite) resolution of the source spectrum (using a FT deconvolution method);

**Ring** tool : calculates Ring effect cross sections (Rotational Raman Scattering approach);

Undersampling tool : generates undersampling cross sections;

#### 4.2 Block-diagram structure of the programme

The overall structure of the programme is schematically described in the following figures. Figure 1 describes the general organization of the processor. The structures of the wavelength calibration and slant column fitting modules are given in Figures 2 and 3, while the coupled linear/non-linear least squares fitting algorithm is sketched in Figure 4.



Figure 4-1. Overall structure of the spectral analysis programme.







Figure 4-3. Structure of the slant column-fitting module.



Figure 4-4. Structure of the coupled linear/non-linear least squares algorithm used in the slant column fitting module and in the wavelength calibration module.

#### 4.3 Slant column fitting

The Beer-Lambert law describes the extinction of the solar radiation in an absorbing atmosphere :

$$I(\lambda) = I_0(\lambda) . exp(-\Sigma \sigma_i c_i)$$
 (eq. 1)

where :

- $I(\lambda)$  is the solar spectrum after absorption;
- $I_0(\lambda)$  is the extraterrestrial solar spectrum;
- $\sigma_{i}$  are the relevant cross sections of the absorbing species, with wavelength and temperature dependent structures;
- $c_i$  are the unknown species column densities.

The logarithm of the ratio of the control spectrum  $(I_0(\lambda))$  and the measured spectrum  $(I(\lambda))$  is denoted optical density (or optical thickness):

$$log(I_0(\lambda)) - log(I(\lambda)) = \Sigma \sigma_i c_i$$
 (eq. 2)

Two fitting analysis methods are implemented (DOAS and intensity fitting) in WinDOAS. They differ in the expression of the fitting function (use of the equation 1 or the equation 2.

- **DOAS fitting** The DOAS approach is a direct application of the equation 2. High frequency spectral structures of the various absorbing species are used to resolve the corresponding contributions to the total optical density. This is obtained using a least-square procedure where the slant column densities (SCD) of the various species are the fitted parameters. Large band contributions to the atmospheric attenuation (Rayleigh and Mie scattering) are accounted for by a low order polynomial function. Simply stated, the DOAS technique is a linear problem. This linearity is unfortunately broken down by the need to account for additional effects, namely:
  - small wavelength shifts between I and  $I_0$  spectra must be corrected using appropriate shift and stretch parameters;
  - possible instrumental and/or atmospheric straylight or residual dark current signal require the introduction of an offset parameter.

In addition to shift and offset, Ring and undersampling effects have to be treated. See section 4.4 to have an overview of all parameters to fit.

Let us consider the modified equation :

$$Log(I(\lambda) - offset(\lambda)) = log(I_0(\lambda)) - \Sigma \sigma_i c_i - P + U$$
 (eq. 3)

with *P*, the polynomial and *U*, the undersampling cross section (Ring effect cross section is simply included as an additional pseudo-absorber).

In DOAS fitting mode, WinDOAS minimises residuals of equation 3 using a Marquardt-Levenberg non-linear least-squares (NLLS) algorithm. The method implements a gradient-expansion algorithm [2], which is based on the iterative combination of a steepest-descent method (suitable for approaching the minimum from far away) and a linearisation of the fitting function.

Linear parameters are determined by a Singular Value Decomposition (SVD) method [11] embedded in the NLLS algorithm.

Intensity fitting WinDOAS also supports the so-called intensity fitting (or direct fitting) method where measured intensities are directly fitted instead of their logarithms. This corresponds to the following equation :

$$I(\lambda) - offset(\lambda) = I_0(\lambda) . exp(-\Sigma \sigma_i c_i - P) + U$$
 (eq. 4)

Two variants have been implemented.

- Full Marquardt In this method, the Beer-Lambert equation is processed without any further linearisation; all parameters are treated as non-linear parameters, with the advantage that (some of) the derivatives of the fitting function with respect to parameters can be calculated analytically. The efficiency of the method depends on the number and type of parameters to fit. Best efficiency is usually obtained when a large number of terms are included inside the exponential.
- Marquardt+SVD This method involves a decomposition of equation 4 in its linear and non-linear parts. In order to maximise the number of linear terms, the equation fitted actually slightly differs from equation 4 (the polynomial term is set out of the exponential) :

 $I(\lambda)$ -offset( $\lambda$ )= $I_0(\lambda)$ . P.exp( $-\Sigma \sigma_i c_i$ )+U

Column densities are still fitted non-linearly but polynomial, offset and undersampling are calculated using SVD. This method is to be preferred over the **Full Marquardt** method when the configuration is such that most parameters are linear (e.g. in the wavelength calibration procedure).

#### 4.4 Fit Parameters

Absorption cross sections The selection of the spectral analysis window determines which absorbers have to be included in the fitting procedure. Several cross sections of a same absorber can be fitted together (for example to account for a temperature dependency of the cross sections). In this case, WinDOAS provides an option by which two or several cross-sections can be mutually orthogonalised (using the same Gram-Schmitt's algorithm that can be also used to generate differential absorption cross sections).

> In addition to this and upon user request, several other manipulations can be applied to absorption cross sections before calculation of the slant column densities. These are described below.

Shift Absorption cross sections are interpolated on the final grid of the control spectrum which may be determined by the programme itself (see section 4.5). Moreover shift and stretch parameters can be fitted in order to obtain a best match of the absorption structures. In automatic reference selection mode (a spectrum is selected in the current file as  $I_0$ , absorption cross sections are aligned with this spectrum based on the determination of the shift between  $I_0$ and a "reference" spectrum (the spectrum with the correct calibration). Interpolation/Convolution In the usual case, cross sections are pre-convoluted and interpolated on an appropriate wavelength grid prior to the analysis. However the programme also authorises the direct use of high-resolution cross sections, which can be convoluted in real-time with a predefined slit function or with the slit function determined by the wavelength calibration process. Standard and  $I_0$  convolution types are supported (see Offline tools described in section 4.6). Differential cross sections The aim of calculating differential absorption cross sections is to separate narrow spectral features from unstructured absorption not useful in the DOAS method. Differential cross sections can be obtained either by orthogonalisation with respect to an orthogonal base formed with the component vectors of the polynomial term or by high pass filtering. In the latter case the same high pass filter is applied to spectra and absorption cross-sections. High-pass filtering is presently only supported in optical density fitting mode (DOAS mode). Low-pass filters can be applied to both spectra and absorption cross sections. A Low-pass filtering large choice of filters is proposed (see section 4.6). Wavelength-dependent Absorption cross-sections can be replaced by geometrically corrected crosssections that take into account the wavelength dependency of the AMF. The AMF correction is based on the following equation:  $log(I_0) - log(I) = \sigma(\lambda).(AMF(\lambda,SZA) - AMF_0(\lambda,SZA_0)).C(SZA)/C(SZA_0)$ where : SZA and SZA0 are respectively the SZA of the measured spectrum I, and the reference spectrum  $I_0$  (if applicable);  $AMF(\lambda,SZA)$  and  $AMF_0(\lambda,SZA_0)$ the wavelength dependent AMF are calculated in SZA and  $SZA_0$ :  $AMF(\lambda,SZA)=a(SZA)+b(SZA).\lambda+$ ...+n(SZA). $\lambda^n$ +A. $\sigma_{hr}$ and  $\sigma_{hr}$  in the formula above, is a highresolution cross section: C(SZA) and  $C(SZA_0)$ are photochemical coefficients calculated respectively in SZA and  $SZA_0$ 

**Polynomial** A polynomial function is required to account for the smooth part of the atmospheric attenuation (Rayleigh and Mie scattering). In WinDOAS, the order of the fitted polynomial can be set up to degree 5.

Differential cross sections can be obtained either by orthogonalisation to an orthogonal base formed with the component vectors of the polynomial (using Gram-Schmitt's algorithm) or by high-pass filtering (see above).

Shift and Stretch Shift and stretch parameters allow correcting for possible misalignment between the various spectral items involved in the data evaluation (i.e. measured and reference spectra as well as absorption cross-sections).

Shift and stretch parameters may be fitted or simply applied to any kind of spectral item, according to the following equation :

$$\Delta = a + b (\lambda - \lambda_0) + c (\lambda - \lambda_0)^2$$

where :

- $\lambda$  the wavelength calibration;
- $\lambda_0$  the centre wavelength of the current spectral window;
- *a* the fitted shift;
- *b* the fitted stretch (first order) ;
- *c* the fitted stretch (second order).

Offset Correction An ideal spectrometer in an ideal atmosphere would measure the part of the sunlight that has been elastically scattered by air molecules and particles in the zenith direction. In a real experiment however a number of possible additional sources of signal may add up to the ideal Rayleigh/Mie contribution leading to "offset" the measured intensity by a certain amount. In addition to the Ring effect, which is to a first approximation a natural source of offset, instrumental sources of offset also need to be considered like stray light in the spectrometer and dark current of the detector. This is the purpose of the offset parameter which we defined as follows:

$$I(\lambda)$$
-offset $(\lambda)$ . $\overline{I} = I_0(\lambda)exp(-\tau(\lambda))$ 

with :

offset(
$$\lambda$$
) =  $a + b.(\lambda - \lambda_0) + c.(\lambda - \lambda_0)^2$ 

where  $\lambda_0$  is the centre wavelength of the spectral analysis window; *a*, *b* and *c* are the fitted parameters and  $\overline{I}$  the mean intensity in the current spectral analysis window. Due to the normalisation by  $\overline{I}$ , the offset values can be easily interpreted relatively to the absolute intensity of the spectrum (percent offset).

**Ring Effect** The so-called Ring effect arises in the atmosphere due to inelastic scattering processes (mainly Rotational Raman Scattering (RRS) by molecular  $O_2$  and  $N_2$ ). Roughly speaking, it manifests itself by a broadening of the solar and atmospheric spectral features present in measured spectra. This broadening typically reduces the depth of thin solar and atmospheric absorption features by several percents. Hence, it has a strong impact on spectroscopic measurements using the DOAS method and requires appropriate correction to be implemented in retrieval algorithms. This is especially true for minor absorbers like *BrO* or *OClO*, for which weak absorption features can be completely masked by Ring structures.

In DOAS, the Ring effect is usually accounted for as an absorber. Ring cross sections can be obtained from different sources (for example Vountas et al. [13], Chance et al. [5]).

WinDOAS implements a tool for the calculation of Ring cross section using the method described by Chance et al. [5]. The approach consists in a simple convolution of a solar source spectrum using calculated RRS cross sections. Refer to section 4.6 for a brief description of the way this method is implemented.

#### **Undersampling Correction** The undersampling is a well-known problem of GOME (see Ref. [4]). It arises from the poor sampling ratio of the GOME instrument (2 to 3 pixels/FWHM of the resolution of the spectrometer) which results in a lost of spectral information when interpolating earthshine spectra during the DOAS fitting process.

To some extent, the problem can be corrected by using an ad-hoc cross section obtained by simulating the effect based on a high-resolution solar reference [4]. Undersampling cross sections can be pre-calculated using the WinDOAS undersampling tool (see section 4.6) or they can be calculated in real time, just after the wavelength calibration procedure using the corrected grid and the determined slit function.

#### Standard Deviations of The Fitted Parameters

Standard deviations are calculated directly from the SVD algorithm for linear parameters (see ref. [11]) and from the Marquardt-Levenberg algorithm for nonlinear parameters (see ref. [2]). They are weighted by the  $\chi^2$  previously divided by the square root of the number of degrees of freedom N:

$$N = \frac{Window\_width}{Filter width} - nFit$$

where :

Window widthis the width of the current spectral window;Filter widthis the total effective width of the filter (low pass and high pass);nFitis the total number of fitted parameters (linear and non-linear).

AMF The conversion of slant column densities (SCD) to vertical columns (VCD) is obtained after division by an appropriate geometrical factor (or air mass factor AMF) :

#### VCD = SCD/AMF

For the moment, AMF are not calculated by WinDOAS but imported from ASCII files (see format in section 9.1). WinDOAS supports options for solar zenith angle dependent AMF, wavelength dependent AMF and climatological AMF.

Graphical Display Of	Let us consider the following definitions:
Spectral Fit Results	$Tc = I(\lambda)$ -

and

 $T = I_0(\lambda).exp(-\tau)$  with  $\tau$  the optical density.

offset( $\lambda$ );

According to the selected fitting method, formulas used to display fitting results are different. Definitions used by WinDOAS are described in the following table :

	Calculated	Measured	
		Optical density fitting	Intensity fitting
Residuals		R = log(T) - log(Tc)	$R = T - Tc \implies R' = T/Tc = 1 + R/Tc \text{ and}$ $R'' = log(1 + R/Tc) = log(T) - log(Tc)$
Cross sections	$\sigma_i.c_i$	$R+\sigma_i c_i$	$R''+\sigma_i c_i$
Polynomial	Р	R+P	R''+P (Full Marquardt) P/R' (Marquardt+SVD)
Offset	log(1-offset/I)	R+log(1-offset/I)	R"+log(1-offset/I)

#### 4.5 Wavelength calibration and slit function characterisation

The wavelength calibration procedure implemented in WinDOAS is based on the alignment of the Fraunhofer structures of the control spectrum with those of an accurately calibrated high-resolution solar reference atlas that is degraded to the resolution of the instrument as part of a fitting procedure. The reference atlas used for this purpose is usually the Kurucz et al. [9] spectrum.

The spectral interval is divided in a number of equally spaced sub-windows in which a NLLS fitting algorithm is applied to fit measured intensities to the solar spectrum. The algorithm can take into account molecular absorption and offset correction. The derived values of the shift are used to reconstruct an accurate wavelength registration.

The instrumental slit function is characterised by fitting the parameters (usually the FWHM) of a selected line shape (see the **Supported line shapes** in next section). The wavelength dependency of the slit function is determined by polynomial fitting through the individual slit function parameters (SFP) found in the different sub-intervals.

By fitting different parameters on both sides of a Voigt profile function, the program accounts for the possible asymmetry of the slit function.

#### 4.6 Off-line tools

Convolution/Filtering Tool

**Definition** The convolution of a spectrum S by an instrumental slit function F is given by the following integral :

$$f(\lambda_0) = \int S(\lambda) \cdot F(\lambda - \lambda_0) d\lambda$$

This integral is calculated by decomposition of the surface area in trapezes. The integral interval is defined by the width of the slit function (6 x FWHM). In order to speed up the convolution algorithm, analytical slit functions are precalculated on a suitable wavelength grid (6 pixels at FWHM) and then, interpolated on the grid of the spectrum.

For Gaussian and error function line shapes, a FFT algorithm is used to calculate analytically the integral in situations requesting optimisation of the processing time (e.g. within the wavelength calibration procedure).

Supported line shapes WinDOAS supports the following line shapes for convolution :

The Gaussian line<br/>shapeThe usual 'standard' expression used to approximate instrumental slit<br/>functions is the Gaussian function. The Gaussian is the exact line shape<br/>in the diffraction limit, i.e. in the case of an infinitely thin entrance slit.<br/>The normalised Gaussian is given by :

$$G(x) = \frac{1}{a \cdot \sqrt{\pi}} \cdot \exp(-\frac{x^2}{a^2})$$

where :

*x* is the distance from the line centre,

$$a = \frac{\sigma}{\sqrt{\ln 2}}$$

and  $\sigma$ =Gaussian half-width.

