

Analyzing & Testing

TGA-GC-MS Coupling

Evolved Gas Analysis with Chromatographic Pre-Separation



Leading Thermal Analysis

Thermogravimetry and Evolved Gas Analysis

Thermogravimetry (TGA) and Simultaneous Thermal Analysis (STA)

Thermogravimetry (TGA) and Simultaneous Thermal Analysis (STA), which is primarily simultaneous TGA-DSC (Differential Scanning Calorimetry), are widely applied analytical methods for the research and guality control of all kinds of inorganic and organic materials and products. Often, it is not sufficient to determine only the mass and enthalpy changes resulting from a thermal treatment; additional information may be required about the volatile products evolved during chemical reactions, thermally induced transitions, evaporation, and decomposition in order to gain deeper insight into the nature, structure and composition of the materials. NETZSCH thermobalances and STA systems are arranged vertically with the sample above the balance, allowing the gases to flow in the natural upward direction. This design is ideal for coupling gas analysis equipment to the top of the furnace.

Mass spectrometry (MS) is the most comprehensive method for gas analysis, providing sensitivity down to the ppb/ ppm level, high resolution at the atomic level, and also high speed of detection. Although the direct coupling of quadrupole mass spectrometers (QMS) - e.g., Aëolos® or SKIMMER to a TG or STA - is a well proven method, it has limitations in determining the individual components of gas mixtures which often hinder clear identification and interpretation. This is where gas chromatography comes into play.

TG information

- Dehydration
- Desolvation
- Binder burn-out
- Decomposition
- Pyrolysis
- Combustion
- Oxidation
- Corrosion
- Evaporation
- Compositional analysis





Gas Chromatography (GC) – Mass Spectrometry (MS)

GC is a high-resolution method for separating volatile and semi-volatile compounds. The gas mixtures are separated based on the differences in component distribution between a stationary phase (e.g., inner coating of a capillary) and a mobile phase (purge gas; e.g., helium, nitrogen, argon, hydrogen).

Gas components with low affinity for the stationary phase but higher affinity for the mobile phase will be rapidly carried away by the purge gas, whereas gases with a high affinity for the stationary phase will follow with a relatively significant time delay ("retention time").

MS is applied as a detection system at the outlet of the GC separation column and will register the time distribution of the separated gas components in the purge gas flow. Because of this pre-separation of the gases by the GC, and the sensitivity and resolution of the MS, detailed structural information is provided, allowing most compounds to be exactly identified. **Mobile Phase**

The mobile phase in GC is a gas, therefore all analytes must be evaporated and introduced into the column in gaseous form.

Stationary Phase

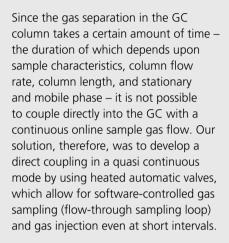
In current technology, the stationary phase in the GC is usually a polymeric coating inside a long and narrow fused silica capillary.

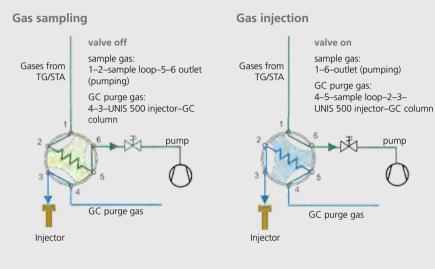
MS information

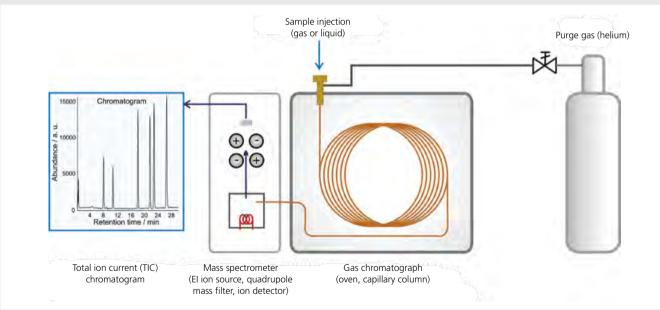
- Gas detection
- Gas identification
- Solid-gas reaction
- Decomposition products
- Pyrolysis gases
- Combustion products
- Flue gas identification

