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REVIEW ON HYPHENATED TECHNIQUES AND THEIR APPLICATIONS

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ABSTRACT:

Hybrid methods have gotten a lot of coverage in recent years as a primary way to solve complex analytical challenges. Combining two (or more) methods to detect and isolate chemicals from solutions is referred to as hyphenated separation techniques. Chromatography is frequently used as the other technique. In chemistry and biochemistry, hyphenated methods are commonly used. Few examples are Gas Chromatography-Mass Spectrometry (GC-MS), Gas Chromatography-Infrared Spectroscopy (GC-IR), Liquid Chromatography-Mass Spectrometry (LC-MS), Liquid Chromatography-NMR Spectroscopy (LC-NMR), Liquid Chromatography-Infrared Spectroscopy (LC-IR) and Capillary Electrophoresis-Mass Spectrometry (CE-MS) etc. The hyphenated techniques have lot of advantages and applications like Environmental monitoring, Food, refreshment, flavor and aroma examination, Forensic and criminal cases, Biological and pesticides detections. This review focuses on few hyphenated techniques and their utility in practice.

Key words: LC-MS/MS, hyphenated techniques, LC-NMR, applications, chromatography

Hirschfeld devised the word "hyphenation" to describe the simultaneous use of a separation technique and one or more spectroscopic detection techniques a couple of decades ago. The hyphenated technique was born out of the union of a separation and a spectroscopic detection technique. Chromatography separates organic molecules in a mixture into pure or nearly pure fractions. Spectroscopy – This technique generates selective data for detection using standards or library spectra. (1)

Hybrid methods have gotten a lot of coverage in recent years as a primary way to solve complex analytical challenges. Over the years, the value of merging separation technologies with spectroscopic techniques for both quantitative and qualitative analysis of unknown compounds in complex natural product extracts or fractions has been demonstrated. Combining two (or more) methods to detect and isolate chemicals from solutions is referred to as hyphenated separation techniques.Chromatography is frequently used as the other technique. In chemistry and biochemistry, hyphenated methods are commonly used. When the name of one of the methods includes a hyphen, a slash is most often used instead of a hyphen. To obtain structural information that will allow the compounds present in a crude sample to also be characterized. Liquid chromatography (LC), usually a high-performance liquid chromatography (HPLC), gas chromatography (GC), or capillary electrophoresis (CE) is linked to

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spectroscopic detection techniques, e.g., Fourier-transform infrared (FTIR), photodiode array (PDA) UV-vis absorbance or fluorescence emission, mass spectroscopy (MS), and nuclear magnetic resonance spectroscopy (NMR), resulting in the introduction of various modern hyphenated techniques.

- Gas Chromatography-Mass Spectrometry (GC-MS)
- Gas Chromatography-Infrared Spectroscopy (GC-IR)
- Liquid Chromatography-Mass Spectrometry (LC-MS)
- Liquid Chromatography-NMR Spectroscopy (LC-NMR)
- Liquid Chromatography-Infrared Spectroscopy (LC-IR)
- Capillary Electrophoresis-Mass Spectrometry (CE-MS)

Advantages of Hyphenated Techniques:

- 1. To solve difficult analytic problems.
- 2. Less time for research
- 3. A wide spectrum of automation
- 4. Increasing flow of samples
- 5. Improved repeatability
- 6. Since it is a closed device, pollution is reduced.
- 7. Improved combined selectivity, resulting in a higher level of information
- 8. Provide excellent separation efficiency as well as on-line complementary spectroscopic data acquisition on a complex mixture's LC or GC peak of interest.(2-11)

GC-MS:

GC-MS is a hyphenated technique that was created by combining GC and MS.It was the first of its kind being used for research and development. This hyphenated technique produces mass spectra that contain more structural details based on fragmentation interpretation. It is possible to combine fragment ions of different relative abundances. Compounds that are sufficiently volatile, thin, and stable at high temperatures can be easily analysed in GC-MS conditions.For GC-MS analysis, polar compounds, especially those with a large number of hydroxyl groups, may need to be derivatized. The conversion of the analyte to its trimethylsilyl derivative is the most successful derivatization technique. A sample is injected into the GC device's injection port, vaporised, removed in the GC column, analysed by the MS detector, and registered in GC-MS.GC requires the analyte to have significant vapor pressure between 30 and 300°C.