If the Gaussian can be used as a good approximation for the slit function when a thin entrance slit is used, then, for a large entrance slit, the slit function can be approximated by the convolution of a boxcar and a Gaussian function. The expression to use is :

$$E(x,a,\Delta) = \frac{1}{4.\Delta} \cdot \left[ erf(\frac{x+\Delta}{a}) - erf(\frac{x-\Delta}{a}) \right]$$

where  $\Delta$  is half the pixel size and erf(x) is the well-known error function usually defined in text books by :

$$\operatorname{erf}(\mathbf{x}) := \frac{2}{\sqrt{\pi}} \cdot \int_{0}^{\mathbf{x}} \exp(-t^{2}) dt$$

The 2n-Lorentz

n=1 n=4 This line shape is a generalisation of the well-known Lorentzian function (n=1). It is useful to approximate slit functions having shapes differing significantly from the Gaussian one. Low order 2n-Lorentz can mimic line shapes having large wings (typically the Lorentzian profile), or line shapes having a plat top (high order degree). The generic expression for this function is:

$$P_{2n}(x) = \frac{\sigma^{2n}}{x^{2n} + \sigma^{2n}}$$

with  $\sigma$  the half-width at half maximum, and *x* the distance from the line centre.

**TheVoigt Profile** Function

The Voigt profile function is the convolution of a Gaussian and a Lorentzian function. This function is used in a wide range of contexts, and the optimisation of its computation has received much attention. The Voigt profile is usually expressed as :

$$K(x,y) \coloneqq \frac{y}{\pi} \cdot \int_{-\infty}^{\infty} \frac{\exp(-t^2)}{(x-t)^2 + y^2} dt$$

where x is the distance from the line centre in units of Gaussian half-widths and y is the ratio of the Gaussian to the Lorentzian half-width. Numerous ways have been proposed to compute the Voigt function. A recent and rather efficient algorithm has been recently described by Kuntz (see ref. [8]).





WinDOAS accepts also any user-defined slit function provided in an ASCII file. Finally the program can account for the wavelength dependency of the slit function characterised by the wavelength calibration procedure (except for the Voigt profile function).

Correction for the Solar The  $I_0$  correction arises because of the difference in cross-sections measured in I<sub>0</sub>-Effect the laboratory with a smooth light source and those 'seen' by the atmosphere with a structured solar light source [Johnston, unpublished results]. Basically, the I<sub>0</sub> spectrum is the highly structured solar Fraunhofer spectrum and the DOAS technique attempts to remove this spectrum by forming the log ratio of two spectra measured at different solar zenith angles thus leaving the atmospheric absorptions. However, because both of the spectra forming this ratio have been filtered by the instrument function before the calculation of the ratio, complete removal of the Fraunhofer structure is not possible. This effect may be dealt with to a good approximation by correcting the laboratory-measured crosssections with the solar Fraunhofer spectrum. Such corrections must be weighted to the column amount of the absorber so a typical maximum column amount of the various absorbers is often used. The corrections to the cross-sections are made according to the following equation:

$$\sigma_{corrected}\left(\lambda\right) = -\frac{1}{scale} \times \ln\left\{\frac{synth(\lambda)}{sol(\lambda)}\right\}$$

where  $synth(\lambda)$  is a synthetic absorbing spectrum, calculated using uncorrected high-resolution cross-section, and subsequently filtered by the instrument function,  $sol(\lambda)$  is the filtered solar Fraunhofer spectrum, *scale* is the column amount of the absorber by which the correction is scaled and  $\sigma_{corrected}(\lambda)$  is the corrected cross-section.

**Deconvolution** The WinDOAS convolution tool implements a de-convolution option. In this case, the convolution is performed using an effective slit function obtained after Fourier transform manipulation of specified convolution and de-convolution functions. The effective slit function is calculated as follows:

$$S_{eff} = \mathfrak{I}^{-1} \left[ \frac{\mathfrak{I}(S_1)}{\mathfrak{I}(S_2)} \right]$$

where  $\Im$  stands for Fourier transform, and S<sub>1</sub> and S<sub>2</sub> are respectively the convolution and de-convolution slit functions. Great care is taken in the algorithm to avoid noise-corruption effects when taking the ratio of the Fourier transforms.

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**Ring Convolution** Because of a double convolution, the algorithm used to calculate Ring cross sections (see the **Ring tool** topic) can be relatively time-consuming, depending on the resolution of the solar spectrum and the grid on which the cross-section has to be calculated. A resolution of 0.05 nm for the solar spectrum is a good compromise.

A solution to speed up the creation of Ring cross sections is to pre-calculate a high resolution (e.g. 0.02 nm FWHM) Raman spectrum which can be further degraded to any intermediate resolution. This feature is currently implemented as an option of the slant column fitting processor.

- Filtering Low-pass filters implemented in WinDOAS correspond to a convolution in the pixels domain. The following filtering functions are supported:
  - Kaiser see description of the algorithm in Kaiser and Reed, 1977 [7].
    - Boxcarconvolution with a rectangle function; this filter<br/>consists in averaging the spectrum over several spectral<br/>points;Gaussianconvolution with a Gaussian function;
    - Triangular convolution with a triangle function
    - Savitsky-Golay the Savitsky-Golay filter uses a least-square linear regression fit of a polynomial of degree k over at least k+1 data points around each point in the spectrum;
  - Binomial convolution with a filter function formed with the binomial coefficients calculated using the recursive Pascal's triangle algorithm;

High pass filters are obtained by subtraction (or division) of low-pass filtered data from the original spectrum.

**Ring Tool** WinDOAS implements a tool to calculate Ring cross section according to the algorithm described by Chance and Spurr, 1997 [5]. The approach consists in a simple convolution of a solar source spectrum using calculated rotational Raman scattering ( $\sigma_{RS}$ ) cross sections. Ring cross sections are calculated from the ratio of the rotational Raman spectrum by the solar spectrum (*Raman/Solar*) according to the following first order optically thin development for Ring effect:

 $I = I_R + RR S = I_0 exp(-\tau)$   $log(I_R + RRS) = log(I_0) - \tau$   $log(I_R) + log(I + RRS/I_R) = log(I_0) - \tau$   $Ring = log(I + a.RRS/I_R) = a.RRS/I_R - b.(RRS/I_R)^2 + \dots$ 

with  $I_R = Rayleigh$  scattered intensity RRS = Rotational Raman scattered intensity

 $\Rightarrow$  Ring cross section (1<sup>st</sup> order) = *RRS*/*I<sub>R</sub>* ~ *Raman/Solar* 

The WinDOAS Ring effect tool calculates the first and the second term of the Taylor expansion. Spectra are convoluted using a user-specified slit function.

WinDOAS can also generate Ring cross sections accounting for molecular absorption (molecular Ring effect) using a Taylor expansion of the Beer-Lambert's law :

RRS ~ 
$$I_0.exp(-\tau_{O3}).\sigma_{RS}$$
  
~  $I_0(1-\tau_{O3}+...).\sigma_{RS}$   
~  $I_0.\sigma_{RS} - I_0.\tau_{O3}.\sigma_{RS}$ 

In this last expression the Ring effect has been separated in a Fraunhofer fillingin contribution ("pure" Ring effect) and a molecular filling-in contribution (molecular Ring effect). Accordingly, molecular Ring effect cross-sections are calculated from a source spectrum given by the product of the solar intensity by the molecular absorption cross-section (of  $O_3$  in this case).

**Undersampling Tool** The problem of the undersampling can be corrected by using an ad-hoc cross section obtained by simulating the effect based on a high-resolution solar reference. Oversampled and undersampled spectra are calculated as follows :

$$U(\lambda + \Delta) = over(\lambda + \Delta) - under^{(\Delta)}(\lambda)$$

where :

over $(\lambda + \Delta)$	is a high-resolution solar spectrum convoluted on its original grid and interpolated on the final grid $\lambda + \Delta$ ;
under $^{\left( \Delta  ight) }\left( \lambda  ight)$	is a high-resolution solar spectrum convoluted on grid $\lambda$ and subsequently interpolated on the final grid $\lambda + \Delta$ .

Residuals are improved by adding a "second phase" of undersampling :

$$U_2(\lambda) = over(\lambda) - under^{(-\Delta)}(\lambda)$$

The above formulas are valid in intensity fitting mode. In DOAS fitting, the optical density is expressed as a function of the logarithm of the ratio of the measured intensities. Hence the two equations above must be replaced respectively by the following ones:

$$U (\lambda + \Delta) = \log (over (\lambda + \Delta) / under^{(\Delta)} (\lambda))$$
$$U_2 (\lambda) = \log (over (\lambda) / under^{(\Delta)} (\lambda))$$

Undersampling cross sections can be pre-calculated using the WinDOAS undersampling tool or they can be calculated in real time, just after the wavelength calibration procedure using the corrected wavelength grid and the characterised slit function.

#### 5 Quickstart

When starting WinDOAS, two empty windows (**Projects** and **Workspace Environment**) appear inside the main program window. This chapter describes how to create a project, insert files and display spectra. Using a predefined configuration provided with the package (setup for BrO retrieval), it guides you step by step towards the complete configuration of a project.

All needed input files (spectra and cross sections) can be found in the directory **Data** on the distributed disk.

#### 5.1 Creating a Project

#### Creating a New Project

To create a 'new project' item in the **Projects** tree :

- 1. select **Projects-New** command from the menu bar;
- rename the created 'new project' item; for example, give it the name 'my first project';
- 3. a dialog box with **Projects properties** appears.

Figure 5-1	DDAS Analysis Program for Windows	_ 8
	DISTRATION TO AND A THE AN	
Creating a new project	Workspace Environment     S     Project     Interview Represented	
	Analysis Vindows Project properties : my first project	
	Spectra Analysis Filtering Calibration Undersampling Instrumental Sitt Function ASCII Results NASA-AMES Results	
	Format ASCII	
	Observation site Not Specified  Detector SizeFormat	
	Image: Construction	
	Calibration file VHarestua/Uvcal.clb	
	QK Qancel Help	
	DDAS Analysis Program for Windows	16:5

The **Projects properties** dialog box includes several categories of options dedicated to the configuration of a project for analysis. Example spectra files are provided in ASCII format. A complete record is defined per file line and includes the following data : the SZA, the measurement date, the fractional time and a 1024 pixels spectrum.

In the **Instrumental** page, specify the **Detector Size** (1024), check the **Line** button in the **Format** frame and check the **Zenith angle**, **Date** and **Decimal time** buttons in the **Data to read with spectra** frame. Use the **Calibration File** 

button to locate the wavelength calibration in the **Harestua** sub-directory of the **Data** folder. If no wavelength calibration file is given, pixels numbers will be used as abscissa.

Click OK.

Inserting Files To insert files under the new project :

- 1. select the predefined **'Raw Spectra'** node and right-click **'Insert files'** to open the usual **Open Files** Windows dialog box;
- 2. locate and select some spectra files (default extension is 'SPE') for example in the **Harestua\Spectra** sub-directory.

File names have been inserted under the predefined **'Raw Spectra'** node. To display the complete file path, select the file name item in the **Projects** tree and right-click the **Properties** option.

Figure 5-2	A DOAS Majors Frugain to Windows Help	
Ŭ		
Spectra files properties	More and a second a secon	
	Raw Spectra File Properties	
	Don't create files names automatically	
	Raw Spectra E:WINDOAS/Doc/SUM/Sample/Harestua/Spectra/HA_2000_058.spe	
	Cancel Help	
	Edit properties of the selected item	12:04

Don't worry about the **Don't create files names automatically** for the moment. When the format includes record names and dark currents, you can check this button to replace default WinDOAS record names and dark currents file names by yours. The format in this example doesn't use record names and dark currents.

Browsing Spectra To browse spectra :

- 1. open **Graphs** and **Data** windows with the **View** command in the menu bar; resize and organise windows at your best convenience.
- 2. select a spectra file item and right-click the command Browse-Spectra.



The first record is displayed in the **Graphs** window. The **Data** window displays available data on the current record. Use appropriate buttons in the toolbar to move in the file.

#### Handling Graphs



You can perform zooms using the mouse. If the main **Graphs** window contains several graphs (this is not the case in this example), you can click in graph frames to redraw spectra in separate windows and then perform zoom.



You can print (bitmap printing) and save, in ASCII or bitmap format, the content of a graph window. In ASCII format, spectra are saved with their wavelength calibration and, by default, a header with the description of the record. You can disable the header safe keeping by selecting the "ASCII Spectra files without header" file type.

Use the **Stop** button (red circle) in the toolbar before coming back to the project configuration.

Saving Your Work Be sure to save your work at regular intervals using the Save button in the toolbar or the Files-Save command from the menu bar. The Save As Windows dialog box appears. Changes are saved in an ASCII file. The default extension for WinDOAS configuration files is *wds*.

#### 5.2 Example : Configuration of a Project for BrO Retrieval

Configuring A Project For Spectra Analysis	The 1.	e configuration of a project requires the following steps : define in the <b>Workspace Environment</b> tree, all relevant symbols related to the cross sections files that will be used in the analysis:
	2.	call back the <b>Projects properties</b> dialog box to parameterise the wavelength calibration procedure and to setup analysis options independent of the spectral window (fitting method, unit for shifts);
	3.	create an 'Analysis Window' item in the Projects tree for each spectral window to process, and parameterise it.
	The	as stors are described through the example file Sample with

These steps are described through the example file **Sample.wds** provided with the package.
The file <b>sample.wds</b> , in the <b>WinDOAS</b> folder, contains the configuration of a project for BrO retrieval. This file should have been copied to your hard disk during the installation procedure (see section 2.2).
Before using the <b>sample.wds</b> file, you may probably change paths. The fastest way to do it, is to edit the file and check the [All paths] section :
Piece of content of the sample.wds configuration file :
[All paths]
<pre>%0=c:\windoas\data\harestua,40 %1=c:\windoas\data\harestua\spectra,7 %4=c:\windoas,2 %6=c:\windoas\data\ohp\spectra,6 %7=c:\windoas\data\ohp,14</pre>

Loading sample.wds file

Use Files-Open command from the menu bar or the equivalent button in the toolbar.



The **Projects** tree includes two projects, **Harestua** and **OHP** corresponding to UV-measurements performed by IASB/BIRA respectively at the ground-based stations of Harestua (Norway, 60°N) and OHP (France, 44°N). Two analysis windows, **BrO** and **OCIO** have been defined for the **Harestua** project; we will focus mainly on the **BrO** one.

In the **Workspace Environment** tree are defined all symbols used in the analysis configuration of both projects.

Figure 5-6

Description of the sample.wds configuration file

Calibra	ation interval : 330-390 nm;							
Spectra	ctral window : 346-359 nm;							
Main (	Cross sections :							
	Source	Shift applied (nm)	Files on the disk (Harestua project)					
O3 NO2	GOME FM, 221K and 241K, $\rm I_0$ corrected GOME FM, 221K and 293K, $\rm I_0$ corrected	0.03 0.03	<b>o3241_</b> i0.xs and <b>o3221_</b> i0.xs <b>no2cb_</b> i0gome.xs and <b>no296</b> i0.xs					
BrO O <sub>4</sub> Ring	Wahner, 228K Hermans et al. created according to ref. [5], using WinDOAS Ring tools	0.17	bro_228wah.xs o4_chris.xs ringxs					
	Calibra Spectra Main ( O <sub>3</sub> NO <sub>2</sub> BrO O <sub>4</sub> Ring	Calibration interval : 330-390 nm; Spectral window : 346-359 nm; Main Cross sections : Source O <sub>3</sub> GOME FM, 221K and 241K, I <sub>0</sub> corrected NO <sub>2</sub> GOME FM, 221K and 293K, I <sub>0</sub> corrected BrO Wahner, 228K O <sub>4</sub> Hermans et al. Ring created according to ref. [5], using WinDOAS Ring tools	Calibration interval : 330-390 nm;         Spectral window : 346-359 nm;         Main Cross sections :         Source       Shift applied (nm)         O3       GOME FM, 221K and 241K, Io corrected GOME FM, 221K and 293K, Io corrected GOME FM, 221K and 293K, Io corrected 0.03         BrO       Wahner, 228K       0.17         O4       Hermans et al.       Ring         created according to ref. [5], using WinDOAS Ring tools       Wavelenoth calibration :					

A preliminary calibration has been performed in laboratory (Uvcal.clb). It is corrected through NLLS fitting of the control spectrum (a spectrum measured in minimal conditions of absorption) w.r.t. the high-resolution solar reference spectrum by Chance and Spurr [5]. In this case, the instrument slit function line shape is best represented by an error function (integrated gaussians). A preliminary wavelength dependency of the slit function has been calculated (sfp0\_full.slf) in order to convolute cross sections to use in the definition of spectral analysis windows.