Coupling MS-GC-STA 449 **F1** Jupiter®

TGA/STA-GC-MS Coupling



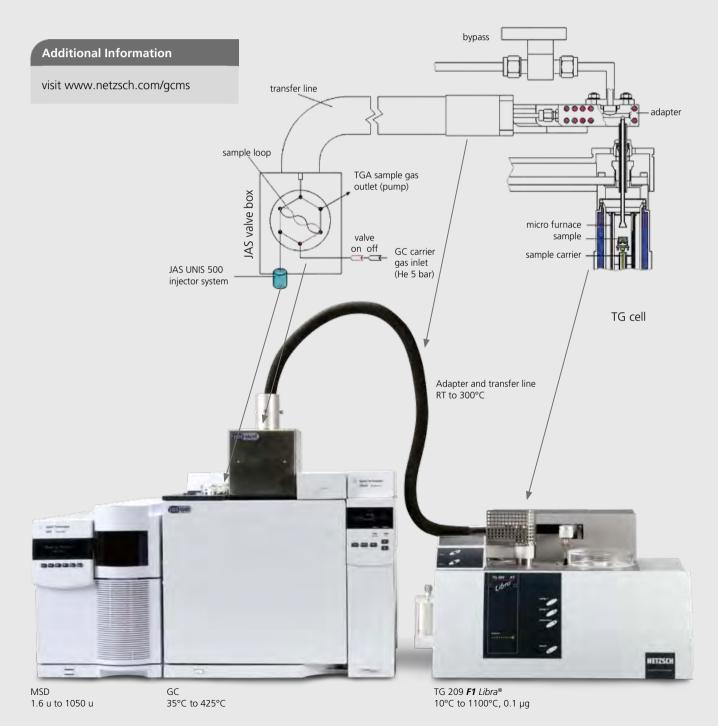




The injected gas mixture of the sampling loop (or parts of it in the split mode) passes the separation column of the GC at the set flow condition (e.g., constant flow of the purge gas 1.5 to 2 ml/min) and arrives, after a final pressure reduction step, as a molecular beam at the ion source of the quadrupole mass spectrometer. The separation column is either held at constant high temperature (for the continuous injection mode), or temperature-programmed from a low start temperature to subsequent high temperature (for the event-controlled mode). The mass spectrometer detects all volatile products as a function of time. The curve of the total ion current (TIC) over time (retention time) is called a chromatogram. Peaks appear in the chromatogram resulting from the sample gas interaction in the stationary and mobile phases, and from the boiling temperature of the individual components. These retention peaks are analyzed by the MS software to determine the contributing substances.



Detailed Diagram of the Functional Elements of the TGA-GC-MS Coupling



Event-Controlled Mode, Continuous Mode

TG /%

100

80

60

40

20

0

.....

290.8 °C

1.3 %/min

290.6 C

376 5 10

NR/SBR, 6.89 mg, 20 K/min, N2, event-control with 3 thresholds

-10.63 %

380 1 C

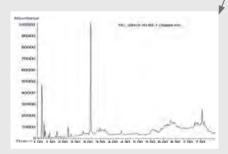
40 Time /min

450.7 °C 6 %/min

50

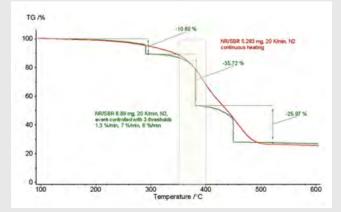
Event-Controlled Mode

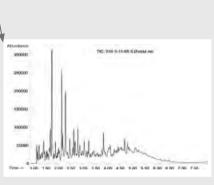
The event-controlled mode uses the automatic detection of mass changes (DTG thresholds) as the trigger for starting the GC-MS. This is called the "reaction-rate-dependent trigger" and operates via the automatic run of the "sequence" function in the ChemStation software. Overlapping reactions can be resolved and the corresponding GC furnace program allows for a nearly complete separation of the evolved gases. Individual thresholds can be selected in every segment of the temperature program.