<u>Retention time-</u> refers to the time between injection and elution (tR). An injection port is located at one end of a metal column (which is mostly filled with a sand-like material to promote optimum separation) and a detector (MS) is located at the other end of the column. The sample is propelled down the column by a carrier gas (argon, helium, nitrogen, or hydrogen, to name a few) propels the sample down the column. The GC distinguishes the components of a mixture over time, while the MS detector provides information that aids in structural detection.(**1&12**).

There are two kinds of GC-MS columns: capillary columns and macrobore and packed columns. The following statements about the GC-MS interface should be carefully considered.

- 1. The interface effectively transfers effluent from the GC to the MS.
- 2. In the interface, the analyte must not condense.
- 3. Before entering the MS ion source, the analyte must not decompose.
- 4. The amount of gas entering the ion source must be within the MS's pumping power.

The most broadly utilized interfaces for a GC-MS are electron sway ionization (EI) and substance ionization (CI) modes. Notwithstanding, in current GC-MS frameworks, different sorts can be utilized that permit recognizable proof of sub-atomic particle. These days, a GCMS is incorporated with

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different on-line MS information bases for a few reference compounds with search abilities that could be helpful for spectra match for the distinguishing proof of isolated parts.(13-14)

Applications of GC-MS

Environmental monitoring

GC-MS has become an enthusiastically suggested device for observing and following natural contaminations in the climate. The expense of GCMS gear has diminished though the dependability has particularly expanded. The assurance of chloro-phenols in water and soil, polycyclic fragrant hydrocarbons (PAH), unleaded fuel, dioxins, dibenzofurans, organo-chlorine pesticides, herbicides, phenols, halogenated pesticides, sulfur in air is extremely advantageous to be screened by this method. It very well may be utilized to screen the debasement results of lignin in bio-mass exploration, pesticides in spinach. Investigation of decacyclene, ovalene and even C60 debasement examination of carbamazepine and its metabolites in treated sewagewater and steroid should be possible without derivatization.(**15,16,17**)

Food, refreshment, flavor and aroma examination

Food varieties and drinks have a few fragrant mixtures existing normally in local state or shaped while handling. GC-MS is solely utilized for the examination of esters, unsaturated fats, alcohols, aldehydes, terpenes and so forth GC-MS is additionally used to identify and gauge foreign substances, deterioration and debasement of food, oil, margarine, ghee that could be destructive and ought to be controlled and checked as directed by legislative organizations. It is utilized in the investigation of piperine spearmint oil, lavender oil, fundamental oil, scent reference norms, aromas, chiral compounds in fundamental oils, aromas, menthol, allergens, olive oil, lemon oil, peppermint oil, yiang oil, strawberry syrup, spreadfatty substances, lingering pesticides in food and wine.(**18,19**)

Forensic and criminal cases

GC-MS can examine the particles from suspect to relate his contribution on the off chance that. The investigation of fire trash utilizing GC-MS can be set up by American Society for Testing Materials (ASTM) standard for fire garbage examination. It is the key instrument utilized in sports hostile to doping research centers to test competitor's pee tests for precluded execution upgrading drugs like anabolic steroids. It is additionally usually utilized in measurable toxicology to discover harms, steroids in natural examples of suspects, casualties, or the expired.(20,21)

Biological and pesticides detections

GC-MS is completely utilized in bio-investigation of blood, pee for the presence of barbiturates, opiates, alcohols, lingering solvents, drugs like sedatives, anticonvulsant, antihistamine, against epileptic medication, narcotic hypnotics, opiates and food things. This method could be utilized for recognizing contaminations, unsaturated fat profiling in microorganisms, presence of free steroids, blood toxins, metabolites in serumorgano-chlorinated pesticides in stream water, drinking water, sodas by head space, pesticides in sunflower oil and so on.(22)

DETECTION OF CHEMICAL AGENTS AGENTS AND PROTECTION

Hazardous identification frameworks have become a piece of all United State air terminals, GC-MS. Is a fundamental piece of synthetic investigation unit. For upgrading ability in country security and general wellbeing readiness, conventional GC-MS units with the transmission quadrupole mass spectrometers, just as those with barrel shaped particle trap (CIT-MS) and toroidal particle trap (T-ITMS) mass spectrometers have been altered for field convenientce and close to continuous discovery of compound fighting specialists (CWA) like sarin, soman, and VX.(23-25)