This analysis setup is mainly based on recommendations established from previous studies (e.g. ref. [1]) and improved according to the type of intstrument and measurements.

**Defining Symbols** The symbols defined in the **WorkSpace Environment** tree must match the first characters of the cross-section files (those characters are immediately followed by an underscore in the cross section file name).

To insert symbols in the **Workspace Environment** tree, select the predefined **'User defined symbol'** node and right-click the **Insert item** command or use the **Insert** key. You can specify a short description for each symbol if you want.

Projects Properties Select the Harestua project node in the Projects tree and right-click the Properties command to open the Projects properties dialog box. Options are applied independently to both spectral windows, BrO and OClO, defined in the selected project. The most important pages in the Projects properties dialog box are :

Analysis selection of the analysis method and the unit for shift and stretch parameters;

Calibration parameterisation of the calibration.

Analysis In the Analysis tab page, we have selected : Analysis Method Optical density fitting;

 Analysis Method
 Optical density fittin

 Shift and stretch units
 Pixels.

Figure 5-7	Project properties : Harestua ? 🗙
<b>Projects properties</b> The <b>Analysis</b> tab page	Instrumental Slit Function ASCII Results NASA-AMES Results Spectra Analysis Filtering Calibration Undersampling
	Analysis Method Optical density fitting
	Least-Squares fit No weighting
	Interpolation spline
	QK <u>C</u> ancel <u>H</u> elp

Wavelength calibration The wavelength calibration interval, specified by the Window limits fields, is subdivided in a number of equally spaced sub-windows. In each of these intervals, the shift and the slit function parameter(s) (SFP) are fitted using a NLLS fitting procedure where a high-resolution solar atlas spectrum, degraded to the resolution of the instrument, is adjusted to the control spectrum. The shift values determined in all sub-windows are then used to reconstruct an accurate wavelength registration. Similarly, the wavelength dependent slit function is determined by polynomial fitting through individual SFP values. The resulting information is subsequently used in the programme, e.g. to convolute cross sections or build undersampling cross-sections.

In the example, the **Calibration** page is configured as follows :

Figure 5-8	Project properties : Harestua
Projects properties The Calibration page	Instrumental Sitt Function ASCII Results NASA-AMES Results Spectra Analysis Filtering Calibration Undersampling
	Solar ref. file UharestuaWur01.ktz Analysis Method Optical density ritting
	✓ Fit SFP Error function
	Display     Polynomial degree     Window limits (nm) -       Image: Spectra Image: Shift Spectra Image: Shift Spectra Image: Shift Spectra Image: Shift Spectra Image: Spectra Image: Shift Spectra Image:
	Number of 10 sub-windows T
	<b>Solar ref. file</b> use this button to locate the high resolution spectrum;

Analysis Method the fitting method used for wavelength calibration can be

Calibration F

		different from the one selected for spectra evaluation in the <b>Analysis tab page</b> (see above);
	Fit SFP	this button is checked in order to fit the resolution of the slit function; in this case, the <b>Error function</b> line shape is selected;
	Display	check boxes in the frame to display the indicated graphs;
	Polynomial degree	the degree of the polynomial used in the final determination of the wavelength registration (shift) and the wavelength-dependent slit function (SFP);
	Window limits	the complete calibration interval;
	Number of sub-windows	the number of calibration sub-intervals;
it Parameters	The <b>Fit Parameters</b> options. The four foll-	button calls another dialog box with several pages of owing pages have been parameterised:

Molecules	specification of the absorption cross sections included in the spectral analysis;
Continuous functions	polynomial fitting - the smooth part of the spectrum;
Predefined parameters	selection of predefined fit parameters;
Shift and Stretch	definition of shift and stretch parameters

### Molecules

In these pages, we specify which absorption cross sections are to be included in the fitting procedure (absorbing correcting terms); in this case, ozone and ring.

For best results (and given the usually limited information content over individual sub-windows), it is recommended to limit the number of free parameters as follows. First try and fit the cross section term(s) freely, looking at the retrieved values in each interval. The usual situation is that spectral signatures are such that the information content largely differs from one interval to another. Look at results in the 'best windows', and fix the absorber amount to a mean value derived from these particular intervals (in our example, the  $O_3$  column density has been fixed to 2.e19 mol/cm<sup>2</sup>). Although this way of working implicitly assumes that the  $O_3$  absorption is constant over the whole calibration interval (which is not necessarily true), this is usually the best compromise.

### Figure 5-9

#### Calibration Fit Parameters Molecules Page 1/2

We take the absorption of O<sub>3</sub> and ring into account. To obtain differential cross sections, orthogonalise them to the orthogonal base defined in the **Continuous functions** page.



To insert a new cross-section in this page, right-click the Insert Cross section

#### command.

In this example, differential cross sections are obtained by orthogonalisation to an orthogonal base formed with polynomial terms defined in the Continuous functions page.

Figure E 40	Analysis windows properties : Calibration description (Harestua project)							? ×	
Figure 5-10									
	Molecules	Continuou	s functions	Predefin	ed parameters	Shift and S	Stretch Gaps	Outputs	
Calibration Fit Parameters							·		2/2
Molecules	Cross	Sections	Co Fit	Cc init	Cc Delta	Cc I0			
Page 2/2	0324	1		2e19	0.001				
8 ,	ring			-0.15	0.001				
The densities of O <sub>3</sub> and ring									
are fixed respectively to 2e19									
and -0.15 mol/cm <sup>2</sup>									
									<b>I</b>
	ОК		Cancel	1	Help				
		- 1	<u>_</u>	. L					

#### **Continuous functions**

The smooth component of the differential absorbance is fitted by a polynomial. The selection of the degree of the polynomial is made in the **Continuous** functions page.

To build an orthogonal base as needed for calculating differential cross sections, right-click Properties command.

Eiguro 5 11	Analysis windows properties : Calibration description (Harestua project)	? ×
Figure 5-11	Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs	
Calibration Fit Parameters		
Continuous function	✓ x0         ✓ x1         ✓ x2         x3         x4         x5         Insert           1 1/x0         1 1/x1         1 1/x2         1 1/x3         1 1/x4         1 1/x5         Hemove           Remove         All items         Properties         Y         Y         Y         Y	
	QKCancelHelp	

#### Predefined parameters

Offset and slit function parameters (SFP) are predefined parameters (i.e. parameters with reserved symbols not defined by the user).

In our example, the selected line shape is the error function (convolution of a gaussian and a boxcar), which is described by 2 SFP parameters : SFP 1 for the gaussian FWHM and SFP 2 for the boxcar width (SFP 3 and SFP 4 are ignored for this type of line shape even though they are checked).

Again in this case, a better accuracy is obtained when limiting the number of free parameters according to the information content of spectra. After optimisation, it has been found that best results are obtained when fixing the boxcar width (SFP 2) to 0.55 nm; the gaussian FWHM (SFP 1) therefore accounting for the wavelength dependency of the slit function is initialised to 0.3 nm (in order to speed up the convergence of the algorithm, it's recommended to initialise fit parameters to a value close to the real one).

Figure 5-12	Analysis windows properties : Calibration description (Harestua project)							
	Molecules Continuou	us functions	Predefined pa	arameters Shift a	nd Stretch 🛛 G	aps Outputs		
Calibration Fit Parameters							1/2	
D 1 C 1	Predefined param	n. Fit	∀al init	Val delta	Fit store	Err store	▲	
Predefined parameters	SFP 1		0.3	0.001				
	SFP 2		0.55	0.001				
	SFP 3	$\checkmark$	0.5	0.001				
	SFP 4	<b>V</b>	0.5	0.001				
							<u> </u>	
	ок	Cancel	1	delp				
			J					
_								

### Shift and Stretch

Shift (and stretch) can be applied either on the control spectrum (symbol **Spectrum** in this case) or on the solar spectrum (symbol **Ref** in this case). In order to avoid the need to interpolate the control spectrum, we recommend for the wavelength calibration, shifting the high-resolution solar spectrum instead of the control one. *Note that, in this case, the cross-section(s) defined in the* **Molecules page** must be shifted together with the solar spectrum.

To change the selection or add another item in the list of shifted quantities, rightclick the **Properties** command.

<b>Figure 5 43</b>	Analysis windows properties : Calibration description (Harestua project)							
Figure 5-13				·				
	Molecules Continuous functions	Predefined parameters	Shift and Stretch	Gaps Outputs				
Calibration Fit Parameters					1/5			
	Cross Sections and spectrum	Shift fit Stre	etch fit Scalin	ig fit				
Shift and Stretch	O3241(Ref;ring	Nor	ie None	Inset	1			
				Remove				
				Remove <u>A</u> ll items				
				Properties				
					<u>.</u>			
	<u>O</u> K <u>C</u> ancel	Help						
		· · · · · · · · · · · · · · · · · · ·						

### Analysis Window Properties

Creating a New Analysis Window Item in the Projects Tree To create a 'new Analysis Windows' item in the Projects tree,

- 1. select the predefined 'Analysis Windows' node and right-click the Insert item command;
- 2. rename the created 'new Analysis Windows' item; for example, give it the name 'BrO';

a dialog box with Analysis window properties appears.

Reference sele	ction Calib	oration	-Fitting int	erval (nm) — Display —		
<ul> <li>Automatic</li> </ul>	Ref	f only	Min 3	46 🔽 Spect	rum 🔽 Polyn.	🔽 Fits
O File			Max 3	159 🔽 Residu	ual 🔽 Predef.	Ref1/F
Files						
Reference 1	\data\harestu	ua\ha99h085i	u_35.ref		E	Browse
Reference 2	80 (	deg +/-	2 deg			
Residuals						Browse
Molecules Cor	ntinuous function	ns Predefin	ed parameters   S	hift and Stretch Gaps	Outputs ]	1/2
Molecules Cor	Intinuous function	ns Predefini	ed parameters   S	hift and Stretch Gaps	Outputs ]	1/2
Molecules Cor Cross Sec No2cb	ntinuous function tions Diff/Orth Different	ns Predefin Iog tial XS	ed parameters S	ihift and Stretch   Gaps   AMF None	Outputs   Fit display	1/2
Molecules Cor Cross Sec No2cb Bro	tinuous function tions Diff/Orth Different Different	ns   Predefin log tial XS tial XS	ed parameters S Interp/Convol Interpolate Interpolate	hift and Stretch Gaps	Outputs	1/2
Molecules Cor Cross Sec No2cb Bro o4	ntinuous function tions Diff/Orthu Different Different None	ns Predefin og tial XS tial XS	ed parameters S Interp/Convol Interpolate Interpolate Interpolate	hift and Stretch   Gaps   AMF None None None	Outputs	1/2
Molecules Cor Cross Sec No2cb Bro o4 ring	tions Diff/Orth Diff/Orth Different Different None Different	ns Predefin og tal XS tial XS	ed parameters S Interp/Convol Interpolate Interpolate Interpolate Interpolate	hift and Stretch   Gaps   AMF None None None None	Outputs	1/2
Molecules Cor Cross Sec No2cb Bro o4 ring	ntinuous function tions Diff/Orth Different Different None Different	ns Predefini log tial XS tial XS	ed parameters S Interp/Convol Interpolate Interpolate Interpolate Interpolate	hift and Stretch   Gaps   AMF None None None None	Outputs	1/2

#### Figure 5-14

Analysis windows properties

### Analysis Windows Properties

This dialog box is dedicated to the configuration of a spectral analysis window. In the above window, we have specified the **fitting interval** for BrO : 346-359 nm.

We have checked the option **Automatic** in the **Reference selection** frame. This means that a different control spectrum is automatically retrieved from the current data file for each twilight. The selection is performed around 80° SZA (cfr the SZA range specified in **Reference**).

The wavelength calibration is applied on the spectrum specified in **Reference 1**; the shift between both reference spectra is determined using a NLLS fitting approach and applied to align cross sections on the control spectrum.

The structure of the **Analysis Windows Properties** pages is very similar to the one used for the calibration (and already defined above).

### Molecules pages

In these pages, the absorption cross sections needed in the BrO fitting interval (see the BrO analysis setup, page 38) are listed and associated parameters are defined. Differential cross sections are generated by orthogonalisation to an orthogonal base defined in the **Continuous function** page. In order to avoid correlation between cross sections of similar shapes (e.g. when treating temperature effects by including two or several cross-sections of the same species), these cross sections can be orthogonalised with respect to each other.

#### Continuous functions page

In this page, the polynomial used for fitting the smooth part of the absorbance is defined. An orthogonal base is built with the main components of the polynomial in order to calculate differential cross sections. To build the orthogonal base, right-click the **Properties** command.

### Predefined parameters page

Among the available predefined parameters, we have selected the components of the offset (second order).

### Shift and Stretch pages

Shift and stretch (1st order) applied to the spectrum are selected for fitting. In these pages, a shift value to be applied (not fitted) to the  $O_4$  cross section is also defined. Note that several cross-sections can be grouped together. To access the configuration menu, right-click the **Properties** command. The units of the shift values are selected in **Projects** properties, in the **Analysis** page.



Figuro 5-16	Molecules Continuous functions	Predefined para	meters Shir	ft and Stretch	Gaps Out	puts	3/5
rigule 5-10							
	Cross Sections and spectrum	Sh init (px)	Stinit	St init (2)	Scinit	Sc init (2)	
Analysis Windows	Spectrum	0.000e+00	U	0.000e+00	0.00	U	
Properties	04	-3.7	0.00	0.00	0.00	0.00	
Shift and Stretch							
Dage 3/5							
1 age 5/ 5	I						
							• •

### Running Analysis

- is To analyse a spectra file, select it in the **Projects** tree and right-click the **Run Analysis** command.
  - Note if you right-click **Run Analysis** command from the parent node, the command will be performed on all spectra files defined in the selected project.



The first step of the program consists in applying in each sub-interval of the calibration window, a NLLS fit between the **Reference 1** spectrum (see **Analysis Windows properties**, page 43) and the high-resolution solar spectrum in order to determine the wavelength dependency of the shift and the fitted SFP (the Gaussian FWHM of an error function in this case) and then to optimise the calibration.

The Graphs window displays :

- the Reference 1 spectrum compared to the high-resolution solar spectrum convoluted with the calculated resolution of the instrument;
- $\succ$  the normalised residual;
- the ring and the ozone contributions to the optical density;
- the wavelength dependency of the shift;
- ▶ the wavelength dependency of SFP 1 (SFP 2 has been fixed);



The program searches for the first spectrum to analyse in the selected file and displays it.



The program then searches in the file for a spectrum to use as control spectrum for the current twilight. In the Data window, you can see the record number and the SZA of the selected spectrum. In this case, the same control spectrum will be used in BrO and OClO analysis windows. The alignment between the control spectrum and the reference is determined using a NLLS fit in both spectral analysis windows and the result of the fit is displayed.





The absorption cross sections are interpolated on the control spectrum grid.

Then, the program starts the spectral fitting in the BrO window according to the method selected in the **Analysis** page of **Projects** properties. Columns densities of all selected species are retrieved. Once the analysis is completed, residuals and fitting results are displayed in the **Graphs** window. The retrieved parameters are displayed in the **Data** window. We have zoomed the BrO fit.

If we go on, the program will perform the spectral fitting in the OCIO window.

## 6 The Workspace Environment Tree

The Workspace Environment tree contains two sections :

- in the Observation Sites section, the user can define a list of observation sites specified by their geolocation coordinates. These are used for correcting zenith angles data or performing a spectra selection (balloons and satellite measurements only).
- in the User-defined symbols section, the user must define the list of all relevant symbols that will be used in the configuration of projects and analysis windows. These symbols are needed to build cross sections files filters, to link AMF and cross section files, to name result files and for internal manipulations.