Automatic detection of the TGA step for additives with high volatility: identification of vulcanization agents, plasticizers, or oils in the rubber sample.







Second step of rubber decomposition with styrene

Event-Controlled Run

Temp. /*C DTG /(%/min)

2 700

0 600

-2

.4

-6

-12 100

-14 0

-25.79 %

70

449.7 °C

500

400

-8 300

-10 200

- Nearly complete identification of evolved gases with exact temperature correlation to TGA and DTG
- Integration of chromatogram
- Library search report → quantification



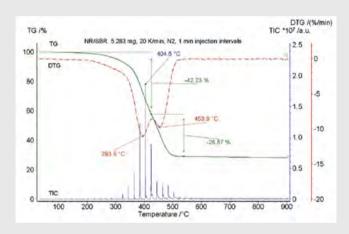
Continuous Mode

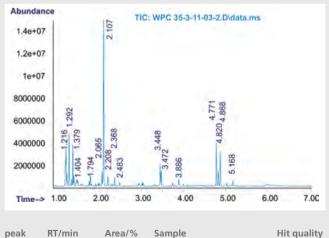
The TG and GC-MS are started simultaneously. Then, when running in continuous mode, the GC-MS operates with a fixed column temperature (e.g. 250°C) and short injection intervals. The evolved gases are identified in good temperature and time correlation with the TGA and DTG results. The total ion chromatogram can be displayed, evaluated, stored and exported together with the thermal analysis curves. A trigger for the start of the GC-MS run can also be programmed in any temperature segment.

- Identification of ions/molecules
- Extraction of ions (shows temperaturetime profile, SIM mode)
- Verification of temperature-time correlation with TGA-DTG curves
- Fast and easy method for routine control of gas composition and comparison of samples

Single Run (or Stand-Alone GC-MS Run)

- Triggered or manual start of GC program
- Complete identification of evolved gases
- Integration of chromatogram
- Library search report → quantification (see table, extract)





can		711007 70	Sample	inc q
	1.216	10.14	Propene	91
	1.292	9.40	Pentane	91
	1.379	6.06	1-Pentene, 2-methyl-	94
	1.404	1.18	2-Pentene, 4-methyl-	91
	1.794	1.82	Heptane, 4-methyl-	94
	2.065	2.38	2-Hexene, 4,4,5-trimethyl-	78
	2.107	32.19	1-Heptene, 2,4-dimethyl-	95

Software: Easy Handling, Reliable Data Evaluations

Programming of Measurements

The start of the combined TGA-GC-MS experiment is programmed in the *Proteus*[®] software together with the usual inputs for the thermal analysis measurement parameters.

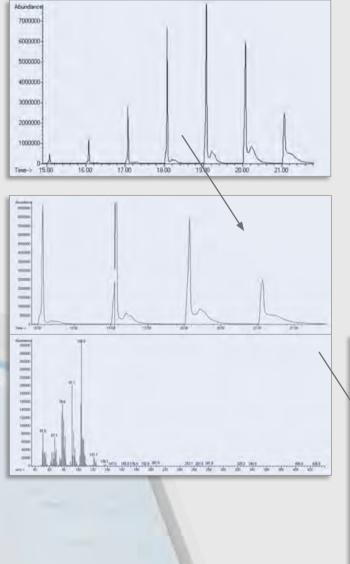
Measureme		Identity:	event-con	trol
 O Correct 		Sample name:	WPC	
() Sample	the meter	Sample mass:	8.030	mg
	+ correction	Crucible mass:	0	mg
C sample	+ conscion	MECO		
		rec.g.	neo.	1
Laboratory:	TG-GC-MS		Device	Value
Project:	Test	Purge	1 MFC	OXYGEN
Operator:	Kal	Purge	2 MEC	HELIUM
Qate:	27.04.2011 10:32:34	Prote	ctive MFC	HELIUM
Mgterial:	composite 🛩	3	-	
			Ch	ange gases
Temperature calbra	dich:			
O will not be used				
() will be used (se				
	eustcal5\TG 209 F1\TG 209 F1-20 K_min.ng	N.		
	antenotia cost tha cost i con Diminik			
emark:				
telium 30:40 Al2O3	crucible witht lid, adaptor 300°C + transfer	Ine heating 280°C		

