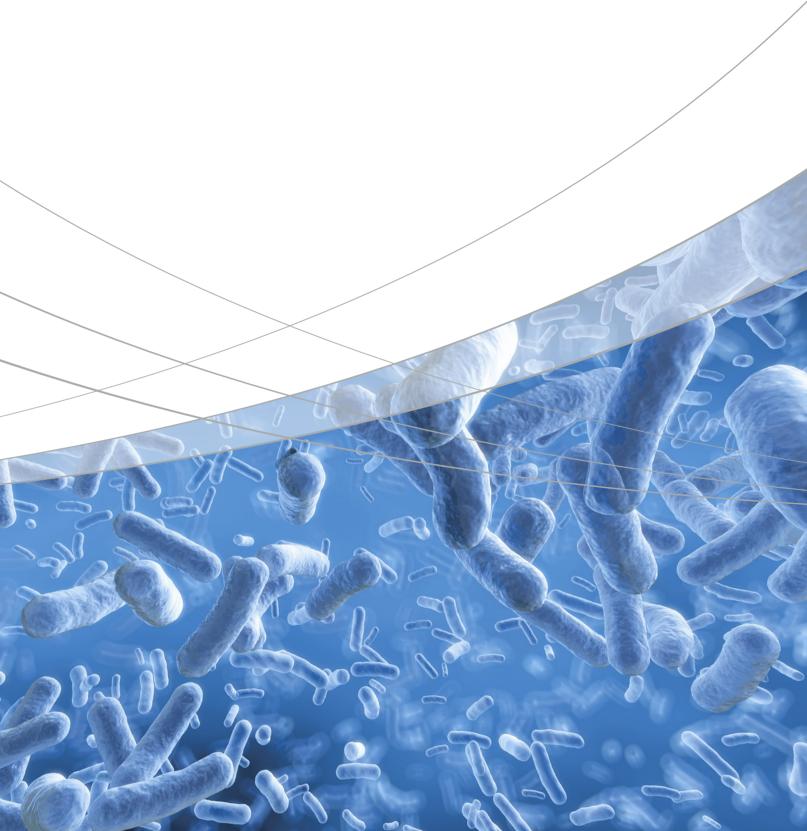
Solutions for





Infectious Diseases

Infectious diseases are disorders caused by harmful microorganisms (pathogens) such as viruses, bacteria, fungi or parasites. The number of cases for these in the U.S. is led by bacteria with viruses a distant second, with the number of bacterial infectious diseases is on the rise in the last ten years (Figure 1). Given this rise, it is of paramount importance that these infections are investigated and examined. This solution guide introduces a variety of analytical instruments to evaluate such tasks.

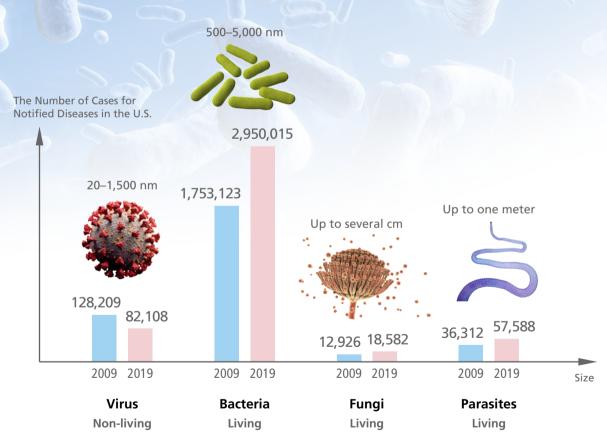


Fig. 1: Four major types of microorganisms

This solution guide introduces three analytical instruments: Liquid Chromatography–Mass Spectrometer (LC-MS), Matrix Assisted Laser Desorption/Ionization Time-of-Flight Mass Spectrometer (MALDI-TOF MS), and Gas Chromatography-Mass Spectrometer (GC-MS). A targeted population will be larger at the avoidance phase compared to the testing and care phases. Click on an image to read a specific application.