This chapter describes how to manipulate and configure **Observation Sites** and **User-defined symbols** items in the **Workspace Environment** tree.

Figure 6-1	🐼 Workspace Environment 🛛 🗖 🗙
riguie o-i	🖃 Observation Sites
	- Jungfraujoch
The Workspace	OHP
Environment Tree	Harestua
	Bruxelles
	User defined symbols
	- 03241
	Bro
	04
	ring
	Oclo
	- 03221
	no2cb
	no296
	03202
	resol
	03273
	i res

## 6.1 Creating a New Item in the Workspace Environment Tree

Creating a new item in the 1. Workspace Environment tree

select the parent node in the tree (Observation Sites or User-defined symbols node);

- 2. right-click to open the associated shortcut menu and select the **Insert Item** option or use the **Insert** button on the keyboard;
- 3. replace the new item text by the name of your choice;
- 4. validate to open the properties dialog box for configuring the object.

## 6.2 Observation Sites Properties

Figure 6-2 : Observation Site Properties	Observation Site properties : Harestua		
	Longitude in degrees (positive eastward)	10.753	
	Latitude in degrees (positive northward)	60.217	
	Site name abbreviation	HA	
	Altitude (meters)	600.000	
	KI	Help	

For geolocation coordinates, WinDOAS applies the following conventions :

- longitudes are positive eastwards; negative westwards;
- latitudes are positive northwards; negative southwards.

The **Site name abbreviation** (two characters only) is used to start the name of overpass files when no output file name is imposed by the user.

The Altitude field is used to calculate moon positions.

# 6.3 User-Defined Symbols Properties

Figure 6-3 : User-defined Symbols Properties

Symbol description	<u></u> K
BrO cross section (Wahner, 228K)	
	Help

Cross sections symbols can be completed with a short description.

# 6.4 Handling Items in the WorkSpace Environment Tree

Note that only one item can be selected at the same time in the tree. Use rightclick shortcut menus to handle items in the **WorkSpace Environment** tree.

The deletion of an item (observation site or user-defined symbol) is possible only if the symbol is not used in the configuration of a project or an analysis window.

## 7 The Projects Tree

This chapter describes the organisation, the handling and the configuration of the different kinds of items in the **Projects** tree. It covers :

- > the management of files and directories structures in the **Projects** tree;
- the configuration of projects;
- ➤ the parameterisation of the analysis;
- > the parameterisation of the wavelength calibration.

## 7.1 Description of the Projects Tree

WinDOAS is based on the notion of projects. A project can be defined as a set of files sharing the same configuration of analysis.

WinDOAS allows defining several projects in a session giving users the possibility to handle several analysis configurations. Files can be organised in user-defined folders and facilities are provided to manage large amount of files by loading only the directory structure in the tree.

The organisation of projects, analysis windows, spectra files and directories in a tree structure completed with the definition of right-click shortcut menus at each level of the tree makes the access, manipulation and configuration of all these objects very easy.



### Figure 7-1

Main Components of The Projects Tree

#### **Directories and Files Management** 7.2

Spectra Files	To insert one or several spectra files in the Projects tree,	
	1. select the predefined 'Raw Spectra' node under the parent project;	
	2. right-click the <b>Insert files</b> command;	
	3. in the usual <b>Open Files</b> Windows dialog box, locate the source path and select the spectra files.	
	Note that the file path doesn't appear in the tree. If you right-click the <b>Properties</b> command, you can display this path.	
User-defined Folders	You can organise spectra files in folders structures.	
	To create a 'new folder' item, right-click the <b>Insert folders</b> from the predefined Raw Spectra' node and give it a name.	
	The new folder item has the same properties as the predefined 'Raw Spectra' node i.e. you can insert spectra files in this folder or create a new level of folders.	
Directories structures	For large amount of files, creation and handling of spectra files and folder items in the <b>Projects</b> tree can become too heavy. In this case, you have the possibility to insert directory structures instead of spectra files.	

To insert a directory structure in the Projects tree, select the predefined 'Raw Spectra' node under the parent project or a user-defined folder and right-click the Insert Path options. The following dialog box appears :

Figure 7-2	Insert/Change	path	
Inserting a Directories tructure in the Projects Tree u can see the impact of this tion on the <b>Projects</b> tree in igure 7-1; spectra are in the <b>Spectra</b> sub-folder)	Path to insert Folder name File filter Files list	E:WINDOAS\Doc\SUM\Sample\Harestua Harestua  *.spe (\/Vildcards characters (?,*) may be used)  Spectra	(Browse paths) OK Cancel Help
	1	C Include subfolders	

Use the Browse paths pushbutton to locate the directories structure on your disk and specify a File filter ('spe' is the default extension for spectra files) to cut down the list of files. The Folder name field indicates the name of the item that will be inserted in the Projects tree and that will give access to the selected path. Check the Include subfolders to extend the files selection to subdirectories. Before handling spectra files items in the directory structure, validate your selection. To call back to the dialog box above and change the path or files selection, right-click the Properties option from the parent directory node.

### Struct (You car insertion

Figure

#### Four actions are possible on a spectra file : Handling Spectra

Browse	browses spectra in the selected file;
Run Analysis	analyses spectra using the configuration of the parent project and analysis windows; this step includes the correction of the wavelength calibration of the reference spectrum if it has been requested in the configuration of analysis windows;
Export in ASCII	saves the content of the original file to a new file in ASCII format; in the Windows Save dialog box, you can specify the name of the output file or 'automatic'. In the latter case, the original file name will be kept but the extension will be changed into ASC;
Run Calibration	uses the configuration of calibration defined in the <b>Project properties</b> (see page 58) to apply it on spectra in the selected file.

According to the selected item in a project, these actions are carried out by the way of shortcut menus :

- ⊳ on the selected spectra file;
- ⊳ on all spectra files of a user-defined folder and/or directories structure;
- ≻ on all spectra files of the selected project.

They can be applied on all projects defined in the Projects tree using the Projects Run Analysis and Run Calibration commands from the menu bar.

And finally, you can call back the Properties of a (sub-)directory item, select spectra files to process in the Files list and right-click to perform action.

some	Path to insert	E: WMNDOAS\Doc\SUM\San	nple\Harestua\spectr	a	
iles in ectory	Folder name File filter	spectra *.spe	(Wildcards character	s (?,*) may be used)	
	Files list	HA_2000_051.spe     HA_2000_052.spe     HA_2000_053.spe     HA_2000_054.spe     HA_2000_055.spe     HA_2000_055.spe     IA_2000_055.spe     IA_2000_055.spe	HA_2000_056.spe HA_2000_057.spe HA_2000_058.spe HA_2000_059. HA_2000_060. HA_2000_061. HA_2000_061.	HA_2000_063 HA_2000_064 HA_2000_065 Browse spectra Export in ASCII Run Analysis Run Calibration	Cancel Help

Fig

Apply analysis or individual spectra the selected sub-di

#### 7.3 Handling Items In The Projects Tree

Use right-click shortcut menus to handle items in the Projects tree. Note that only one item can be selected at the same time in the tree.

Common options	Remove	removes the selected item from the Projects tree;	
	Rename	renames the selected item;	
	Remove all items	removes all child items under the selected node;	
	Hide	hides the selected item (files, folders, directories or analysis windows) without loosing its properties; hidden items are not considered for the spectra analysis;	
	Show hidden items	recovers under the selected node, all previously child items (files, folders, directories or analysis windows);	
	Copy/Cut/Paste	these operations are supported for analysis windows and projects nodes but not for files and folders; when pasting an item in another configuration file, the list of symbols is updated if needed.	
Projects	Right-click the <b>Properties</b> option to open the <b>Project properties</b> dialog box to configure the selected project.		
	Select the following Pr	1g <b>Projects</b> commands in the menu bar :	
	New	to create a "new project" item in the Projects tree;	
	Remove All	to remove all projects items from the <b>Projects</b> tree;	
Analysis Windows	View cross sections	displays, in the main <b>Graphs</b> window, all cross sections defined in the selected analysis window;	
	Properties	opens the <b>Analysis windows properties</b> dialog box and parameterise the selected analysis window;	
	Insert item	inserts a new 'analysis windows' item in the parent project;	
	Sort items	sorts the analysis windows list in alphabetic order;	

Spectra files See previous section.

# 7.4 Project Properties

Options for configuring projects are separated according to the category they belong to, among the following pages of the **Projects Properties** dialog box :

Spectra	selection of spectra to analyse;	
Analysis	selection of the analysis method;	
Filtering	selection of a filtering method to apply to spectra and absorption cross sections;	
Calibration	wavelength calibration and slit function characterisation;	
Undersampling	selection of a method for calculating undersampling cross sections;	

Instrumental	specification of the spectra files format and instrumental corrections;
Slit function	definition of a slit function for convoluting cross sections in real-time;
ASCII results	output file configuration;
NASA-AMES results	output configuration (specific NDSC format).

Figure 7-4 : Project	roject properties : Harestua
Properties	Instrumental Silt Function ASCII Results NASA-AMES Results
	Spectra Analysis Filtering Calibration Undersampling
The Spectra Tab	
Pane	Display SZA range (deg)
i age	Spectra 🔽 Data Min 0.00
	✓ Fits Fits/page (∀xH) 4 X 4 Max 90.00
	I Pause Detta 0.00
	Spectra Nº range
	Min 0
	Geolocations Max 0
	Mode Circle area
	Radius 0.00 km Files
	Longitude 0.00
	Latitude 0.00

Spectra selection	The <b>Spectra</b> tab page is dedicated to the selection of spectra to display and/or to analyse. The selection of spectra can be made on :	
	SZA	specify the SZA range and the SZA step in degrees;
	Spectra N°	specify a range of record numbers;
	Geolocations	specify geolocation coordinates.
Geolocation coordinates	The geolocation coordinates condition is useful for limiting the selection of spectra to a geographic area (balloons and satellites measurements):	
	Circle area	circle area defined by its centre coordinates and its radius;
	Rectangle area	rectangle area defined by longitudes and latitudes ranges;
	Circle area using obs. sites list	defines circle area around locations defined in the <b>Observation Sites</b> list of the <b>Workspace environment</b> tree (overpasses for satellites measurements).

Let geolocation fields at "0.0" if you don't want to perform this kind of selection.

Files option If the selected spectra files format supports record names and dark currents, the Names and Dark current buttons must be respectively checked. If these data must be retrieved from files different from the original spectra one, the default file path and name are then, the same than the original spectra ones but with the respective "nms" and "drk" extensions. The default record names and dark current files names can be changed in the properties of the associated spectra file item in the **Projects** tree.

### Display During spectra analysis, you can display :

- ➢ spectra;
- ➢ fits;
- data.

By default, WinDOAS pauses between each display. Spectra browsing and analysis can be performed in a non-stop way by unchecking the **Pause** button in the **Display** frame. In this case, you can speed up spectra processing by disabling also the **Spectra, Fits** and **Data** displays.

The number of graphs to display depends on the number of parameters to fit in your application. For each defined analysis window, you can select individually the fits to display (see section 7.6, page 67). By default, WinDOAS displays one graph at a time in the main **Graphs** window and you pass from one graph to the other using appropriate toolbar buttons. But you can change at your best convenience, the number of graphs to display vertically (V) and horizontally (H) in one time in the main **Graphs** window by filling the fields **Fits/page** in the **Spectra** tab page of **Project properties** dialog box. The maximum number of graphs per window is limited to 20.

Figure 7.5 · Project	Project properties : Harestua
Properties	
Topenies	Instrumental Slit Function ASCII Results NASA-AMES Results
	Spectra Analysis Filtering Calibration Undersampling
The Analysis Tab	
Pane	
r ago	Analysis Method Optical density fitting
	Least Squares fit No weighting
	Shift and stretch units Pixels
	OK Cancel Help

The Analysis Method In the Analysis tab page, you can select the retrieval method among three possible choices :

- Optical density fitting;
- Intensity fitting (Marquardt-Levenberg+SVD);
- ▶ Intensity fitting (Full Marquardt-Levenberg).

These methods differ by the way the Beer-Lambert's equation is expressed and solved. Details are given in section 4.3.

- Shift and stretch units Selection of the unit, **Pixels** or **Nanometers**, in which shift and stretch parameters are expressed.
  - Interpolation Linear and cubic Spline interpolations are implemented. Spline interpolation is selected by default.

Least-Squares Fit The fit can be weighted if uncertainties on measurements are known. Instrumental and Statistical weighting options are foreseen but not yet implemented.

Figure 7.6 · Dupingt	ect properties : Harestua	? ×	
Figure 7-6 : Project		deres	
Properties	Instrumental Slit Function ASCII Results NASA-AMES Results		
	Spectra Analysis Filtering Calibration Undersampling		
The Filtering Tab			
Page	Low Pass Filtering		
	Fitter Type Kaiser 💌		
	Cutoff freq. 0.60 Tolerance (dB) 80.00		
	Pass band 0.10		
	Number of iterations		
	High Pass Filtering		
	Filter Type None 💌		
ſ	OK Cancel Help		

Spectra and cross sections can be filtered using one of the filters proposed in the **Filtering** tab page :

- ➢ Kaiser;
- ➢ Boxcar;
- Gaussian;
- ➢ Triangular;
- Savitzky-Golay;
- Odd-even pixels correction;
- Binomial.

According to the selected filter, different fields must be filled in. The selected

filtering can be disabled for individual cross sections in the **Analysis windows** properties dialog box (see page 70).

Select filter type None if you want to disable spectra and cross sections filtering.

The program supports low-pass and high-pass filtering. In the latter case, cross sections are filtered by subtracting a fitted polynomial or a smooth spectrum calculated by filtering the original cross sections a high number of times. Differential cross sections can be obtained by this way. For the moment, high-pass filtering is supporting only in optical density fitting mode.

Figure 7-7 : Project	Project properties : Harestua
Properties	Industrated Cliffornation CCCII Results NICCA AMEC Results
	Spectra Analysis Filtering Calibration Undersampling
The Calibration Tab	
Page	Solar ref. file e:\data\xs-hr\kurucz\Kur01.ktz
	Analysis Method Optical density fitting
	Fit SFP Error function
	Display Polynomial degree Window limits (nm)
	I Spectra I Fits Shift 3 Min 330
	Residual Shift/SFP SFP 3 Max 390
	Number of sub-windows 10
	Fit parameters
	QK Qancel Help

Wavelength Calibration And Slit Function Characterization Wavelength calibration and instrumental slit function characterisation are performed using a non-linear least-squares (NLLS) fitting approach where measured intensities are fitted to a high-resolution solar atlas spectrum degraded to the resolution of the instrument.

The wavelength interval used for this calibration must be specified in the **Window limits (nm)** box. This interval is subdivided in a number of equally spaced sub-windows and, shift and slit function parameters (SFP) are fitted using the NLLS approach in each interval. The shifts values determined in all sub-windows are then used to reconstruct an accurate wavelength registration. Similarly, the wavelength dependent slit function is determined by polynomial fitting through individual SFP values. These two information are subsequently used e.g. to build undersampling cross-sections.