Selection of the Trigger Mode

- Trigger (for continuous mode)
- Reaction-rate-dependent trigger (for event-controlled mode), with delay time for the duration of the experiment

tode			
O Trigger			
Reaction rate depe	ndent trigger		
Delay: 15	<160> min		
erivative value preser	ntabion		
in percent			1.3
absolute			
arameters of TG signa	al derivative Derivative Range /mg/	min Eilter Factor	
	500	75	10
2000	50	150	
2000	50 5	150 400 1000	10



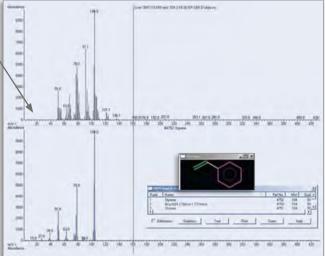


Evaluation of the GC-MS-TIC Chromatograms

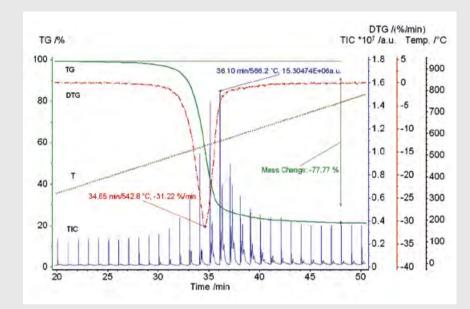
This example shows NR/SBR rubber measured in the continuous mode with 1-min injection intervals and 250°C column temperature.

The MS spectra below every retention peak can be viewed by double-clicking (example spectrum at 19.054 min, 403.5°C).

The next double-click displays the hit list of the integrated MS library search program, including the identity, molar mass, structure, and stored spectrum of the chemical compound found. Here, styrene is identified at 19.054 min (403.5°C) with the high hit quality of 91%.

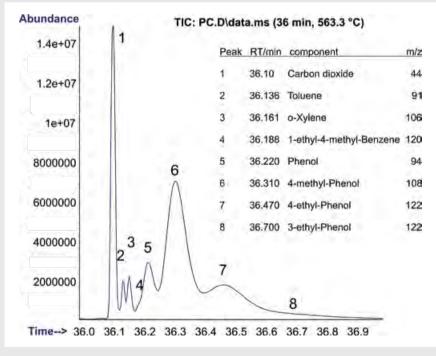


Pyrolysis of Polymers



Polycarbonate (PC)

The pyrolysis of 12.42 mg of PC (Lexan 144R) in helium leads to one-step decomposition with the DTG peak at 543°C. The TIC chromatogram of 1-min injection intervals at a column temperature of 250°C (split 10:1) indicates different stages in gas evolution, with the highest intensity at 566°C.



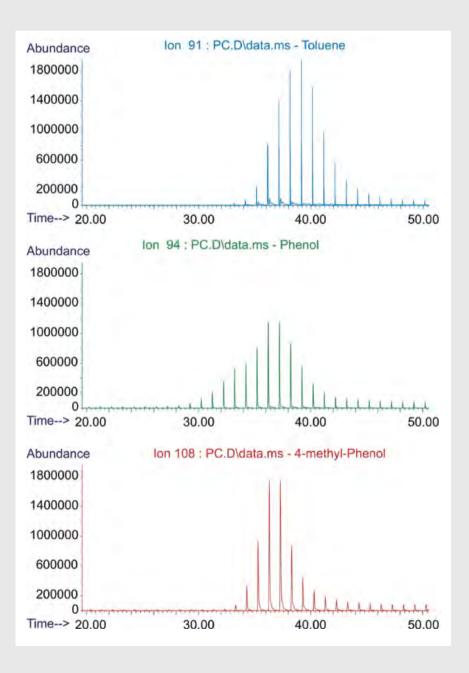
The analysis of the underlying MS spectra of the TIC, beginning at 36 min, leads to the identification of 8 main components with the NIST library search.