Target base



Whole population

Those with symptoms



Your work place

Chemical Manufacturing

Hospital Academia

Hospital

LC, LC-MS



MALDI-TOF MS



GC, GC-MS







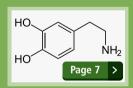
Vaccine active ingredients



Alcohol purity testing

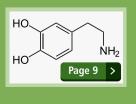


Testing





Biomarker search



Care



Ion-Pair Reversed-Phase LCMS-9030 Mass Spectrometer for Separation and Identification of Oligonucleotides



Issue

Shorter oligonucleotides (less than 20 bases) can be easily resolved by HPLC, but the separation of those with longer sequences becomes progressively more challenging.

Solution

Q-TOF LC-MS can be employed to capitalize on its high mass accuracy.

System Description

System Configuration

HPLC System: Nexera

Column: Shim-pack Scepter HD-C18-80

 $(2.1 \text{ mm I.D.} \times 100 \text{ mm}, 1.9 \mu\text{m})$

LC-MS System: LCMS-9030 lonization Mode: ESI (Negative)

Sample Information

DNA oligonucleotide standard solution

Benefits

- High resolution is achieved for the separation of 10 to 60 base oligonucleotides.
- Highly accurate masses are obtained for oligonucleotides.

Conclusion

10–60 base oligonucleotides were separated and identified.

Representative Data

Table 1: The most abundant mass accuracy of oligonucleotides on LCMS-9030

Standards	M.W. (Da)	Cal. <i>m/z</i>	Meas. m/z	ppm	Adduction
10-mer	3043	1013.1763	1013.1773	0.99	[M-3H] ³⁻
15-mer	4634	925.7547	925.7557	1.08	[M-5H] ⁵⁻
20-mer	6117	763.5005	763.5020	1.96	[M-8H] ⁸⁻
25-mer	7642	954.1569	954.1585	1.68	[M-8H] ⁸⁻
30-mer	9191	834.4997	834.5014	2.04	[M-11H] ¹¹⁻
40-mer	12274	817.1994	817.2013	2.33	[M-15H] ¹⁵⁻
50-mer	15379	853.3599	853.3615	1.87	[M-18H] ¹⁸⁻
60-mer	18493	1026.3331	1026.3349	1.75	[M-18H] ¹⁸⁻

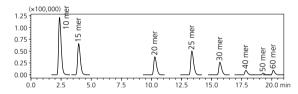


Fig. 2: Separation of 10 to 60 base oligonucleotides

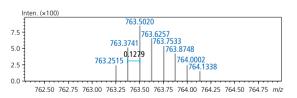


Fig. 3: High-resolution mass spectrum of 20-base oligonucleotide



Quadrupole Time-of-Flight Liquid Chromatograph Mass Spectrometer

LCMS-9030

The LCMS-9030 quadrupole time-of-flight (Q-TOF) mass spectrometer is a powerful instrument that integrates the world's fastest and most sensitive quadrupole technology with TOF capabilities for accurate mass measurement.



Product

Quality Control of Synthetic Peptides Using the MALDI-8030 Dual Polarity Benchtop MALDI-TOF Mass Spectrometer



Issue

Quality Control (QC) is critical in providing high-purity products.

Solution

Negative ionization mode can be used in QC to preserve the integrity of species with labile functional groups.

System Description

System Configuration

MALDI System: MALDI-8030

Matrix Solution: alphacyano-4-hydroxycinnamic acid

(CHCA), 5 mg/mL in 1:1 acetonitrile / water, with/without acid (0.1% TFA) depending on the instrument polarity.

Sample Information

Synthetic peptides provided by a client

Benefits OOO

- Peptides with labile functional groups can be analyzed and the detection of the intact species will be viable.
- Mass spectra become cleaner as salt adducts won't be detected.

Conclusion

Cleaner mass spectra were obtained and QC was conducted.