RESEARCH IN ASTRO CHEMISTRY AND GEOCHEMISTRY

Several GC-MS have left earth for the astro chemistry studies. Two were taken to Mars planet by the Viking program. Scientist analysed the atmosphere of Venus with GC-MS. The Huygens probe of the Cassini-Huygens mission landed one GC-MS on Saturn's largest moon, Titan. Significantly enhanced molecular ions, major isomer and structurally significant mass spectral peaks, extended range of low volatility hydrocarbons that are amenable for analysis and unique isotope ratio information make GC-MS valuable for organic geochemical applications.(26,27)

Petrochemical and hydrocarbons analysis

Significantly enhanced molecular ions that are always observed, isomer and structurally significant mass spectral peaks and extended range of low volatilite hydrocarbons that are amenable for analysis including waxes up to C74H150 makes the GC-MS a most valuable technique. Broad range of petrochemicals, fuels and hydrocarbon mixtures, including gasoline, kerosene, naphthenic acids, diesel fuel, various oil types, transformer oil, biodiesel, wax and broad range of geochemical samples can be analysed by GC-MS.(28,29)

Clinical toxicology

Improved sub-atomic particles, expanded scope of mixtures managable for examination, prevalent affectability for compounds and quicker investigation are the primary alluring highlights of the clinical toxicology. The poison and toxins are recognized by GC-MS. It is widely utilized in clinical toxicology.(30)

Academic research

As a one of a kind and amazing innovation, the GC-MS gives an uncommon chance to play out the investigation of new mixtures for portrayal and ID of orchestrated or derivatized compound. It is broadly utilized in unadulterated and applied sciences like Chemistry, Polymers, Nanotechnology and Biotechnology and so forth It yields valuable data that can be utilized in research distribution globally.(**31**)

INDUSTRIAL APPLICATIONS

GC-MS is utilized in businesses for the examination of sweet-smelling solvents, inorganic gases, amino liquor in water, pollutions in styrene, glycol, diols, xylene, allergens in beautifiers and so on GC-MS is utilized for the portrayal of formic corrosive in acidic corrosive for mechanical use. In Industries acidic corrosive is significant transitional in coal synthetic blend. It is utilized in the creation of poly ethylene, cellulose acetic acid derivation and poly vinyl just as synthetic fiber and textures. By nature of its broad range of applications, GC-MS has ushered in a new era of research and elevated the impactful presentation and characterization of chemicals to new heights.GC-MS is a sophisticated technology that cannot be compared to other modern analytical tools, however it can be combined with a mass spectrophotometer to produce GC-MS/MS. It has a wide range of uses, including academic research, quality control, and industrial applications. Its simple, efficient, and automated system produces quick, repeatable, and effective outcomes that help develop Science and Technology. (32,33)

LC-IR:

The hyphenated procedure created from the coupling of a LC and the discovery strategy infrared spectrometry (IR) or FTIR is known as LC-IR or HPLC-IR. While HPLC is quite possibly the most impressive partition procedures accessible today, the IR or FTIR is a valuable spectroscopic method for the recognizable proof of natural mixtures, in light of the fact that in the mid-IR locale the constructions of natural mixtures have numerous ingestion groups that are normal for specific functionalities, e.g., - OH, - COOH, etc. Notwithstanding, blend of HPLC and IR is troublesome and the advancement in this hyphenated strategy is very lethargic on the grounds that the hyphenated method's 237 ingestion groups of the versatile stage dissolvable are so gigantic in the mid-IR district that they regularly dark the little sign created by the example segments. Moreover, as a discovery procedure, IR is substantially less touchy contrasted with different other location methods, e.g., UV and MS. One is a stream cell approach and the other is a dissolvable disposal approach. The methodology utilized with the stream cell in LC-IR is like that utilized in UV-vis and other ordinary HPLC finders. For this situation, assimilation of the portable stage prompts the obstruction of the identification of test segment ingestion groups, however some straightforward district of the mid-IR range produces discovery probability. For the most part, KBr or KCl salts are utilized for the assortment of test segments in the eluent, and warming up the medium before IR identification kills the unpredictable versatile stage solvents. There are two kinds of interfaces for the dissolvable disposal approach: diff use-reflectance infrared Fourier change (DRIFT) approach and buff er-memory strategy. A bound together interface for GC, HPLC, and SFC hyphenation to FTIR applying IR minuscule method is likewise accessible today. (34,35) In the first LC/IR systems flow cells were used in a fashion

analogous to LC with on-line UV/VIS absorption detection. In order to circumvent interfacing difficulties related to the IR absorptions of the mobile phase, in 1979 Kuehl and Griffiths. developed the first solvent elimination based LC/IR set-up in which the eluent is evaporated prior to IR detection. Since then two approaches can be discerned in LC/IR, namely, the flowcell approach and the solvent-elimination approach. In the contemporary practice of LC/IR both approaches are applied, although the detection limits and spectral information obtained with either approach may differ considerably. The principles, applications, merits and limitations of flow-cell and solvent-elimination LC/IR have been reviewed in a number of books and papers..(**36-38**)

