

Hiden EGAsoft Data Acquisition software for TGA-MS Applications

Auto Sequencing Software

etup	A	Sequence			
BAR Scans		N2 02 C02		MID	
1-20 PIC	BAR	N2 O2		MID	
1-506	BAR	CO2		MID	
1-506_BAR_fast	BAR	1-506		BAR	
1-506_BAR_slow	BAR				
1-50	BAR				
1-50_PIC	BAR				
📒 MID Scans					
agas 2gas	MID				
	MID				
- Air	MID				
- CO2	MID				
	MID				
N2_fast	MID				
<mark></mark>	MID				
N2_normal	MID				
N2_02_C02	MID				
N2_02	MID				
N2_slow	MID				
water_H2_28_PIC	MID				
water_PIC	MID				
🛁 Sequences					
					_
setup: N2_02_C02			l l	A 6	
cription:			[(ave

Configure a sequence of optimised scans for multiple stage analysis. Scans separated by start/stop signal

Simple Stop/Start Command Setup

置 Engineering Setti	ngs	—	×	
Connection Environment Scan Speeds External Inputs File locations	Trigger Input 101 Viggering Vacuum			Auto Start/Stop setup configured by simple switch control

Configure Multiple Bar Scans

BAR Scan Setup										_		×
	Start Mass	Stop Mass	Detector	Autorange High	Autorange Low	Start Range	Electron Energy	Emission Current	Settle Time		Dwell Time	
Enabled	1	50	Faraday 🗸	-5 🜩	-10 🜩	-5 🔹	70	400	Normal	~	Normal	~
Subscan 2	1	50	Faraday 🗸	-5 🔹	-10 🜩	-5 🔹	70	400	Normal	~	Normal	~
Subscan 3	1	50	Faraday 🗸	-5	-10 🖨	-5 🔹	70	400	Normal	~	Normal	~
				Save	Loa	ad						

Bar scan setup – run up to 3 bar scans at a time.

The mass regions of interest are scanned, the parts of the mass range of no

interest are ignored, reducing the data acquisition time.

Mass spectrometer parameters can be optimised for each part of the scan.

MID Setup

miD Scan Setup											- 🗆 X
Save 📔	Load	Clear]								Advanced View Scan Background
								I	LED Filter	Off	Background 🗸
Nitrogen	\sim	28	14 29	15	0	0	0	0	0		0
		1000	72 8	0.7	0	0	0	0	0		
Oxygen	\checkmark	1000	114 4	1							0
		40	20 36	38	0	0		0	0		
Argon	\checkmark	1000	107 3	1	0	0	0	0	0	0	0
Carbon dioxide	~	44	28 16	12	45	22	46	29	13	0	0
		1000	114 85	60	13	12	4				
I	\checkmark		0 0	0							0
		0	0 0	0	0	0	0	0	0	0	
	V	0	0 0	0	0	0	0	0	0	0	U
	\checkmark	0	0 0	0	0	0					0
			0 0								
	\checkmark		0 0	0	0	0	0	0	0	0	0
		0	0 0	0	0	0	0	0	0	0	

MID scan setup Automatic removal of spectral overlaps

3D Bar Scan Viewer



Real time 3D data viewer for easy determination of bar scan trends



MID View

Simple MID setup - display up to 16 trends at a time

Bar View



Quadrupole Mass Spectrometers for Advanced Science

Data Analysis Software



Peak integration/fitting routines

Data Export Facility



- Export to correct format for import to TGA software
- Export to Excel
- Export to NIST MS library: Two methods included:
 - 1. Complete spectra from selected BAR scan
 - 2. Selected peaks from MID view

NIST export from a selected BAR scan



Quadrupole Mass Spectrometers for Advanced Science

NIST import from a selected BAR scan



When importing a Bar scan the whole mass spectrum is used in the NIST search routine

MID View - export to NIST



Simple MID setup - display up to 16 trends at a time

NIST with imported selected MID peaks



Importing MID peaks allows the user to choose groups of peaks to narrow the NIST search