Representative Data

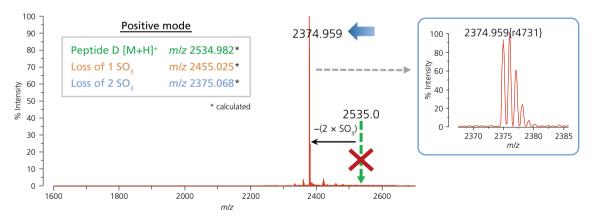
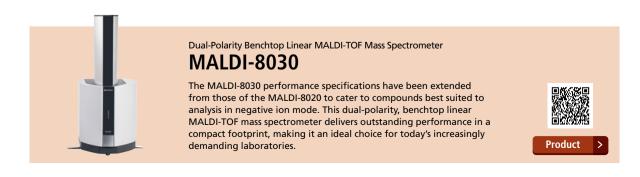


Fig. 4: Positive mode MALDI spectrum of Peptide D: only the species corresponding to the loss in both sulfo groups was detected.



Determination of Alcohol Content in Hand Sanitizers by Headspace GC-FID



Issue

Each country has a minimal purity for ethanol and isopropanol used in hand sanitizers. Sanitizers can contain colorants and thickening agents that can contaminate the GC, however, and render results inaccurate.

Solution

Headspace GC-FID can be employed instead of a liquid injection system.

System Description

System Configuration

GD-FID: Nexis GC-2030AF

Headspace Autosampler: HS-20

Column: SH-Rxi-624Sil MS (0.32 mm I.D.x 30 m, 1.8 µm)

Sample Information

Commercially available hand sanitizers

Benefits

- Headspace will keep your instrument clean.
- The longer uptime means a higher throughput.
- The cleaner instrument means stable data.

Conclusion

Hand sanitizers were analyzed with good repeatability and linearity according to EPA 5021.

Representative Data

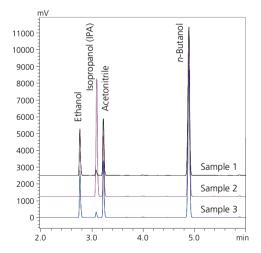


Fig. 5: Overlaid chromatograms of the hand sanitizer samples

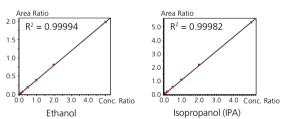


Fig. 6: Calibration curves of ethanol and IPA at 0.1 – 5 % (v/v)

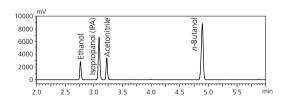


Fig. 7: Separation of the alcohols from the acetonitrile (IS) both at 2 % (v/v) with a SH-Rxi-624Sil MS column



Gas Chromatograph with Headspace Sampler

Nexis GC-2030 with HS-20 NX series

The HS-20, developed as the best solution for volatile component analysis, has been improved and introduced as the NX series. The excellent basic performance and user-friendly design provide a powerful solution for scientists in both research and quality control laboratories.



Product >

Protein Biomarker Quantitation from Human Blood and Plasma Using Novel Collection Technology by LC/MS/MS



Issue

It takes time and expertise to analyze a wide range of peptides at once.

Solution

A dedicated system and its accompanying kits (PeptiQuant Comprehensive kit and Biomarker Assessment kit) can be used to analyze a wide range of peptides at once.

System Description

System Configuration

LC-MS System: LCMS-8060

Column: Phenomenex Aeris C18

(2.1 mm I.D. × 150 mm, 1.7 µm)

Sample Information

Dried plasma spots and dried blood spots

Benefits OOO

Matrix effects can be reduced by using dried plasma spots as a sample compared to dried blood spots.

Conclusion

27 peptides were monitored from dried blood/plasma spots.

Representative Data

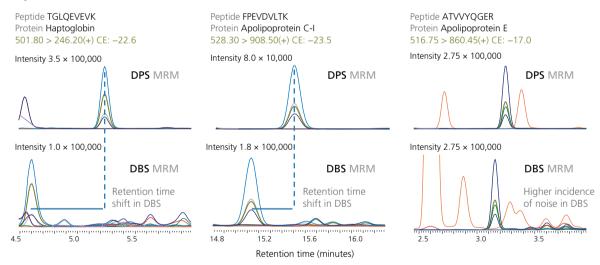


Fig. 8: MRM chromatograms for three peptides; haptoglobin, apolipoprotein C-I and apolipoprotein E.