LC-MS:

LC-MS or HPLC-MS refers to the coupling of a LC with a mass spectrometer (MS). One of the most important techniques of the last decade of the twentieth century was hybrid liquid chro matography-mass spectrometry (LC-MS). The rapid pace of growth, as well as its widespread accepta nce and adoption, It's incredible, particularly given the price tag. LC-MS has become the technique for decision for analytical support in numerous phases of medication advancement inside the drug business.(39) Willoughby and coworkers broke down the advancement of LC-MS innovation, just as its acknowledgment as an insightful technique.at present we are as yet in the beginning phase of acknowledgment and utilization of LC-MS in true applications. While new advancements in instrumentation are easing back down, huge advancement is made in the various fields. This stage puts various requests on the advancement of LC-MS as an insightful method, as is for example demonstrated by the as of now developing revenue in programming development, particularly for more effective information handling after LC-MS examination. (40)The chemical separation capacity of LC is combined with the ability of an MS to selectively detect and confirm molecular identity in an LC-MS. MS is one of the most sensitive and selective methods of molecular analysis, providing information on the analyte molecule's molecular weight as well as its fragmentation pattern. For validating the identification of analyte molecules, the information gained from MS is important. (41-43). One of the most significant issues with LC-MS is the response efficiency. The type of interface used is highly dependent on a number of factors.

Three significant troubles are met in joining the two incredible scientific procedures, LC and MS:

(I) the clear stream rate contrariness as communicated in the need to present 1 ml/min of a fluid emanating from a regular LC segment into the high vacuum of the mass spectrometer,

(ii) the dissolvable creation contradiction as consequence of the successive utilization of non-unstable versatile stage added substances in LC division advancement.

(iii) the ionization of non-unstable and additionally thermally labile analytes.(44-46).

QUANTITATION

It is for the most part hard to perform quantitative judgments utilizing outright MS reactions. This is a direct result of the huge number of elements that impact the outright MS reaction, for example, the neatness of the particle source, particle optics and the crash cell, particle concealment, particle source stream rates, crash cell pressure and a definitive MS vacuum. It is hard to control these variables and, as an outcome, outright MS reactions are dependent upon critical everyday variety. Along these lines, inward norms are typically needed to accomplish dependable and exact quantitative outcomes. Stable isotope variants of the analyte are ideal inside principles as they have practically indistinguishable substance properties however are effortlessly recognized during MS. Besides, they right for any misfortunes or shortcomings in the example planning measure and right for particle concealment. This strategy is named stable isotope weakening and is equipped for giving examines that are extremely exact and exact. Thus, stable isotope weakening LC-MS tests are regularly appropriate as reference techniques. This is done in certain tests where creating calibrators isn't straight-forward e.g., dried blood spot tests. Nonetheless, this methodology depends on the steady isotope inward standard being 100% unadulterated and having a similar molar reaction as the analyte. For precise outcomes, adjustment bends plotting analyte inside standard reaction proportion versus analyte focus are as yet required. Metabolomics is mainly concerned with the detection and quantification of small

molecule metabolites (It helps in the comprehension of biological and biochemical processes in comple x systems). (47) MS offers quantitative examination of metabolites with high affectability and selectivity and potential to recognize metabolites. For instance, the accessibility of different climatic

pressing factor ionization (API) techniques in both positive and negative modes [e.g., electrospray ionization (ESI), climatic pressing factor synthetic ionization (APCI), what's more, environmental pressing factor photoionization (APPI) empowers ionization of different classes of metabolites.(48,49)

METHOD OPTIMIZATION

The development of mass spectrometer (MS), pretreatment, and liquid chromatography (LC) separation technologies has enhanced the sensitivity and depth of proteome research, making it feasible to examine lowly expressed proteins, including those of interest. In setting up a LC-MS examine countless conditions and boundaries should be considered and improved. The genuine conditions are exceptionally subject to the idea of the analyte and the LC division, making it hard to give conventional conditions. (50-52)Each analyte requires singular improvement. Albeit distributed strategies are a significant beginning stage, test execution and ideal conditions can fluctuate extraordinarily between various instruments and test networks. Affectability is exceptionally reliant on the instrument utilized and the test conditions. Instrument producers are persistently improving the affectability of their mass spectrometers and by and large offer a scope of models with various sensitivities. It is in this manner imperative to survey if an instrument has the fundamental affectability to accomplish the ideal furthest reaches of identification.