The current tab page comprises the following fields and buttons for configuring calibration :

	Solar ref. file	2	the solar reference spectrum (see remarks below);
	Analysis Me	thod	the fitting method (can be different from the one used for spectra analysis);
	Fit SFP		uncheck the <b>Fit SFP</b> button if you don't want to characterise the slit function;
	Display		specify which graphs to display at the end of the procedure;
	Polynomial	degree	the degree of the polynomial used in the final determination of the wavelength registration (shift) and the wavelength-dependent slit function (SFP);
Window limits Number of sub-windows		its	the wavelength window on which the calibration correction must be applied;
		ſS	the number of calibration intervals;
	Fit Paramete	ers	This pushbutton gives access to a dialog box using a property sheet similar to the one defined in <b>Analysis windows properties</b> , in which you can specify parameters to fit and atmospheric absorption to take into account for the fit evaluation. This dialog box is detailed in section 7.7.
Solar reference spectrum	There are two	o cases to	consider :
	<ol> <li>You known character with a so slit funct</li> </ol>	ow the sl rise it onc olar refere tion on th	lit function of your instrument and you don't want to the again; then you have to provide the <b>Solar ref. file</b> field ence spectrum previously convoluted with the instrument e grid of the spectrum to calibrate.
	2. You wan provide Because high-reso	nt to chara the <b>Sola</b> WinDOA plution so	Acterise the slit function of your instrument; then you must r ref. file field with a high-resolution solar spectrum. AS uses FT in order to speed up the NLLS algorithm, a lar spectrum sampled on a constant grid is expected.
Line Shape Selection	When the button <b>Fit SFP</b> is checked, you have to select in the list on the right one of the basic line shapes to fit (see page 27). Extra data can then be requested according to your selection. The configuration of parameters to fit is done in the dialog box called by the way of the <b>Fit Parameters</b> pushbutton.		
Display	Spectra	compare	es the observed and calculated spectra;
	Fits	displays	fits of Ring effect and/or atmospheric absorptions if any;
	Residual	displays	the residual of the fit;
	shift/SFP	displays paramete	the wavelength dependency of the shift and slit function ers fitted over the selected wavelength window.

Figure 7-8 : Project	Project properties : GOME				
Properties	Spe	ctra	Analysis	Filtering	Calibration
The I Indersampling	Ur	idersampling	Slit F	unction	ASCII Results
Tab Page	Sol	arref. file	f:\gwindoas.cd	i-r\data\Nevvkpno_v	ac.ktz
Ū	Method	Fixed phas	ie 🔻		
	Shift	0.008			
	Sint Cool				
	<u></u> K		ancel	Help	

Method See the Undersampling topic in 4.6 section to get further details about undersampling correction.

In this page, three methods are proposed :

From file	Undersampling cross sections are provided in files like usual cross sections. These files might be created using the WinDOAS <b>Undersampling</b> tool (see 8.5);
Fixed phase	WinDOAS uses the information derived from the NLLS wavelength calibration and slit function determination to create the undersampling cross sections, with a fixed value of the shift. The selection of this method is recommended;
Automatic phase	The undersampling cross sections are calculated at each iteration of the analysis procedure, using the fitted value for the shift between control and measured spectra. This method is rather time consuming and only applicable for testing purposes.

- Shift The Shift field operates only in Fixed phase method.
- Solar ref. file In Fixed phase and Automatic phase, undersampling cross sections are calculated from a high resolution solar spectrum convoluted on an oversampled and an undersampled grid with the instrument slit function (see the Supported line shapes, page 27).

### If you don't want to correct for undersampling :

Select **File** method and for all the **Analysis Windows** (see page 72) defined under the current project, uncheck the fitting of undersampling cross sections in the **Predefined parameters** page in intensity fitting mode or remove **Usamp1** and **Usamp2** cross sections symbols from the **Molecules** page in optical density fitting.

Figure 7-9 : Project	Project properties : Harestua	? ×
Properties	Spectra Analysis Filtering Calibration Undersampling	1 1
	Instrumental Slit Function ASCII Results NASA-AMES Results	12
The Instrumental Tab		1
Page	Format ASCII	1003
	Observation site Not Specified	
	Detector Size Format Data to read with spectra 1024 Column Data to read with spectra Zenith angle Decimal time Date (DD/MM/YYYY) lembda	
	Calibration file	
	Instr. function	
	<u>QK</u> <u>Cancel</u> <u>H</u> elp	

In the **Instrumental** page, you must specify the **format** of the spectra files to process. The fields to fill in depend on the selected format. For a detailed description of a specific format or to include a new spectra files format, contact authors.

A preliminary calibration is expected in the **Calibration file** field. Instrumental corrections can be applied on spectra (those are divided by the vector given in the **Instr. function** field).

The **Observation site** field is used to (re-)calculate SZA from geolocation coordinates given in the **Workspace Environment tree** and to build automatic output file names (see **The ASCII Results Tab Page**, page 63).

Figure 7-10 : Project	Project properties : Harestua
Properties	
	Spectra Analysis Filtering Calibration Undersampling
	Instrumental Slit Function ASCII Results NASA-AMES Results
The Slit Function Tab	
Page	Solar ref. file C:Wy Data\Cross sections and AMF\XS-HR\Kurucz\Kur
	Slit Function type Error function, wavelength dependent
	Slit Function C: Wy Packages WinDOAS Data Harestua Sfp0_full.slf
	Boxcar width 0.55 nm
	Apply Fwhm correction
	OK Cancel Help

WinDOAS gives the possibility to convolute cross sections in real time using the information provided by the wavelength calibration procedure.

If you don't want to characterise the slit function, cross sections can be convoluted before spectra analysis using the line shape specified in the **Slit Function** tab page. Note, that in this case, it is preferable to use pre-convoluted cross sections obtained using the **Convolution/Filtering** tool (see section 8.3). Refer to the same section for a list of all possible slit function types.

Fields to fill in depend on the selected slit function type.

The **Solar ref. file** entry field must be filled in if the convolution with  $I_0$  correction is requested for at least one cross section.

If the dependence of the slit function with the detector temperature is known, WinDOAS can determine the difference of resolution between the control spectrum and each spectrum to analyse and use this information to convolute cross sections in real-time. To apply this, select as **Slit Function type**, a wavelength and temperature dependent one, enter the appropriate file and check the button **Apply Fwhm correction**. Note that this method is heavy in processing time.

Figure 7-11 : Project Properties	Project properties : Harestua	×
	Spectra Analysis Filtering Calibration Undersampling Instrumental Sitt Function ASCII Results NASA-AMES Results	a and a second
The ASCII Results Tab Page	Output Path or File	$\sim$
	Analysis Fluxes 330;350;380;	
	Calibration Cic 380/330;	
	Spec No     A       Name     Date & time (YYYYMMDDhhm       Date & time (YYYYYMMDDhhm     >>       Date (DD/MM/YYYY)     >>       Time (hr.mm:ss)     <	
	QK Cancel Help	

Output Files Format This page is dedicated to the selection of calibration and analysis results to output :

- when the Analysis button is checked, the selection of the results file and the selection of fields to output are enabled. The program produces tabbed separated columns ASCII files that you can easily load in spreadsheets like Excel. There is one line per processed spectrum and the first line is reserved for columns titles;
- when the Calibration button is checked, wavelength calibration results are saved before analysis results. Lines start with the semicolon (;) character.

Automatic Creation Of The Output Path or File pushbutton allows you to locate the path in which results have to be saved. You can specify the name of the output file. If you The Output File Name specify "automatic" as the file name, WinDOAS will create automatically output files names depending on the measurement date and the observation site specified in the Instrumental tab page (see page Error! Reference source not found.Error! Bookmark not defined.) as follows : <output path>\XXYYYYMM.ASC where : output path is the user-defined path; XX is the abbreviation of the observation site selected in the Instrumental tab page; if no observation site is specified, the abbreviation XX is used; YYYY is the year of measurement; MM is the month (zeros padded) of measurement; ASC is the default file extension for ASCII files; Fluxes and Colour A flux is the mean signal calculated over seven pixels around the indicated indexes wavelength. To output Fluxes, specify the central wavelengths separated by the semi-colon (;) character. For example, 330; 350; 380; will calculate and output fluxes at 330, 350 and 380 nm. A colour index is the ratio of two fluxes. To output colour indexes (Cic), just write the ratios separated by the semi-colon (;) character as in the example below: 380/330; 380/350; In WinDOAS, the output is fully configurable. The selection of analysis results **Output Selection** (e.g. slant column densities or fitted non-linear parameters) is made individually in Analysis windows properties dialog boxes (see section 7.6). The ASCII Results tab page proposes the following list of spectra-related information items. Select those that are available in your spectra files. Spec No the record number; Name the record name; Date & time (YYYYMMDDhhmmss) measurement date and time in the specified format; Date (DD/MM/YYYY) measurement date; Time (hh:mm:ss) measurement time; Year the year of data; Day number the day number; Fractional day the fractional day number; Fractional time the fractional time;

Scans	the number of scans;
Tint	the exposure time;
SZA	SZA;
Chi Square	the $\chi^2$ for each analysis window;
RMS	the RMS for each analysis window;
Tdet	the detector temperature;
Sky obs	information of the sky during the current measurement;
Ref SZA	the SZA of the control spectrum if this data is known;
Longitude	longitude of the observation site;
Latitude	latitude of the observation site;
Altitude	altitude of the observation site;

Figure 7-12 · Project	Project properties : Harestua			
Properties		N		
	Spectra Analysis Filtering Calibration Undersampling			
	Instrumental Slit Function ASCII Results NASA-AMES Results	14		
The NASA-AMES		14		
Results Tab Page	Results nath E: (AINDOAS)Doc/SLM(Semple)			
rtesuits rub ruge				
	Save Instrument Reticon PDA EG&G	and a		
		Jung		
	NO2 Not saved 🔽 OCIO Not saved 💌			
	O3 Not saved  BrO BrO			
	,			
	OK Cancel Help			

The NASA-AMES (Research Centre) is an ASCII based reporting system which includes several categories of self-described data formats. The Ames format is used to provide for example, the NDSC database.

WinDOAS can output results in the Ames format for four molecules, NO<sub>2</sub>, O<sub>3</sub> (slant and vertical columns), BrO and OClO (slant columns only) using the template NASAAMES.HDR provided with the program.

For each molecule, you can select the analysis window in which the results to output are calculated. The selected analysis window must include a symbol whose syntax is exactly the same as the indicated one.

As for **ASCII results**, you can specify in the usual Windows Save dialog box, the name of an output file or just the word **"automatic"**. In the latter case, WinDOAS will create automatically output files names according to the syntax described above but will use the UVV extension instead of the ASC one.

The **Instrument** and **Experiment** fields must be filled in. They will be used in place of \$INSTR and \$EXPERIMENT macros (edit the NASA-AMES.HDR file) when building the Ames file header.

## 7.5 Analysis Windows Properties

Figure 7-13 : Analysis ties : BrO (Harestua project) ? × Analysis wind Windows Properties Calibration Fitting interval (nm) Display Reference selection Automatic 346 Spectrum 🔽 Polyn. 🔽 Fits Ref only • Min O File Max 359 Residual V Predef. Ref1/Ref2 Files ..\data\harestua\ha99h085u\_35.ref Browse Reference 1 80 Reference 2 deq +/- 2 deg Browse Residuals Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 1/2 Cross Sections Diff/Orthog Interp/Convol AMF Fit display -None Differential Interpolate 2 Differential XS Bro Interpolate Insert 1/Rel o4 Interpolate Remove None Differential XS Interpolate ring V -Properties F oĸ Cancel Help Right-click shortcut menus

> This dialog box is dedicated to the parameterisation of the main spectral analysis. The possibilities of analysis configuration are very large. This section describes the main components of the **Analysis windows properties** dialog box. Next section will guide you in the definition of parameters to fit.

Reference Spectra A reference spectrum is assumed to be correctly wavelength calibrated. If a wavelength calibration correction is applied anyway on this spectrum, cross sections are re-interpolated on the new grid.

WinDOAS gives the possibility to define two reference spectra (**Reference 1** and **Reference**). If only one reference spectrum is specified, it is used as control spectrum i.e. the  $I_0$  spectrum in the Beer-Lambert law. If two reference spectra are given, the wavelength calibration is applied only on the first one (**Reference 1**); the shift between both reference spectra is then determined using a NLLS fitting approach in order to align cross sections on the second reference spectrum (**Reference**).

WinDOAS allows selecting different control spectra from the current data file by specifying a range in SZA (option **Automatic** in the **Reference selection** frame). For each twilight, the spectrum in the file with the SZA closer to the one specified by the user will be selected as control spectrum. If the SZA range is lower than SZA supported by the current file, the spectrum in the file with the minimum SZA will be selected for both twilights.

Calibration To calibrate the reference spectrum, select **Ref only** in the list proposed by the **Calibration** frame. The wavelength interval on which the calibration procedure is applied is selected in the **Calibration** page of **Projects properties** (see page 58). It can be different from the one selected for spectra analysis.

The option **Spectra only** is useful for tropospheric measurements where no reference spectrum is needed.

The option **Ref+spectra** allows applying a calibration correction on both reference spectrum and spectra to analyse. This option has been designed to handle spectra recorded with unstable (unthermostated) instruments where the spectral resolution can vary a lot from one spectrum to another. In this case, the resolutions of both spectra are matched to the resolution of the less resolved spectrum, and the absorption cross-sections can be convoluted in real-time to the same effective resolution (see 7.6).

Select None if you don't want to correct the wavelength calibration of the reference spectrum.

Fitting interval This is the spectral interval where the current analysis configuration is applied.

- Display After completion of the analysis, WinDOAS can display graphically:
  - ➤ the normalised spectra before analysis;
  - $\succ$  the calculated residuals;
  - the fitting results for each species and for predefined parameters (offset, undersampling);
  - the continuum part fitted by a polynomial.

Check the appropriate buttons for selecting the graphs you want WinDOAS to display. An individual selection of cross sections and predefined parameters can still be performed (see section 7.6).

Files	Reference	Reference spectra (see above)
	Residuals	If you want to perform further studies on residuals, you can specify the name of an output file in which calculated residuals will be saved. A line of the resulting file contains the calculated residual for a spectrum record.
itting Parameters	The user must define the fitting parameters in the appropriate pages of property sheet at the bottom of the <b>Analysis windows properties</b> dialog b The different pages are :	
	Molecules	definition and configuration of the list of cross sections to fit;
	Continuous functions	specification of the degree of the polynomial fitting the continuous component of the absorbance;
	Predefined parameters	this page proposes several predefined parameters such as offset, undersampling ;

F

Shift and Stretch	shift and stretch can be applied to any spectral item;
Gaps	gaps can be introduced in the fitting window (e.g. to eliminate bad pixels);
Outputs	selection of the calculated columns densities to save in the output file.

According to the associated type of parameters, the selected page proposes several columns of options to fill in, to check or to select from a multiple choice. The conviviality of the analysis parameterisation is enhanced by the use of rightclick shortcut menus to handle lists of items in the different tab pages. Because of the complexity of the whole set of options, the structure is detailed page after page in the following section.

## 7.6 Configuration of the Fitting Parameters

### The Molecules Pages

The options for defining and configuring the list of cross sections to fit are distributed in two **Molecules** pages (see figures below) :

Figure 7-14 : Analysis	Molecules Continuou	us functions Prede	efined parameters Shi	ft and	Stretch Gaps Outp	uts	1/2
Windows Properties	Cross Sections	Diff/Orthog	Interp/Convol		AMF	Fit display	
	Bro	Differential XS	Interpolate		SZA only		
Molecules	04	None	<u>I</u> nsert	•	None		
Page 1/2	ring	Differential XS	<u>R</u> emove		None	<b>V</b>	
	03241	Differential XS	Hemove <u>All items</u>		None	<b>V</b>	
	I		<u>P</u> roperties	•	Cross section file		<b>_</b>
					AMF file 😽		T

Figure 7-15 : Analysis Windows Properties

> Molecules Page 2/2

Cross	Sections	Filter	Cc Fit	Cc init	Cc Delta	Cc I0	_
Bro			<b>L</b>	0.	0.01		
o4			4	0.	0.01		
ring			1	0.	0.01		
03241			<b>V</b>	0.	0.01		

Molecules are characterised by their cross section. They are represented by symbols previously defined in the **Workspace Environment** tree (see chapter 6). WinDOAS uses these symbols to build cross section files filters :

- Cross section files names must imperatively start with the symbol name as prefix followed by the underscore character !
- There is no constraint on the cross section file extension; the default one used by WinDOAS for creating cross sections files filters starts with "xs".