The upper MRM chromatograms correspond to DPS extracts; the lower MRM chromatograms from
DBS extracts highlight retention time shifts and a higher incidence of noise in the MRM channel.



Application of MALDI-TOF MS for Discrimination of Species Processing **Highly Conserved Ribosomal RNA Gene Sequences**



Issue

Identification can be difficult due to a high homology rate of 16S rRNA gene sequences.

Solution

A dedicated system and its accompanying database can be used to identify microorganisms.

System Description

System Configuration

MALDI System: **AXIMA Microorganism Identification**

System

Sample Information

Harvested microorganisms

Benefits



A dedicated database allows you to compare an unknown mass spectrum to a reference for a more confident identification.

Conclusion

E. Coli species were identified.

Representative Data

Table 2: Agreement rate of rRNA gene sequences (From Muroi et al. (2011) with permission from Pharmaceutical Society of Japan)

Species	Strains	No.	1	2	3	4	5	6	7	8	9	10	11
E. coli	NBRC 12734	1											
	NBRC 3972	2	99.1										
	NBRC 12713	3	99.7	99.3									
	NBRC 13168	4	99.4	99.5	99.3								
	NBRC 13891	5	99.9	99.3	99.8	99.5							
	NBRC 13893	6	99.7	99.3	100.0	99.3	99.8						
	NBRC 3301	7	99.7	99.0	99.7	99.2	99.5	99.7					
	NBRC 102203	8	99.5	99.0	99.5	99.2	99.7	99.5	99.3				
	NBRC 14237	9	99.7	99.2	99.7	99.3	99.8	99.7	99.4	99.6			
E. fergusonii	NBRC 102419	10	99.6	98.9	99.5	99.3	99.6	99.5	99.7	99.3	99.5		
E. hermanii	NBRC 105704	11	97.4	97.4	97.3	97.5	97.5	97.3	97.1	97.2	97.4	97.1	
E. blattae	NBRC 105725	12	95.9	96.0	96.0	96.0	96.0	96.0	95.8	95.8	95.9	96.0	95.8



MALDI-TOF MS System

AXIMA Microorganism Identification System

The AXIMA Microorganism Identification System presents a new solution for the identification of microorganisms in the pharmaceutical, food, science, energy, environment, clinical and health fields.



Detection of Tuberculosis Biomarkers by Multidimensional GC-MS



Issue

Tuberculosis infects one third of the world population, and biomarkers are required for a rapid diagnosis. Matrix can be complex and pose an analytical issue.

Solution

Heart-cut GC-MS was used to mitigate adverse effects from matrix interferences.

System Description

System Configuration

GC-MS System: Multi-Dimensional GC-MS System

Sample Information

Sputum

Benefits OOO

Sample matrix can be eliminated for an easier detection of biomarkers.

Conclusion

20 potential biomarker compounds were found.

Representative Data

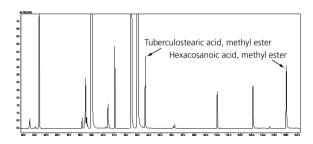


Fig. 9: The 2nd dimension GC-MS chromatogram of the sputum sample. The clean chromatogram allows accurate quantification of the 20 biomarkers.

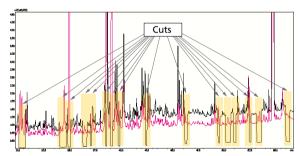


Fig. 10: The 1st dimension FID chromatogram of the sputum sample in standby mode (Pink) and in cut mode with 16 cut positions for 20 target compounds (Black).



Multi-Dimensional GC/GCMS System

MDGC/GCMS-2010 series

The complete separation and quantification of specific compounds can be difficult in samples that contain many compounds, such as petroleum products and fragrance products. Multi-Dimensional GC (MDGC) uses columns with different separation characteristics to achieve highly accurate separation of target components from such complex matrices.