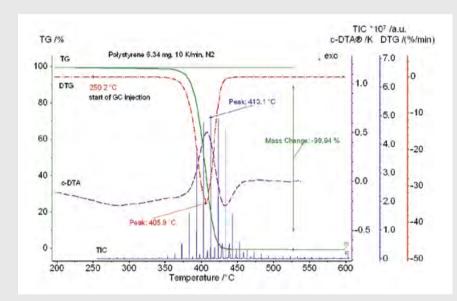
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Extracting individual molecule-ions from the total ion chromatogram allows for the monitoring of the temperature and time dependence of gas evolution of individual ions (SIM mode). Clearly visible is the later evolution of toluene compared with phenol and 4-methyl-phenol.





Pyrolysis of Polymers



Polystyrene (PS)

The NIST SRM 705a decomposes in a nitrogen flow in one step at 406°C (DTG peak) and the TIC chromatogram (maximum 413°C) indicates at least two ranges of gas evolution.

1e+07	1	6		TIC: 06	PS.D\data.ms (17 min, 42	2.6 °C)
16+07			Pea	k RT/min	component	m/z
9000000		2	1	17.041	Toluene	91
8000000			2	17.058	Styrene	104
7000000			3	17.098	1-phenyl-1,3-Butadiene	130
7000000			4	17.161	1-methylethenyl-Benzene	118
6000000			5	17.230	dipheny-Methane 8	168
5000000			6	17.293	1,1-diphenyl-Ethylene	180
			7	17.447	1.1'(1.3-propanediyl)bis-Benzene	196
4000000			8	17.636	4-Phenyl-1-butyne	130
3000000	1		9	17.773	1,2-diphenyl-Cyclopentane	220
2000000			10	17.842	1,1'-(1-butene-1,4 diyl)bis-(Z)-Benz	ene 208
1000000		3	4	5 6	7	9 10

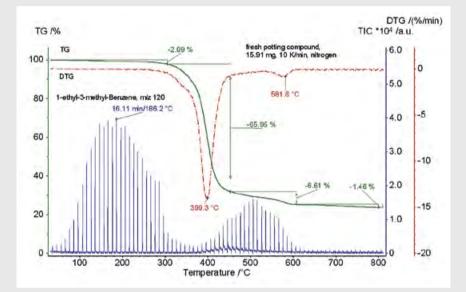
The evaluation of the chromatogram at 17 min (423°C) reveals styrene (m/z 104) as the dominant product (of the whole decomposition), and the styrene dimer (m/z 208) as a component of the second decomposition stage.

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Evaporation of Solvents

Potting Compound

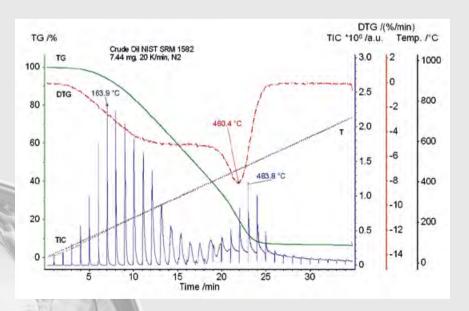
This potting compound for the encapsulation of electronic components, declared to be solvent free, evolves a benzene compound right from the start of the experiment. The selected ion m/z 120 for 1-ethyl-3-methyl-benzene shows the maximum intensity at 186°C, in a range where the TGA curve displays no significant mass loss. The gases evolved during the decomposition of the potting compound were not part of our focus for this plot.