Particle source boundaries and crash energy can be upgraded during constant implantation of a weaken arrangement of the analyte, ideally in a similar versatile stage utilized for the LC partition. (53) Decision of section particle to screen requires some idea as the most bountiful piece may not generally be the most ideal decision. On the off chance that other huge part particles happen, it is fitting to likewise assess them as they may really give cleaner chromatograms better sign than commotion. It is additionally fitting to screen a second part particle to check for potential impedances. In the event that an obstruction is available, the proportion of the two section particles is distinctive to that of the norm. utilizing the case of methylmalonate estimation in mouse tissues. Derivatization to shape butyl esters was utilized to improve the affectability. The main drawbacks of this discrete method selection strategy are that

(1) creating and validating a set of focused prep LC methods is laborious, and

(2) the results can be suboptimal for compounds with prepare RTs around the boundary of contiguous focused prep LC methods. (53-56)

Another thought is whether to utilize single MS or couple MS. Couple MS really brings about a decline in supreme sign in view of misfortunes in the crash cell and the way that the sign might be spread across a few parts when just a couple are being identified. According to the World Health Organization, 70% of the world's population has used non-conventional medications at some point in their lives, including medicinal plant therapies. The market for medicinal and aromatic plants (MAPs) is continually growing, thanks to rising consumer demand around the world. With an annual growth rate of 6.1 percent from 2017 to 2022, the global market for botanical and plant-derived pharmaceuticals is predicted to increase from USD 29.4 billion in 2017 to about USD 39.6 billion by 2022. (57)

APPLICATIONS

BIOCHEMICAL SCREENING FOR GENETIC DISORDERS

To detect metabolic disorders, blood samples from newborn. Newborns are examined using LC-MS. In newborn screening, second-tier LC-MS testing has been utilised to corroborate the results of first-tier immunoassays. The early work of Millington et al. set up the capability of evaluating neonatal dried blood spots for a wide scope of intrinsic mistakes of digestion (IEM). The advancement of ESI sped up this cycle and strategies for handling the huge number of tests needed for testing all new-borns were created. Australia took a lead here and all Australian infants are right now tried utilizing this procedure. Many professionals working in various parts of newborn screening have been captivated by the idea of universal population-based newborn screening for rare inborn metabolic abnormalities. At the same time, it continues to cast doubt on many of the newborn screening ideas that have guided the industry for the past four decades. Newborn screening (NBS) for treatable "hidden" genetic metabolic abnormalities was launched about 50 years ago to detect phenylketonuria in blood spots obtained from newborn babies using a bacterial inhibition assay.(**58**)It is viewed as a feature of acknowledged medical care in all nations in the created world. Numerous public wellbeing labs in North America added evaluating for intrinsic hypothyroidism during the 1970s to their menu of illnesses evaluated for.

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During the 1980s and 1990s new advancements in electrospray ionization also, pair mass spectrometry permitted the improvement of fast, high throughput examinations of tests separated from dried blood spots (DBS). As new strategies have been grown new difficulties have been uncovered, with respect to which sicknesses to evaluate for, how to affirm analyze rapidly and precisely and how to circle back to patients distinguished through the program. (59-62)

THERAPEUTIC DRUG MONITIRING AND TOXICOLOGY

Disappointment with the significant expense of business immunoassays utilized in helpful medication checking and their variable cross-reactivity with metabolites has prodded the advancement of LC-MS examines as choices. Remedial medication observing (TDM) is needed to improve treatment of basic portion drugs with a limited helpful reach where there is a decent possibility of either overdosage or underdosage. Checking the medication fixation can direct the medication dosage to improve helpful adequacy while limiting the side impacts. TDM has been performed for a longtime utilizing immunoassay yet it is perceived that immunoassay techniques can endure with vague obstruction from related mixtures, metabolite impedance or lattice impacts. Fluid chromatography–couple mass spectrometry (LC–MS/MS) has been in routine use in clinical research centers for barely 10 years. LC-MS, which automates sample preparation and reduces analytical run times, is likely cost-effective despite high initial investment costs. This single example exemplifies the majority of the benefits of LC-MS techniques and argues that they are suitable for everyday TDM. (63-66)