To insert a new cross section in the <b>Molecules</b> pages :
<ol> <li>right-click the Insert option from the property sheet; the usual Open Files Windows dialog box appears with, as Files of type, the list of available cross sections; this list is build from the list of symbols previously defined in the Workspace Environment tree and is updated after each new insertion in order that two identical symbols are not defined in the same analysis configuration;</li> </ol>

Inserting A New Cross	Open Cross s	ections Files			?	×
Section	Look jn: [	🔄 Xs	•	1	8-0- 8-0- 8-0-	
List of files depending on the selected cross section	iningxs     ining_2.xs     ining_g.xs     ining_g.xs     ining_g.xs     ining_g.xs     ining_g.xs     ining_g.xs     ining_g.xs	ining_kgv.xs ining_t.xs				
	File <u>n</u> ame:				<u>O</u> pen	
List of available cross sections (Com, Usamp1 and Usamp2	Files of <u>t</u> ype:	ring Cross Sections (ring_*.xs*) Com Cross Sections (Com_*.xs*) Usamp1 Cross Sections (Usamp1_*. Usamp2 Cross Sections (Usamp2_*. Usamp2 Sections (Tom_*.vs*)	(S*) (S*)		Cancel <u>H</u> elp	
are predefined symbols)		OclO Cross Sections (OclO_*.xs*)	k			

- 2. select in the Files of type list, the cross section you want to insert;
- 3. in the cut down list of files, select the file containing the cross section to fit.
- To control or modify the cross section file associated to a cross section symbol Handling cross sections defined in the Molecules pages, right-click the Properties of the selected items in the list symbol; To remove a cross section from the list, select the cross section symbol and right-click the Remove option. The option is greyed if another cross section is orthogonalised to the selected one or if the selected symbol is used in the Shift and Stretch pages. Differential cross sections Differential cross sections can be generated either by orthogonalisation or highpass filtering according to the definition or not of an orthogonal base formed with the component vectors (generally, a base of order 2) of the polynomial defined in the **Continuous functions** page (see the next section, page 70). There are three possible choices : None use the original cross section; **Differential XS** generate a differential cross section : ≻ by orthogonalisation if an orthogonal base is defined in the Continuous functions page;

	a cross section in the proposed list	the selected cross section is orthogonalised to the orthogonal base (if defined) and to another cross section defined in the list (orthogonalisation in cascade is allowed).
	In the latter case, you example, two O <sub>3</sub> cross available cross section <b>Molecules</b> page excep are added in or remove	can avoid any correlation between two cross sections (for sections measured at different temperatures). The list of ns includes all cross sections symbols defined in the ot the one selected. It is updated as cross sections symbols ad from this page.
Interpolation/Convolution	This column describes	the action to perform on the selected cross section :
	None	the selected cross section is assumed to be correctly aligned on the reference spectrum grid; so, no action has to be performed on the cross section (for example, a user-defined undersampling cross section in optical density fitting);
	Interpolate	this is the usual choice; the selected cross section (a pre-convoluted or a high resolution one) has to be interpolated on the grid of the reference spectrum;
	Convolute Std	this option gives the possibility to convolute a cross section in real-time using either the information on calibration and slit function provided by the wavelength calibration procedure or the user-defined slit function specified in the <b>Slit Function</b> tab page of <b>Project</b> <b>properties</b> (see page 62). If the wavelength calibration/slit function characterisation is applied on the control spectrum and spectra to analyse (see page 66), the cross section with <b>Interp/convol</b> action set to <b>Convolute Std</b> is convoluted with the poorer resolution;
	Convolute I0	the cross section is convoluted with $I_0$ correction using the concentration defined in the column <b>Cc I0</b> (see below);
	Convolute Ring	in the same way, the program can generate a ring cross section; the expected cross section file must be a pre- convoluted ring cross section calculated by the WinDOAS <b>Ring tool</b> (see section 8.4) on a high- resoluted grid;
AMF	If you want to calculate using wavelength deper <b>AMF</b> column (option right-clicking the <b>Prop</b>	e vertical columns or if you want to correct a cross section endent AMF, select one of the options proposed in the different from the <b>None</b> one) and specify an AMF file by <b>erties</b> of the selected cross section.
	Possible choices :	
	SZA	this is the option usually selected to calculate vertical columns; the AMF depends only on the SZA;
	Climatology	select this option to introduce climatological AMF; the AMF depends on the SZA and the calendar day;

Wavelength select this option to correct the selected cross section using wavelength dependent AMF (modified DOAS).

The format of the requested AMF file depends on the option selected in this column. The formats to use for AMF files are described in **Annex**.

- Fit display This column is active only if the Fits button is checked in the Display frame of the current dialog box. It allows selecting which cross sections fits are to be displayed.
  - Filter This column is active only if a low-pass filter has been selected in the Filtering tab page of **Project properties** (see page 57). It allows defining individually which cross sections are to be filtered.
  - Cc Fit If you uncheck the fit of a cross section, the weight of the cross section in the optical density is fixed at the concentration value given in the Cc Init column (see the next column).
  - **Cc** Init If the selected cross section is not fitted (the **Cc** Fit button is unchecked), this is the fixed concentration value to take into account in the optical density. This is also the initial concentration value for the non-linear algorithm in intensity fitting.
- **Cc Delta** This is the delta value used by the non-linear algorithm to calculate numerically partial derivatives of the fitting function for the selected non-linear parameter in intensity fitting. It determines the accuracy with which the algorithm approaches the minimum for this parameter. The default value shouldn't be modified except if the system seems not to converge.
  - Cc 10 This column is active only if the Convolute I0 option has been selected for a cross section. This is the concentration of the cross section used to calculate the theoretical optical depth in convolution with  $I_0$  correction.

### The Continuous Functions Page

In this page, check the components in  $x^n$  or  $1/x^n$ , of the polynomial fitting the continuous part of spectra to analyse.

	▼ ×0	<b>▼</b> x1	<b>▼</b> x2		□ ×4			
inuous functions	🗖 1/x0	1.0x1	☐ 1/x2	 1/x3	<b>1</b> /x4	1 <i>b</i> x5		
							-	

If you want to calculate differential cross sections by orthogonalisation, you must create an orthogonal base (see page 69). For that, right-click the **Properties** option to open the following dialog box and select in the list of available components, the ones that will be used to create the orthogonal base.

Figure 7-18 : Analysis Windows Properties	Orthogonal base		
	Available	Selected	
Creating an orthogonal base	×3	x0 x1 x2	Use <u>set</u> and <u>set</u> pushbuttons to move items from one list to the other.
	QK	Help	

### The Predefined Parameters Pages

WinDOAS proposes several predefined fitting parameters:

- ➤ offset;
- undersampling (Usamp1 and Usamp2);
- scaling factor for the control spectrum (Sol);
- common residuals (Com).

Options for configuring these parameters are distributed in two pages (see figures below) :

Figure 7-19 : Analysis Windows Properties

Predefined parameters Page 1/2

Predefined param.	Fit	Val init	Val delta	Fit store	Err store	_
Offset (Constant)		0.	0.001			
Offset (Order 1)	1	0.	0.001			
Offset (Order 2)	<b>V</b>	0.	0.001			
Com		0.	0.001			

Check buttons in the Fit column for selecting the fitting parameters.

Val init and Val delta are respectively the initial value and the convergence factor, parameters used by the NLLS algorithm. In general, these values should not be modified, except in case of convergence problems.

Fit store and Err store columns are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). In these columns, you can check individual results and standard deviations of fitted parameters.

Figure 7-20 : Analysis	Molecules	Continuous	functions	Predefined parameters	Shift and Stretch	Gaps Outputs	2/2
Windows Properties	Prede	fined param.	Val min	Valmax			
	Offse	t (Constant)	0	0			
Predefined parameters	Offse	t (Order 1)	0	0			
Page 2/2	Offse	t (Order 2)	0	0			
	Com		0	0			•
							T

	If the non-linear algorivalues for the selected condition is applied.	thm does not converge, you can fix the range of plausible parameter. If <b>Val min</b> and <b>Val max</b> are set to 0, no range
Fitting the offset	WinDOAS allows fitti of the dark current, or 2 are supported. Sele buttons in the <b>Fit</b> colu	ng an instrumental offset (e.g. due to incorrect correction residual stray-light). Polynomial offsets of order 0, 1 and ct the degree of polynomial by checking the appropriate mn.
Fitting the undersampling	The way the undersam on the fitting metho correction method in page 60) is set to <b>File</b> :	pling correction is implemented slightly differs depending d (DOAS or intensity fitting). If the undersampling the <b>Undersampling</b> tab page of <b>Project properties</b> (see
	in DOAS fitting	enter undersampling cross sections files as any other cross sections in the <b>Molecules</b> page using <b>Usamp1</b> and <b>Usamp2</b> predefined symbols;
	in Intensity fitting	right-click the <b>Properties</b> option from <b>Usamp1</b> and <b>Usamp2</b> symbols in the <b>Predefined parameters</b> page to specify or update the user-defined undersampling cross sections files.
	Check <b>Usamp1</b> and/ undersampling correct	or <b>Usamp2</b> buttons in the <b>Fit</b> column in all cases of ion except in <b>File</b> method under DOAS fitting.
Fitting the common residual	In case systematic structures section can be c undersampling correction	actures are identified in the residuals, a common residual reated and introduced in the spectral analysis. As for the ion, the implementation depends on the fitting method:
	in DOAS fitting	enter the common residual cross section file as any other cross sections in the <b>Molecules page</b> using the <b>Com</b> predefined symbol;
	in Intensity fitting	right-click the <b>Properties</b> option from the <b>Com</b> symbol in the <b>Predefined parameters</b> page to specify or update the user-defined common residual cross section file.
	Check the <b>Com</b> buttor	n in the <b>Fit</b> column in intensity fitting only.
Fitting a scaling factor for the control spectrum	Check the <b>Sol</b> button i	n the <b>Fit</b> column of <b>Predefined parameters</b> page.
The Shift and Stretch Pages	In WinDOAS, you can any spectral item (spec (five pages in total), yo one) and cross sections	n fit or apply a shift, a stretch and/or a scaling factor for ctra and cross sections). In the <b>Shift and Stretch</b> pages u can select spectra (the spectrum to analyse or the control s to shift.
Inserting a cross section To fit or apply a shift and/or a stretch to a cross section, right-click the Insert option from the Shift and Stretch pages to open the following dialog box.

Figure 7-21 : Analysis	Set of cross sections for shift and stretch					
Windows Properties Selecting a cross section to shift	Available Bro No2cb 03221 03241 Ref res resol rina	Selected				
	<u>Ō</u> ĸ	<u>C</u> ancel <u>H</u> elp				

This dialog box presents the list of symbols defined in the **Molecules page** and completed with the **Spectrum** and **Ref** symbols (these latter two symbols usually refer respectively to the spectrum to analyse and the control one).

You can select the cross section to shift in this list. If you select several symbols, the same shift and stretch parameters will be applied to all items of the selection.

After validation of the selection, WinDOAS automatically updates the list on the left so that a symbol can not be selected twice.

Note that when a cross section symbol has been selected in the **Shift and Stretch** page, it can not be removed anymore from the **Molecules** page.

Modifying the selection Y

You can modify at any time a selection of symbols in the **Shift and Stretch** page by right-clicking the **Properties** option from this selection to call back the dialog box in figure 7-21.

Figuro 7-22 : Analysis	Molecules	Continuous functions	Predefined parame	eters	Shift and S	tretch	Gaps	Outputs	1/5	5
Windows Properties	Cross	Sections and spectrum	Shift fit	Stret	ch fit	Scaling	g fit			_
	Spect	rum		1st o	rder	None				
Shift and Stretch	03221	;03241		None	•	None				
Page 1/5	Bro			None	•	None				
									•	١

Selecting fitting When you have selected the appropriate symbols, you must specify which parameters to fit for this selection in columns of page 1/5. Stretch and scaling parameters can be fitted using a polynomial expression of order 1 or 2. Select option None in columns Stretch fit and Scaling fit if you don't want to fit these parameters. Uncheck the button in the Shift fit column if you don't want to fit the shift.

Note that when you disable the fit of one of these parameters, the associated initial value defined in page 3/5, is still applied to the selected cross section or spectrum.

Windows Properties         Shift and Stretch Page 2/5         Output selection         Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties         Shift and Stretch Page 3/5	Figure 7-23 : Analysis	Molecules Continuous functions	Predefined para	meters Shift an	d Stretch Gaps	Outputs	2/5
Shift and Stretch Page 2/5       Section         Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Caps Outputs 3/5         Shift and Stretch Page 3/5       Continuous functions Predefined parameters Shift and Stretch Caps Outputs 3/5	Windows Properties	Cross Sections and spectrum	n Shistore	St store	Sc store	Err store	
Shift and Stretch Page 2/5       03221;03241 Bro         Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties         Shift and Stretch Page 3/5		Spectrum					
Page 2/5       Bro         Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs         Shift and Stretch Page 3/5       Shift and Stretch Output Shift and Stretch Gaps Outputs         Bro       0.03       0.00       0.00	Shift and Stretch	o3221;O3241		П	Г	Г	
Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5         Figure 7-24 : Analysis Shift and Stretch Page 3/5       Image: Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5         Figure 7-24 : Analysis Bhift and Stretch Page 3/5       Image: Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5         Figure 7-24 : Output Stretch Page 3/5       Image: Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5         Figure 7-24 : Analysis Windows Properties       Shift and Stretch Gaps Outputs       3/5         Shift and Stretch Page 3/5       Image: Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5         Bro       0.03       0.00       0.00       0.00       0.00       0.00	Page 2/5	Bro	Г	П	Г	Г	
Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5         Figure 7-24 : Analysis Windows Properties       Shift and Stretch Page 3/5       Notecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs       3/5							
Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375		l					
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Output selectionColumns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.Figure 7-24 : Analysis Windows Properties Shift and Stretch Page 3/5Molecules Continuous functions Output Shint (nm) St init (nm) St init (2) Sc init Sc init (2) Sc init (2) Sc init Sc init (2)							
Output selection       Columns of this page are activated only if the Analysis button is checked in the ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Shift and Stretch Page 3/5       Output Shinit (nm) St init St init (2) Sc init Sc init (2)         Shift and Stretch Page 3/5       0.00       0.00       0.00							
ASCII Results tab page of Project properties (see page 62). You can check individual results to save in the output file : shift (Sh store), stretch (St store) and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Analysis Windows Properties       Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 375         Figure 7-24 : Output file :       0.00       0.00       0.00       0.00         Shift and Stretch Page 3/5       Pro       0.17       0.00       0.00       0.00	Output selection	Columns of this page a	re activated	only if the	Analysis bu	itton is chec	ked in the
Figure 7-24 : Analysis         Windows Properties         Shift and Stretch Page 3/5             Molecules             Predefined parameters    Shift and Stretch Page 3/5               Molecules             Predefined parameters    Shift and Stretch Page 3/5              Molecules             Molecules             Molecules             Molecules             Molecules             Molecules             Molecules             Molecules             Predefined parameters             Shift and Stretch Page 3/5             Bro             Molecules             Molecules             Molecules             Molecules                  Molecules             Molecules		ASCII Results tab pa	ge of <b>Proje</b>	ct properti	i <b>es</b> (see page	e 62). You	can check
and scaling (Sc store). If the Err store column is checked, the standard deviations of fitted parameters (shift, stretch and/or scaling) are saved in the output file.         Figure 7-24 : Analysis Windows Properties         Shift and Stretch Page 3/5		individual results to sa	ve in the ou	tput file : :	shift ( <b>Sh sto</b>	re), stretch	(St store)
Figure 7-24 : Analysis       Molecules       Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs       375         Windows Properties       Cross Sections and spectrum       Sh init (nm)       St init (2)       Sc init (2)       Sc init (2)         Shift and Stretch       Page 3/5       0.00       0.00       0.00       0.00       0.00		and scaling (Sc store)	). If the l	Err store	column is c	hecked, the	standard
Figure 7-24 : Analysis       Molecules       Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs       375         Windows Properties       Cross Sections and spectrum       Sh init (nm)       St init       St init       Sc i		deviations of fitted pa	, rameters (sł	nift, stretch	and/or sca	ling) are say	ved in the
Figure 7-24 : Analysis Windows Properties Shift and Stretch Page 3/5 Note: The second stretch of the second st		output file.	X	,	,	0/	
Molecules       Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs       3/5         Windows Properties       Cross Sections and spectrum       Sh init (nm)       St init       St init       St init       Sc init		1					
Molecules       Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs       3/5         Shift and Stretch Page 3/5       Cross Sections and spectrum       Sh init (nm)       St init       St init       St init       Sc init <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th>							
Figure 7-24 : Analysis       Molecules       Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs       3/5         Windows Properties       Shift and Stretch       Gaps       Outputs       3/5         Shift and Stretch       Page 3/5       0.00       0.00       0.00       0.00       0.00       0.00       0.00         Bro       0.17       0.00       0.00       0.00       0.00       0.00       0.00		×	×				
Windows Properties         Cross Sections and spectrum         Sh init (nm)         St init         St init         Sc init	Figure 7-24 : Analysis	Molecules Continuous functions	Predefined para	meters Shift an	d Stretch Gaps	Outputs	3/5
Spectrum         0.00	Windows Properties	Cross Sections and exectrum	n Shinit (nn)	Stipit St	init (2) So init	So init (2)	
Shift and Stretch         o3221;03241         0.03         0.00         0.00         0.00         0.00           Page 3/5         Bro         0.17         0.00         0.00         0.00         0.00		Spectrum	0.00	0.00 0.0	0 0.00	0.00	
Page 3/5 Bro 0.17 0.00 0.00 0.00 0.00	Shift and Stretch	03221;03241	0.03	0.00 0.	00.00	0.00	
	Page 3/5	Bro	0.17	0.00 0.	00.00	0.00	
						-	
<u> </u>							• •

Initial values of the fitting parameters In this page, you can specify the initial value of the fitting parameters (these are used by the NLLS algorithm). Shifts are expressed in the unit selected in the **Analysis** tab page of **Projects properties** (see page 57). There is one column for each kind of parameter : shift, stretch order 1, stretch order 2, scaling order 1, scaling order 2.