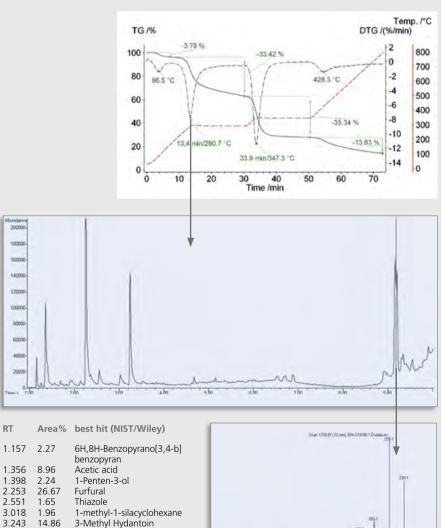
Fuels

Crude Oil Pyrolysis

The continuous heating shows the ranges of distillation (characterized by a nearly constant mass loss rate of 5%/min up to 400°C) and pyrolysis of the hydrocarbons (up to 550°C).



Biomass



Oak Wood Pyrolysis

4.370 mg of ground oak wood (quercus robur) was pyrolyzed in nitrogen in the event-controlled mode. The DTG threshold of 8%/min resulted in two TGA steps with isotherms at 291°C and 347°C. The gases automatically injected at the GC at these temperatures were separated with the GC furnace program: 60°C for 0.5 min, heating to 310°C at 25 K/min.

The TIC chromatograms for the two events (pyrolysis steps) with peaks in the full range of retention times between 1 min and 10 min are easily evaluated by the library search report. Based on the selected integration parameters, the report shows the major peaks with retention time, percentage of peak area, best hit for the identified molecule, and hit quality (not shown in table). Any individual peak analysis can be carried out manually by a few mouse clicks, e.g., identifying substituted phenanthrene at 9.209 min retention time in the chromatogram for the gas evolved at 291°C (see bottom figure).



6.881

9.156

9.209

9.298

1.70

16.97

9.01

1.42

Benzenamine, N-methyl-N-phenyl-2-lsopropyl-10methylphenanthrene

Phenanthrene,

1-methyl-7-(1-methylethyl)-Benzene, 1-methyl-4-[(4-pro pylphenyl)ethynyl]-



Rush Grass (Juncus) Pyrolysis

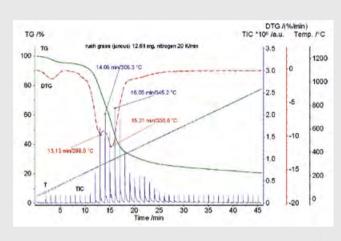
After drying (4.6% mass loss), the rush grass decomposes in a nitrogen atmosphere in several steps which are not well separated. The gases evolved represent the cellulose and lignin content of the sample.

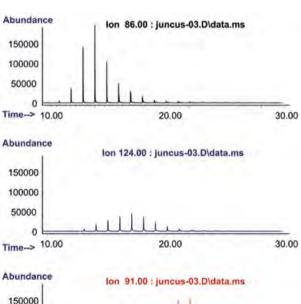
The selected fragment ions (figure right) show significant differences in their temperature distributions (the displayed time range of 10 min to 30 min corresponds to the temperature range 226°C to 626°C):

■ m/z 86 for digitoxose (from cellulose)

- m/z 124 for guaiacol (from lignin)
- m/z 91 for toluene (from lignin)





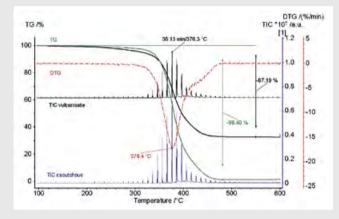


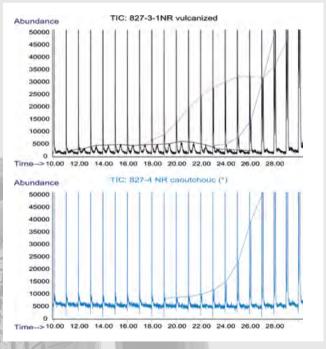
20.00

100000 50000 0 Time--> 10.00 LINS

16

Rubber Pyrolysis





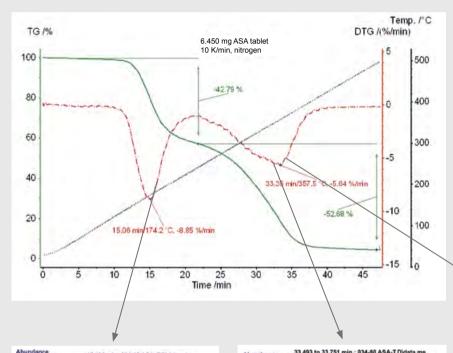
Differentiation of NR and Vulcanized NR

The difference between the pure NR (natural rubber) caoutchouc and the vulcanized NR is shown by the TIC chromatograms, mainly in the range from 132°C to 294°C (27 min), where TG indicates 0.3%/min difference in the mass loss rate.