Some likely traps and mis-IDs have been noted however these can be wiped out via cautiously coordinating with maintenance times and mass spectra to a norm and utilizing qualifier particle proportions. The affectability of current instruments likewise permits the investigation of oral liquids and hair tests. (67-70)

Application of LC/ESI-MS in in forensic sciences

LC-MS is utilized for assurance of harmfulness, in drug examination and furthermore in follow investigation. By utilizing modest quantity of test, the poisons in various material can be resolved with LC-MS. Any harmful metabolites in food or drinks can be controlled by utilizing LC-MS. E.g., Identification of cleanser added into squeezed orange can be dictated by investigating by the juice and cleanser test. The standard surfactant alkyl diphenyl ether sulphonic corrosive is utilized. Both juice and cleanser tests are dissected in same chromatographic conditions. The mass chromatograms and mass spectra acquired from the juice and cleanser tests are indistinguishable with the reference spectra of standard surfactant (alkyl diphenyl ether sulphonic corrosive).Liquid chromatography/mass spectrometry (LCNS) has recently gained popularity in a variety of fields because it allows for the confirmation study of polar or non-volatile compounds without derivatization. There are a variety of effective LC-MS interfaces available. Atmospheric pressure chemical ionisation (APCI), electrospray (ESI), and thermospray are some of them (TSP)Interfaces for Frit-fast atom bombardment ionisation (Frit-FAB), Frit-fast atom bombardment ionisation (Frit-FAB), and particle beam (PB) interfaces.Applying LC-MS to forensics, the authors have successfully analyzed various illicit drugs and their metabolites, quaternary ammonium salts 5.6, benzodiazepines, as well as aqueous degradation products of pesticides.(71,72)

LC-NMR:

Among the spectroscopic methods accessible to date, NMR is likely the most un-delicate, but it gives the most helpful primary 240 Sarker and Nahar data toward the construction, clarification of normal items. Mechanical advancements have permitted the immediate equal coupling of HPLC frameworks to NMR, offering ascend to the new down to earth procedure HPLC-NMR or LC-NMR, which has been generally known for more than most recent 15 years. The first on-line HPLC-NMR explore utilizing superconducting magnets was accounted for in the mid-1980s. In any case, the utilization of this hyphenated procedure in the logical research centers began in the last piece of the 1990s as it were. (73)LC-NMR vows to be of incredible worth in the examination of complex combinations, everything being equal, especially the investigation of normal items and medication related metabolites in bio-fluids. LC-NMR tests can be acted in both consistent stream and stop-stream modes. A wide scope of bio analytical issues can be tended to utilizing 500, 600, and 800 MHz frameworks with 1H, 13C, 2H, 19F, and 31P tests. The fundamental requirements for online LC-NMR, notwithstanding the NMR and HPLC instrumentation, are the nonstop stream test and a valve introduced before the test for recording either persistent stream or halted stream NMR spectra. A MS can likewise be joined to the

framework through a splitter at the yield of the LC-NMR interface. In a large portion of the LC-NMR activities, switched stage sections are utilized, utilizing a twofold or tertiary dissolvable blend with isocratic or slope elution. (74-77) The protons of the solvents of the versatile stage cause extreme issues for getting a sufficient NMR range. The collector of the NMR spectrometer isn't exactly ready to deal with the serious dissolvable signs and the frail substance flags simultaneously. To beat this issue, dissolvable sign concealment can be accomplished by one of the three significant strategies: pre-immersion, delicate - beat numerous illumination or water concealment upgrade through T1 impacts (WET) pre-immersion utilizing a z-inclination. This problem can also be minimized by considering the following guidelines:

1. Using eluents that have as few 1H NMR resonances as possible, e.g., H2 O, ACN, or MeOH.

2. Using at least one deuterated solvent, e.g., D2O (approx.\$290/L), ACN-d3 (approx.\$1600/L), or MeOD (approx.\$3000/L).

3. Using buffers that have as few 1H NMR resonances as possible, e.g., TFA or ammonium acetate.

4. Using ion pair reagents that have as few 1H NMR resonances as possible, e.g., ionpairs with t-butyl groups create an additional resonance.