Note that in case the fit of these parameters is disabled in page 1/5, non-zero initial shift values are still applied.

Figure 7-25 : Analysis	Molecule	s Continuous functions	Predefined parar	neters Shif	t and Stretch	Gaps Ou	tputs	4/5
Windows Properties	Cro	ss Sections and spectrum	Sh delta (nm)	St delta	St delta (2)	Sc delta	Sc delta (2)	
	Spe	ctrum	0.001	0.001	0.001	0.001	0.001	
Shift and Stretch	032	21;03241	0.001	0.001	0.001	0.001	0.001	
Page 4/5	Bro		0.001	0.001	0.001	0.001	0.001	
								I.

**Delta values** In the fourth page, you can specify a convergence parameter (delta) for each parameter to fit. This value is used by the NLLS algorithm, e.g. to numerically calculate partial derivatives of the fitting function.

Figure 7-26 : Analysis	Molecules Continuous functions Predefined parameters Shift and Stretch Gaps Outputs 5/5
Windows Properties	Cross Sections and spectrum         Sh min (nm)         Sh max (nm)           Spectrum         0.00         0.00
Shift and Stretch Page 5/5	03221;03241 0.00 0.00 Bro 0.00 0.00
	<u></u>
Convergence control	In case of convergence problems or for safety reasons, the range of values allowed for the shift can be limited to a specified interval.
The Gaps Page	Gaps can be defined to eliminate pixels or groups of pixels from the fitting interval.
Figure 7-27 : Analysis Windows Properties Gaps	Molecules     Continuous functions     Predefined parameters     Shift and Stretch     Gaps     Outputs       Gap     Min Value (nm)     Max Value (nm)       Gap 355.3-355.9     355.3     355.9
Insert a gap	To insert a gap, right-click the <b>Insert</b> option and fill in <b>Min Value</b> and <b>Max Value</b> columns with the gap limits expressed in nanometers. The first column is automatically updated after validation of the entry fields.
Remove a gap	To remove a gap, select it in the Gap column and right-click the Remove option.
The Outputs Pages	In these pages, you can select, for each cross section defined in the <b>Molecules pages</b> (see page 67), which analysis results to save in the ASCII output file(s). Columns are activated only if the <b>Analysis</b> button is checked in the <b>ASCII Results</b> tab page of <b>Project properties</b> (see page 62).
Figure 7-28 : Analysis Windows Properties Outputs Page 1/2	Molecules       Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs       1/2         Cross Sections       AMFs       Residual Col       Sint Col       Sint Err       Sint Fact       Image: Col       Image: Col

Slant Columns and AMFs Check buttons of Slnt Col and Slnt Err columns if you want to output respectively the calculated slant column and the standard deviation of the selected species. Buttons in AMFs column are activated only if an AMF dependency has been selected in the Molecules pages (see page 69). In the Slnt Fact column, you can specify scaling factors that will be applied on slant columns and standard deviations before safe keeping in the output file. The Residual Col column gives you the possibility to enter a value for the residual column amount of the selected species in the reference spectrum (this value is used in the calculation of the vertical column).

Figure 7-29 : Analysis	Molecules Continuou	is functions	Predefin	ed parameter	rs   Shift and Stretch	Gaps	Outputs	2/2
Windows Properties	Cross Sections	Vrt Col	Vrt Err	Vrt Fact				
Outputs	Bro	N		1.				
Page 2/2	04							
: ugo 2/2	ring							
	03241							_
	J							
								••

Vertical Columns Buttons in Vrt Col and Vrt Err columns are activated only if an AMF dependency has been selected in the Molecules pages (see page 69). In Vrt Fact column, you can specify scaling factors to apply on vertical columns and standard deviations.

### 7.7 Configuration of the Wavelength Calibration procedure

The wavelength calibration facility developed in WinDOAS is based on a NLLS fitting procedure where the shift between the spectrum to calibrate and a highly accurate solar reference spectrum is determined on a series of equally spaced short intervals. The procedure also allows characterising the instrumental slit function, through fitting of user-defined Slit Function Parameters (SFP). The wavelength calibration can be applied to any kind of spectrum (the reference spectra or spectra to analyse).

The wavelength calibration uses the same property sheet as the one described in **Analysis windows properties.** You can access this property sheet by clicking the **Fit parameters** button on the **Calibration** page of **Project properties** (see page 58). Refer to previous section (7.6) for a detailed description of options, page per page. This section describes aspects particular to the parameterisation of the calibration.

The Molecules pages The Molecules pages allow introducing correcting absorbers that may optimise the accuracy of the wavelength calibration, e.g. the Ring effect (when calibrating a scattered-light spectrum) and O<sub>3</sub> absorption can be taken into account in the fit. Note that these pages are lightened compared to the equivalent pages in the **Analysis Windows properties** dialog box.

### Figure 7-30 : Analysis Windows Properties

Calibration Molecules page 1/2

This configuration assumes the presence of O<sub>3</sub> and ring structures in the fit

ns Predefined parameters Shift and Stretch Gaps Outputs  1/2 hog Fit display Interp/Convol  tital XS  Interpolate	dows pro	perties : Ca	libration d	lescription (H	larestua project]	)		3
1/2 hog Fit display Interp/Convol tial XS I Interpolate Interpolate	Continuou	us functions	Predefined	d parameters	Shift and Stretch	Gaps	Outputs	
hog Fit display Interpi/Convol tial XS I Interpolate Interpolate Interpolate								1/2
ntial XS Interpolate Interpolate Interpolate	Sections	Diff/Orthog		Fit display	Interp/Convol			
ntial XS 🛛 Interpolate	1	Differential	XS		Interpolate			
		Differential	XS	$\overline{\mathbf{v}}$	Interpolate			
ده.								
<u>*</u> >								
								<u>.</u>
Cance		dows proj	dows properties : Ca Continuous functions Sections Diff/Orthog Differential Differential	dows properties : Calibration of Continuous functions Predefine Sections Diff/Orthog Differential XS Differential XS	dows properties : Calibration description (F         Continuous functions       Predefined parameters         Sections       Diff/Orthog       Fit display         Differential XS       Image: Cancel         Cancel       Help	dows properties : Calibration description (Harestua project       Continuous functions     Predefined parameters       Sections     Diff/Orthog       Diff/Orthog     Fit display       Interp/Convol     Differential XS       Differential XS     Interpolate       Differential XS     Interpolate	dows properties : Calibration description (Harestua project)         Continuous functions       Predefined parameters       Shift and Stretch       Gaps         Sections       Diff/Orthog       Fit display       Interp/Convol         Differential XS       Interpolate         Differential XS       Interpolate         Differential XS       Interpolate	dows properties : Calibration description (Harestua project)         Continuous functions       Predefined parameters       Shift and Stretch       Gaps       Outputs         Sections       Diff/Orthog       Fit display       Interp/Convol         Differential XS       Interpolate         Differential XS       Interpolate

To generate cross section (column **Diff/Orthog**), see page 69. The **Fit display** column is activated only if the **Fits** button is checked in the **Display** frame of the **Calibration** page of **Project properties** (see page 59). The selected cross section is usually **interpolated** on the grid of the control (reference) spectrum. If the **Interp/Convol** action is set to **Convolute Std**, the selected cross section is convoluted in real time either with the fitted slit function if **Fit SFP** button is checked in the **Calibration** page of **Project properties** (see page 59) or the slit function defined in the **Slit Function** page of **Project properties** (see page 62). Standard and I0 convolutions are supported; ring cross sections can be generated in real-time using option **Convolute Ring**.

### Figure 7-31 : Analysis Windows Properties

#### Calibration Molecules page 2/2

In this configuration, the O<sub>3</sub> column density is fixed at 2e19 mol/cm<sup>2</sup>

Analysis win	dows prop	perties : Ca	libration	description (I	Harestua pi	oject)			? ×
Molecules	Continuou	is functions	Predefine	d parameters	Shift and St	retch Gaps	Outputs	200	_
Cross 03241	Sections	Co Fit	Cc init 2e19	Cc Delta	Cc 10				
ring			-0.15	0.001					
								<u>.</u>	
<u></u> к		<u>C</u> ancel		Help					

In order to optimise the accuracy of the calibration (and given limitations inherent to the method), it is recommended to carry out two passes when a cross section is introduced in the fit :

- 1. in the first pass, the column density is fitted (button in the **Cc Fit** column is checked) over all the calibration interval; a mean value of the concentration is then determined from the sub-intervals where the fitting of the cross section has a sense (where the spectral information is largest);
- 2. in the second pass, the concentration is fixed at the mean value determined at the previous pass.

See section 7.6 for a complete description of the columns in this page.

The Continuous Functions Page In this page, you define the polynomial fitting the continuous component of the spectrum to calibrate. To generate differential cross sections by the orthogonalisation method, don't forget to build an orthogonal base by right-clicking the **Properties** option from this page (see page 70).

# The Predefined Parameters Page

In the parameterisation of the wavelength calibration, you can fit offset and slit function parameters (SFP).

### Figure 7-32 : Analysis Windows Properties

A

Calibration Predefined parameters

Parameterisation for an error function line shape

naly:	sis windows prope	rties : Cali	ibration desc	cription (Harest	Ja project)		? ×
Mol	ecules Continuous	functions	Predefined pa	rameters Shift a	nd Stretch G	aps   Output	s
							1/2
	Predefined param.	Fit	Val init	Val delta	Fit store	Err store	▲
	SFP 1		0.3	0.001			
	SFP 2		0.55	0.001			
	SFP 3	$\checkmark$	0.5	0.001			
	SFP 4	$\checkmark$	0.5	0.001			
	l						
ſ	ōĸ	<u>C</u> ancel		leip			
			·				

There are 4 predefined items for SFP : SFP 1, SFP 2, SFP 3 and SFP 4. According to the line shape selected in the **Calibration** page of **Project properties** (see page 59), these parameters are used or ignored :

	SFP 1	SFP 2	SFP 3	SFP 4
Gaussian	FWHM	Ignored	ignored	Ignored
Error function	FWHM	Boxcar width	ignored	Ignored
2n-Lorentz	FWHM	Ignored	ignored	Ignored
Voigt profile	FWHM <sub>L</sub> (G)	$G/L_L$	FWHM <sub>R</sub> (G)	G/L <sub>R</sub>

**SFP 1** is generally used for fitting the FWHM of the selected line shape. The Voigt profile function is the convolution of a Gaussian and a Lorentzian line shapes. It can be used for fitting asymmetric line shapes by fitting differently on the left and the right sides, the FWHM of the Gaussian (resp.,  $FWHM_L(G)$  and  $FWHM_R(G)$ ) and the Gaussian/Lorentzian ratio (resp.  $G/L_L$  and  $G/L_R$ ). If there are several SFP to fit, it is generally recommended to fix some of them to an estimated value in order to avoid numerical instabilities.

# The Shift and Stretch Pages

For the configuration of these pages, refer to the previous section (7.6, page 72). In order to avoid the interpolation of the spectrum to calibrate, it is recommended to fit the shift of the high-resolution solar spectrum, represented by the symbol **Ref**.

In this case, all cross sections defined in the **Molecules** page have to be shifted with the solar spectrum.

Eigure 7 22 - Analysia	Analysis win	dows properties : Ca	libration descript	ion (Harestua	a project)		? ×
Windows Properties	Molecules	Continuous functions	Predefined parame	ters Shift and	d Stretch Gaps	Outputs	
Calibration							1/5
Shift and Stretch Page	Cross	Sections and spectrum	Shift fit	Stretch fit	Scaling fit		
onni and otreten i age	0324	1;Ref;ring		None	None		
The O <sub>3</sub> and ring cross sections defined in the <b>Molecules</b> page are shifted with the solar spectrum.						-	
							••
	<u>o</u> k	Cancel	Help				

- The Gaps Page This page is not active for the wavelength calibration procedure in the current version of the programme.
- The Outputs Page If the calibration is applied on all spectra (**Run Calibration** option in **Analysis Windows Properties**), this page allows checking individual fitting results (slant columns and standard deviations).

### 8 The WinDOAS Tools

This chapter covers the main features supported by WinDOAS about convolution and describes the three off-line tools included in the software :

- the Convolution/Filtering tool;
- > the **Ring** tool which allows creating Ring effect cross sections;
- the Undersampling tool for creating undersampling cross sections.

### 8.1 Supported line shapes

The three tools propose to select the type of the convolution slit function in the following list :

- 1. File;
- 2. Gaussian;
- 3. 2n-Lorentz;
- 4. Voigt;
- 5. Error function;
- 6. Gaussian, wavelength dependent;
- 7. 2n-Lorentz, wavelength dependent;
- 8. Error function, wavelength dependent.

In short, WinDOAS :

- accepts any user-defined slit function provided in an ASCII file (choice 1).
- supports the four basic line shapes described in section 4.6 page 27 (choices 2 to 5);
- can take into account the wavelength dependency (except for the Voigt profile function) of the resolution of the slit function (choices 6 to 8). In this case, a file expressing the wavelength dependency of the resolution is requested; it can have been previously produced by the wavelength calibration and slit function characterisation procedure;

The parameterisation of the slit function dialog boxes depends on the selected slit function type.

### 8.2 About Files Formats

Input files used in the configuration of WinDOAS tools are ASCII files. The supported file formats are summarised in Annex 9.1.

Output files are ASCII files including the wavelength grid and the convoluted/filtered spectrum. As these files are generally intended to be used for spectra analysis, it is preferable to foreseen for cross sections files a file name in the format imposed by WinDOAS as described in section **Configuration of the Fitting Parameters**, page 67.

WinDOAS inserts in the output file a header describing how the convoluted/filtered spectra have been created. To disable header writing, check the **Remove header** button in tools dialog boxes.