Only the vulcanized NR shows the additives benzothiazole (12 - 24 min) and xylene (start at 17 min, 193°C). Significantly, the decomposition of the isoprene (representative for NR) starts in both sample tests at 23 min (254°C).



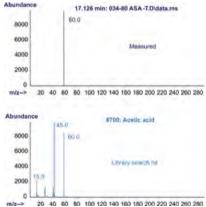
Pharmaceuticals



Acetylsalicylic Acid (Trade Name: e.g. Aspirin®)

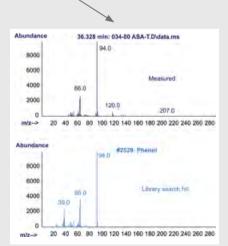
The evaporation and decomposition of the powdered acetylsalicylic acid tablet sample in nitrogen yields the main products acetic acid (1st TG step) and acetylsalicylic acid and phenol (2nd TG step).

The GC-MS coupling to the sensitive TG is also an ideal instrument combination for the analysis of residual solvents and impurities in pharmaceutical drugs and excipients.



Abundance				
9000	92	0 120.0		
7000	1		Measure	be
6000 5000				
4000	65.0	138.0		
3000	1.00	152.0	207.0	
2000	46.0		1 6.4	281
1000	L L BB.	· 11 11 "	9,0	. 1
m/z-> 20 4	40 60 80 1	00.120 140 160 1	180 200 220 24	10 260 28
	10 60 80 1			
Abundance	10 60 80 1		Acetylsalicylic	
	10 60 80 1	#43668: A		
Abundance 9000 8000 7000		#43668: A		
Abundance 9000 8000 7000 6000 3	3,0	#43668: A 1i 135/0		acid
Abundance 9000 8000 7000 6000 4 5000		#43668: A 1i 135/0	Acetylsalicylic	acid
Abundance 9000 8000 7000 6000 3	3,0	#43668: A 1i 135/0	Acetylsalicylic	acid
Abundance 9000 8000 7000 6000 4 5000 4000	3,0	#43668: A 1: 135.0	Acetylsalicylic	acid

20 40 60 80 100 120 140 160 180 200 220 248 260 280



Expertise in Service



Our Expertise – Service

All over the world, the name NETZSCH stands for comprehensive support and expert, reliable service, before and after sale. Our qualified personnel from the technical service and application departments are always available for consultation.

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- Installation and commissioning
- Hotline service
- Preventive maintenance
- Calibration service
- IQ /OQ/PQ
- On-site repairs with emergency service for NETZSCH components
- Moving/exchange service
- Technical information service
- Spare parts assistance

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Our Expertise – Applications Laboratories

The NETZSCH Thermal Analysis Applications Laboratories are a proficient partner for nearly any thermal analysis issue. Our involvement in your projects begins with proper sample preparation and continues through meticulous examination and interpretation of the measurement results. Our diverse methods and over 30 different state-of-the-art measuring stations will provide ready-made solutions for all your thermal needs.

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The three Business Units – Analyzing & Testing, Grinding & Dispersing and Pumps & Systems – provide tailored solutions for highest-level needs. Over 2,500 employees at 130 sales and production centers in 23 countries across the globe guarantee that expert service is never far from our customers.

When it comes to Thermal Analysis, Adiabatic Reaction Calorimetry and the determination of Thermophysical Properties, NETZSCH has it covered. Our 50 years of applications experience, broad state-of-the-art product line and comprehensive service offerings ensure that our solutions will not only meet your every requirement but also exceed your every expectation.

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