The remainder is sent to the MS through a 12 m capillary and NMR. The valveswitching interface, also known as the valveswitching interface, is a versatile alternative. A powerful alternative is the valve-switching interface termed the BNMI (BRUKER NMR MASS-SPECTROMETRY INTERFACE). This BNMI is essential in the LC–NMR–MS loop storage mode, in which a portion of the loop contents can be stored in a delay loop after transfer to the NMR. At this timeFollowing NMR acquisition, the dilutor slowly infuses analytes into the sample, into MS.Late advances in both equipment and programming for the immediate coupling of LC and NMR have given another life to this hyphenated procedure.(**78,79**)These advancements incorporate new curl and stream cell plan for high affectability, new RF framework for various dissolvable concealment and improved unique reach angle elution capacity, and programmed top picking/putting away abilities. **APPLICATIONS**

1. Identification of drug degradation products.

2. Low level impurities can be isolated and identified.

3. This technique is used for tracking pesticides, herbicides & organic pollutant for environmental monitoring.

4. Differences in electrophoretic motilities and structural information.

5.It provides information toward the structure elucidation of natural products.

6. The analysis of complex mixtures of all types, particularly the analysis of natural products and drug-related metabolites in biofluids. (80-81).

CE-MS:

CE an automated separated technique presented in the mid-1990s. CE investigation is driven by an electric field, acted in limited cylinders, and can bring about the quick partition of a large number of various mixtures. The adaptability and the numerous ways that CE can be utilized imply that practically everything atoms can be isolated utilizing this incredible technique. It isolates species by applying voltage across support filled vessels, and is by and large utilized for isolating particles that move at various paces when voltage is applied, contingent upon their size and charge. The solutes are viewed as tops as they go through the indicator and the territory of each pinnacle is corresponding to their fixation, which permits quantitative conclusions. Investigation incorporates immaculateness assurance, measures, and follow level judgments. The enhancement of the interfacing of CE with MS can be a genuine test due to the low stream rates (10–100 mL/min) needed in CE, which is accomplished by a make-up fluid.Couplings via I ESI, (ii) MALDI, or (iii) ICP are the most commonly used CE to MS interfaces. In recent years, there has been a noticeable increase in research effort in the development of highly advanced versions of IEF. ITP is primarily used as a transient phenomenon for determining effective sample concentrations in separation capillaries prior to MS identification.CE-MS interfaces are developed in two ways: as sheath-flow devices or as sheath less devices. Both had their

advantages and disadvantages, and neither won. A significant dilution of samples is paid for the versatility in the composition of spray liquids mixed in sheath-flow interfaces.(82-89).

APPLICATIONS

Urinary biomarkers for renal diseases:

CE-MS investigation of urinary tests from patients with different kinds of ongoing renal infections brought about the foundation of boards that comprised of 20 to 50 urinary polypeptide markers that permitted finding and separation of IgA nephropathy, central segmental glomerulosclerosis (FSGS), membranous glomerulonephritis MGN), and insignificant change sickness. Albeit those underlying investigations showed the capability of urinary proteome examination, they did exclude a blinded approval set. In resulting examines that utilized an improved and strong example arrangement convention and proper measurable assessment of the individual biomarkers, those underlying discoveries were affirmed and approved. (90,91)

For Synthetic in-vitro Glycolysis Studies:

A synthetic in-vitro glycolysis was remade from ten purified Escherichia coli (E. coli) proteins to acquire a superior comprehension of the guideline of successive enzymatic responses. The versatility of glucose as a starting material for chemical synthesis has generated a lot of interest in developing natural and medicinal compounds from microbial cells.(92,93) Capillary electrophoresis combined with electrospray ionization-mass spectrometry has been reported as one method for thorough investigation of intracellular metabolites. Spectrometry (CE-ESI-MS) has developed as a particularly useful too a strong new technique for analyzing many charged objects at the same time compounds. Using CE-ESI-MS, we demonstrate that ten naturally designed enzymes can collaborate in a synthetic pathway in vitro, resulting in the synthesis of consecutive glycolytic metabolites and. According to the incubation time, both DHAP and pyruvate accumulate at a high rate.(94-96)

Characterization of Monoclonal Antibodies:

MAbs are highly heterogeneous proteins, thereby requiring a battery of sophisticated analytical technologies for their complete characterization. Mass spectrometry (MS) has become an essential analytical tool for the structural characterization of mAbs. Monoclonal antibodies (mAbs) are tetrameric glycoproteins with a molecular mass of about 150 kDa, two heavy chains and two light chains interconnected by multiple disulfide links, and a molecular mass of around 150 kDa.possessing at least one N-glycosylation site that is conserved. These orthogonal analytical methods are designed to distinguish the primary isoform of an antibody from micro-variants.Micro-variants are commonly observed whenmAbs are analyzed by charge-based separation techniques such as isoelectric focusing gel electrophoresis(IEF), capillary isoelectric focusing gel electrophoresis (CIEF).(97,98)