Results of the **Convolution** and the **Ring** tools are displayed in the **Graphs** window.

## 8.3 The Convolution/Filtering Tool

The **Convolution/Filtering** tool gives the possibility :

- $\blacktriangleright$  to apply a convolution with  $I_0$  correction;
- > to introduce a shift by shifting the calibration grid before convolution;
- to create an effective slit function taking into account the (finite) resolution of the source spectrum (using a FT deconvolution method):
- to filter the convoluted cross section before saving it.

Convolution/Filtering tool options are distributed in three pages :

generalgeneral options (convolution type, input/output files...)slit functionselection and parameterisation of the slit function;filteringselection and parameterisation of the filter;

Figure 8-1 :	Convolution/Filtering tool
Convolution/Filtering Tool	General Silt function Filtering
The General Tab Page	Convolution type     I0 correction ▼       Conversion     Vac to Air ▼         Shift     0.03       nm       Conc       5e+19
	Input file C:\Data\XS-HR\03\03241_vac
	Output file\data\harestua\o3_vac2air.xs_i0
	Calibration\data\harestua\Uvcal.clb
	Solar spectrum\data\harestua\kur01.ktz
	Qk Cancel Help
The Convolution Type	Three options are proposed for convolution :
	None no convolution; the original cross section is jus interpolated on the final grid;

Standard	usual convolution; convolution integral is calculated using
	method of trapezes.

 $I_{\theta}$  correction standard convolution with  $I_{\theta}$  correction;

**Conversion** The program gives the possibility to convert before convolution, the original wavelength calibration of the input cross section file from the air to the vacuum or inversely, from the vacuum to the air.

Requested Files Names	Input file the name of the high resolution cross section file to interpolate or convolute;						
	Output file	tput file the name of the resulting interpolated or convoluted cross section file;					
	Calibration	the wavelength grid on which the original cross section (input file) must be interpolated or convoluted;					
Shifting the Convoluted Cross Section	You can specify in field <b>Shift</b> , the shift in nm to apply to the convoluted cross section. In order to avoid interpolation after convolution, this shift is applied on the calibration grid before convolution.						
$I_0$ correction	The fields <b>Solar sp</b> resolution solar spec constituent. Both a formula of $I_0$ correct correction is not used	<b>rectrum</b> and <b>Conc</b> are respectively the name of the high ctrum file and the scaling column density of the concerned are used for calculating the synthetic optical density in the tion convolution (see page 29). These fields are hidden if $I_0$ d.					
Figure 8-2 : Convolution/Filtering Tool	Convolution/Filtering tool	<b>?</b> ×					

The S	Slit F	unc	tion	Tal	b
			P	ade	е

Slit function type Gaussian FWHM	e Gaussian 🔽 1 0.165 nm
Deconvolution	
Slit function type	File
Slit function file	

This page is dedicated to the selection and the parameterisation of the convolution slit function. Refer to section 8.1 for a list of possible choices. .

**Deconvolution** If the selected convolution slit function doesn't imply a wavelength dependency of the resolution, a deconvolution slit function can also be defined. In this case, the high resolution cross section is convoluted using an effective slit function obtained from the FT of convolution and deconvolution slit functions.

If you don't want to specify a deconvolution slit function, select option **File** as **Deconvolution slit function type** and let the **Deconvolution slit function** field empty.

Options in this page are ignored if the convolution type **None** is selected in the **General** tab page of the current tool. According to the selected convolution and deconvolution slit functions types, WinDOAS presents different fields to fill in.

Figure 8-3 : Convolution/Filtering Tool	Convolution/Filtering tool General Slit function Filtering	? ×
The Filtering Tab Page	Low Pass Filtering       Filter Type     Kaiser       Cutoff freq.     0.50       Pass band     0.30       Number of iterations     1	
	High Pass Filtering	
	Filter width 5 Pixels (should be an odd number)	
	Number of iterations 300 © Substract C Divide	
	Remove header	
	Ok Cancel Help	

The presentation of this tab page is the same as the **Filtering** one of **Project properties** (see page 57). The parameterisation is different according to the selected filter type. The filter type **None** disables the filtering of the convoluted cross section.

Low-pass and high-pass filtering are supported. In the latter case, the user can choose between dividing the original spectrum by the smoothed one or subtracting the smoothed spectrum from the original one.

# 8.4 The Ring Tool

Figure 8-4 : The Ping Tool	Ring tool		
Figure 8-4. The King Tool	Ring type		
	O Solar 💿 Molecular		
	Files		
	Output file (1st order) f:\gome\xs\ringxs	1.95	
	Output file (2nd order)		
	Solar spectrum e:\data\xs-hr\kurucz\Kur05.ktz		
	Calibration f:\gome\xs\Calib.clb		
	Cross section e:\data\xs-hr\no2\ulbWo2_cb.air		
	_ Sit Function		
	Slit Function type Gaussian		
	Gaussian FVVHM 0.5 nm		
	Remove header		
	Qk <u>C</u> ancel <u>H</u> elp		

Requested data	The <b>Ring tool</b> dialog box is simpler than the <b>Convolution/Filtering</b> one. Requested data are :				
	Ring type	selection of the type of ring cross section to calcular (solar or molecular);			
	Output file	the name of the calculated ring cross section file; the second order is optional;			
	Solar spectrum	the high resolution solar spectrum;			
	Calibration	the calibration on which the ring cross section has to be calculated;			
	Cross section	this field is requested for calculating molecular ring cross sections;			
	Slit Function type	the convolution slit function to select in the list 8.1; parameters to fill in depend on the selected slit function type.			
About the algorithm	The ring cross section is calculated as the ratio of the rotational Raman spectrum by the solar spectrum $(R/S)$ (see section 4.6, page 30).				
	The algorithm is described in ref. [5]. Because of a double convolution, this algorithm can be very slow, depending on the resolution of the solar spectrum and the grid on which the cross-section is calculated. A resolution of 0.05 nm for the solar spectrum is a good compromise.				
	The possibilities to ca sections have been im	lculate the second order ( $R2/S2$ ) and molecular ring cross plemented.			
The output file	The output file contain	ns four columns :			
	<ol> <li>the input waveler</li> <li>the calculated ring</li> <li>the interpolated F</li> <li>the convoluted so</li> </ol>	igth calibration; g cross section; Raman spectrum; olar spectrum.			
	When this file is used the two first columns requested. In this case	as cross section for spectra analysis, WinDOAS loads only except if the <b>Convolution Ring</b> (see section 7.6) action is e, the program uses the Raman and the solar spectra			

# 8.5 The Undersampling Tool

Figure 9.5 · The	Undersampling tool	
Undersampling Tool	Files	
	Output file (Phase 1)	xs\Usamp1xs
	Output file (Phase 2) f:\gome	xsVJsamp2xs
	Solar spectrum	xs-hr%urucz%ur01.vac
	Calibration f:\gome	xs\Calib.clb
	Slit Function	
	Slit Function type Gaussian	۰ <u>۲</u>
	Gaussian FWHM 0.165	nm
	Analysis	
	Analysis	
	Analysis method Optical d	ensity 💌
	Shift (nm) 0.008	
	Remove header	
	Qk Cancel	Help

The problem of undersampling (ex : for GOME spectra) is detailed in ref. [4]

This tool gives the possibility to pre-calculate undersampling cross sections using specified slit function and calibration. These cross sections can be introduced in the analysis parameterisation by selecting the method **File** in the **Projects Properties** (see page 60).

These cross sections can also be created with the present tool and introduced in the analysis parameterisation as cross section files.

Requested data The Undersampling tool dialog box requests the following data :

Output files	output files names for both phases 1 and 2 of undersampling;
Solar spectrum	the high resolution solar spectrum;
Calibration	
Slit Function type	
Analysis method	

Shift

# 9 Annexes

# 9.1 Files format

Calibration	File name	no restriction;		
	File extension	<i>clb</i> by default, but it is not restrictive;		
Tools	Format	<b>Column 1</b> the wavelength calibration		
	Example	ANYTHING	G.CLB	
Cross Sections Analysis windows properties	File name	ror cross sections implied in the definition of analysis windows, the file name must start by the user-defined relevant symbol followed by an underscore; this restriction does not apply to input cross sections in WinDOAS tools.		
TOOIS	File extension	<b>xs*</b> by defau	lt, but it is not restrictive;	
	Format	Column 1 Column 2	the wavelength calibration the cross section	
	Example	BrO_W228.5	ζS	
Solar Spectrum	File name	no restriction	;	
	File extension	<i>ktz</i> by defaul	t, but it is not restrictive;	
Projects properties Tools	Format	Column 1 Column 2	the wavelength calibration the solar spectrum	
	Example	KUR_01.KT	Z	
Deference en estrum	File name	no rostriction		
Reference spectrum		no restriction;		
Analysis windows	File extension	<i>rer</i> by default	, but it is not restrictive;	
properties	Format	Column 1 Column 2	the wavelength calibration the reference spectrum	
	Example	80302191.REF		

AMF Analysis windows	File name	The name of AMF file associated to cross sections implied in the definition of analysis windows must start by the user-defined relevant symbol followed by an underscore:			
properties	File extension	<i>AMF_SZA</i> for SZA dependent AMF; <i>AMF_CLI</i> for climatology dependent AMF; <i>AMF_WVE</i> for wavelength dependent AMF;			
	AMF_SZA Format	Column 1 Column 2	SZA AMF		
	AMF_CLI Format	0 (JE SZA AM	D-1)/365 IF		
		Climatology of matrix whose number grid.	dependent AMF have to be provided in a first column is SZA grid and first line, day		
	AMF WVE	Wavelength d	ependent AMF file should contain 2 parts :		
	Format	<ul> <li>the SZA dependent parameters : SZA A<sub>0</sub> A<sub>1</sub> A<sub>n</sub> X<sub>1</sub> X<sub>n</sub> C</li> </ul>			
		<ul> <li>▶ the wavelength calibration and a cross section column for each parameter X<sub>i</sub>.</li> <li>The formula used for calculating wavelength dependen AMF is :</li> <li>AMF(λ)=((A<sub>0</sub>(SZA<sub>pe</sub>)-A<sub>0</sub>(SZA<sub>re</sub>))+(A<sub>1</sub>(SZA<sub>pe</sub>)-A<sub>1</sub>(SZA<sub>re</sub>))*λ++(A<sub>n</sub>(SZA<sub>pe</sub>)-A<sub>1</sub>(SZA<sub>re</sub>))*λ++(A<sub>n</sub>(SZA<sub>pe</sub>)-A<sub>n</sub>(SZA<sub>re</sub>))*λ+(X<sub>0</sub>(SZA<sub>pe</sub>)-A<sub>n</sub>(SZA<sub>re</sub>))*X<sub>1</sub>+(X<sub>N</sub>(SZA<sub>pe</sub>)-X<sub>0</sub>(SZA<sub>re</sub>))*X<sub>2</sub>+(X<sub>N</sub>(SZA<sub>pe</sub>)-X<sub>N</sub>(SZA<sub>re</sub>))*X<sub>N</sub>)*C(SZA<sub>re</sub>)-X<sub>N</sub>(SZA<sub>re</sub>))*X<sub>N</sub>*C(SZA<sub>re</sub>)/C(SZA)</li> </ul>			
		In <b>Waveleng</b> SZA of the re and C is ignor	<b>th 2</b> option, the formula is the same but the efference spectrum is not taken into account red.		
	Example	BrO_W228.A	MF_SZA		
Slit functions	File name	no restriction	;		
	File extension	<i>slf</i> by default	, but it is not restrictive;		
Analysis windows properties	Format	Column 1 Column 2	the wavelength calibration the line shape		
		For waveleng column is the	th dependent slit function types, the second parameter dependent on the wavelength;		

slit.slf

Example

### 9.2 Troubleshooting

This section describes error messages that are popup by WinDOAS. Most of them come from a bad configuration or a bad WinDOAS objects handling. If you can not solve a problem or if you have detected a bug, it's important to contact authors. The WinDOAS.LOG log file produced by WinDOAS is an important source of information for us. A small application with a detailed description of the trouble and the sequence of manipulations leading to it can help us to solve the problem quickly.

### Tree items handling

#### 4001 Impossible to insert '%s' item due to a system limit

You have consumed all WinDOAS resources allocated for the kind of object you wanted to insert.

4007 Item '%s' already exist !

You have tried to insert in the selected tree, an item that was already existing.

### Dialog boxes settings

4004/5018/5037/

Field is out of %d..%d range

The pointed variable is out of the specified range.

- 4005 Field '%s' is empty or invalid
- 4010 Polynomial degree of '%s' should be a positive integer and a multiple of 2
- 4011 Filter width should be an odd strictly positive number
- 4012 Filter order should be an even strictly positive number
- 4013 %s should be strictly positive
- 4015 Format filter is invalid
- 4018 The number of points used for computing a polynomial should be strictly larger than the requested degree for the polynomial

The indicated field must be filled in or contains an invalid value.

### **Directory structures handling**

### 4016 You should validate path changes before selecting files

You can not manipulate files in a directory structure before inserting the path in the **Projects** tree; use the OK pushbutton to validate the path selection.

4017 Selection of files and files only is requested

In **Paths Properties** dialog boxes, right-click shortcut menus are enabled for files items only.

### **Files handling**

<b>5007</b>	File 9	∕₀s n	ot f	found
-------------	--------	-------	------	-------

5013 Can't open file %s in writing mode

WinDOAS can't open the indicated file; verify the path file.

5010 File %s is empty or not large enough

The indicated file is empty or doesn't contain the expected data.

### Analysis error messages

5014 No automatic reference selection can be perform on this type of file

The selected format doesn't support automatic reference selection mode (see page 43) because the SZA information is not available.

- 5019 Division by 0
- 5020 Exp Overflow
- 5021 Pow Overflow
- 5022 Log error
- 5023 Sqrt argument error
- 5028 Invalid window for calibration (%g,%g)
- 5038 Deconvolution slit function width must be lower than convolution one
- 5039 Slit function should be gaussian or integrated gaussians type for resolution adjustment
- 5040 Slit function should be gaussian or integrated gaussians type and temperature dependent for resolution adjustment
- 5041 Can't apply fitting of slit function parameters with calibration and resolution correction
- 5042 Error with erf function (%s)
- 5043 Matrix inversion failed
- 5046 Incompatible options for %s
- 5047 Number of degree of freedom <= 0 : reduce the number of parameters or increase the width of the analysis window or reduce the width of filter
- 5049 Voigt function failed (x=%g,y=%g)

These errors occur generally in case of bad configuration or bad input. Contact authors if you don't find the origin of the error.

5024 Bad arguments

This error is generally produced after an internal error. Contact authors.

#### 5025 No convergence after %d iterations

#### 5026 Ill conditioned matrix - too small singular value found %g

The algorithm doesn't converge. This error can occur occasionally when evaluating a bad spectrum. If it is produced systematically, it's generally because of a bad configuration or bad input (verify the content of your cross sections files and the initial values of parameters to fit).

### 5032 Zenith angle %.2f not found in file %s; AMF not computed

Interpolation error on SZA occurs when calculating AMF.

- 5034 Bad wavelength scale in file %s
- 5035 Calibration of %s is not the same as reference one
- 5036 Interpolation error because of non increasing abscisses

The wavelength calibration of the indicated file is not the expected one. Verify the size and the content of the wavelength calibration file. The file mustn't contain blank lines.

5048 The slit function is sampled at higher resolution than the original cross section. The cross section will be interpolated on the slit function grid. Continue anyway?

For convolution, the slit function is expected to be interpolated on the grid of the cross section except if the resolution of the grid of the cross section is poorer than the one of the slit function (in this case, the cross section must be interpolated).

### Allocation error messages

- 5005 Allocation error (%s,%ld,%d)
- 5006 Free memory error (%s,%08x)

Internal problem in the allocation/release of some WinDOAS objects; contact authors.