Product >

Simultaneous Analysis of Remdesivir and Metabolites in Human Plasma Using Fully Automated Sample Preparation LC/MS/MS System



Issue

Manual sample extraction can be labor-intensive.

Solution

A sample preparation system can be fully automated.

System Description

System Configuration

UHPLC: Nexera X2

Column: Shim-pack Scepter C18-120

(2.1 mm l.D. × 50 mm, 1.9 μm)

LC-MS System: LCMS-8060 Ionization Mode: ESI (Positive)

Sample Information

Commercially available human plasma

Benefits

- Sample extraction can be automated.
- Automation reduces method variability, the chance of sample mix-ups and the risk of exposure to samples.

Conclusion

Automation not only allowed for walk-away times but yielded good accuracy and reproducibility.

Representative Data

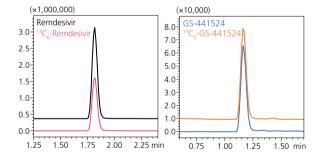


Fig. 11: MS Chromatograms of Remdesivir, [U-Ring-¹³C₆]-Remdesivir (Left) and GS-441524, [¹³C₅]-GS-441524 (Right)

Table 3: Repeatability of Remdesivir and GS-441524 in plasma

		Spiked	Intra-Assay $(n = 6)$				
Compounds	QC Sample	Conc. (ng/mL)	Average Conc. (ng/mL)	Precision %RSD	Accuracy %		
	LLOQ	100	90.5	2.0	91		
Remdesivir	Low	750	797	1.7	106		
Kerridesivii	Medium	1000	1045	0.9	105		
	High	3750	3393	2.0	91		
	LLOQ	5	4.51	3.1	90		
GS-441524	Low	37.5	33.2	2.5	89		
G3-441324	Medium	50	45.2	2.3	90		
	High	187.5	171.7	3.6	92		



Fully Automated Sample Preparation Module for LC-MS

CLAM-2040

Operational improvements in LC-MS systems have made LC/MS an advantageous technique for clinical research, thanks to its specificity, its accuracy and its capability to analyze several targets simultaneously. The CLAM-2040 is an online automated sample preparation module that brings LC-MS smoothly into your laboratory.



Product >

JP- and EP-Compliant Analysis of Impurities of COVID-19 Drug Dexamethasone



Issue

Pharmaceutical companies need to control impurities strictly from a safety perspective.

Solution

LC-MS can be used to monitor impurities.

System Description

System Configuration

HPLC System: Nexera lite

Column: Shim-pack VP-phenyl

(4.6 mm I.D. × 250 mm, 5 µm)

Detection: UV 254 nm

Sample Information

Dexamethasone standard solution

Benefits

System suitability tests are satisfied for both Japanese and European Pharmacopeia using the triple quad LCMS-8060NX.

Conclusion

Impurity testing was carried out according to Japanese and European Pharmacopeia.

Representative Data

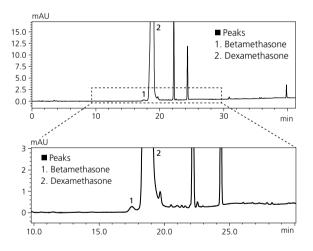


Fig. 12: Chromatogram of Reference Solution (a) (European Pharmacopoeia)

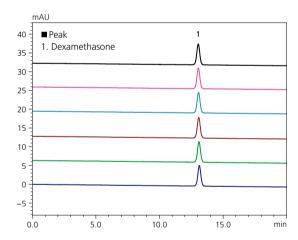


Fig. 13: Comparison of Chromatograms in 6 Repeated Tests with Standard Solution (5.94 mg/L) (Japanese Pharmacopoeia)



Ultra High Performance Liquid Chromatograph

Nexera series

The Nexera series is a family of HPLC systems that marries Al and IoT enhancements to set new industry standards. The Nexera lite is a conventional HPLC model with excellent basic functions and many options for expansion. Cutting-edge Al functions increase the reliability of established analysis workflows and adaptable to any application system.



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