For Forensic Analysis:

Recent advancements in the field of DNA testing have aided law enforcement significantly. Biological evidence from a crime scene can now be linked to the culprit by forensic laboratory.(99)It can reliably rule out people who have been wrongfully charged. InNumerous advancements in DNA testing methods have happened over the last two decades, most notably amongthem the development of polymerase chain reaction(PCR)-based typing methods. The use of short tandem repeat (STR) markers has become mainstream in the forensic DNA typing field. Forensic DNA analysis has relied extensively on short tandem repeat (STR) variations. They were chosen because they are extremely polymorphic and discriminative among people. The Combined DNA Index System has adopted these loci as reference loci (CODIS)as well as assisting in the global development of crime national DNA databases (NDNADs).(100)

Applications of forensic DNA testing:

DNA databasing:

DNA databasing has become a valuable measurable device and as more examples are added to the data set the likelihood of a case-to-case match or case to sentenced guilty party match increments. One issue

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confronting generally criminological research facilities in the United States concerning the data set is the excess of indicted wrongdoer tests holding up to be handled and gone into the data set.(101-103)

Identification of Anthraquinone Colouring Matters in Natural Red Dyes:

For the recognizable proof of anthraquinone shading segments of cochineal, lac color and madder, characteristic red dye stuffs regularly utilized by old painters. With the end goal of such investigation, ESI-MS was discovered to be a substantially more suitable location procedure than DAD one attributable to its higher affectability (identification limits in the reach $0.1-0.5 \mu \text{gml}-1$) and selectivity.(**104**)The strategy created made it conceivable to distinguish unequivocally carmine corrosive and brief acids A, B and E as shading matters in the inspected arrangements of cochineal and lac color, separately.Because of it's selective and sensitive detection capabilities of ESI-MS, it is possible to identify anthraquinone natural colouring materials isolated by CE without a doubt, even when they are not anthraquinone. Hydrophobic alizarin, purpurin, and other hydrophobic alizarins are examples of well-separated alizarins, emodin.When more polar chemicals, such as laccaic acids or carminic acid, must be recognized, the developed approach comes in handy. CE allows you to tell the difference between laccaic acids and other acids which is absolutely impossible with LC. (**105,106**)

Determination of Drugs in Human Plasma:

Utilizing CE/MS techniques, the portrayed API III Quadra post framework gave a worthy particle current electropherogram from sub picomole levels of the focused on intensifies stacked onto the chip.(107) The comparing electropherogram for the standard arrangement of carnitines at the 1–500 μ g/mL level was acquired through SIM CE/MS methods (R2 > 0.99). Furthermore, investigations of sustained examples of imipramine desipramine were estimated comparative with their relating d3 inward principles to get adjustment bends going from 5 to 500 μ g/mL in human plasma (R2 > 0.99). The intra-measure exactness went from 4.1 to 7.3% RSD. The intra-measure precision went from 94.0 to 104%.(108,109)

Analysis of Inorganic Species:

Capillary electrophoresis (CE) has gained a positive response as a method for separating tiny and big organic molecules. The selectable level of particle adduct DE grouping and sub-atomic fracture in the MS interface district permits the framework to be worked as a natural analyzer or as a sub-atomic locator reasonable for oxidation state conclusions. Both inorganic anions and cations (counting salts, basic earth, change metals, and lanthanides) are dissected by CE–MS.(**110**). International intercomparison exercises on rain water samples performed in the framework of the European Union (EU) AQUACON project. Once nonselective detection is combined with CE or IC (e.g., indirect UV or conductivity detection), all of the target analytes must be resolved from each other.interfering matrix components) to give a favorable outcome. confirmation, as well as the use of external calibration methods. (**111,112**)

Developments for Profiling Metabolites of Steroid Hormone Metabolism:

Thirteen steroids were remembered for the technique advancement, and the chose were metabolites engaged with significant pathways of steroid biosynthesis. Albeit just eight of them could be isolated and recognized with UV, they could be distinguished by ESI-MS utilizing chosen particle checking (SIM) strategy.Endogenous adrenal steroids are difficult to analyze with CE and related methods. Corticosteroids seldom dissociate below pH 12 because of their poor hydrophilic structure. Couple MS spectra were additionally gathered. The most reduced restrictions of discovery were 10-100 ng/mL for cortisone, corticosterone, hydrocortisone, and testosterone. Different steroids could be identified at 500-1000 ng/ml. The distinguishing proof of cortisone, corticosterone, hydrocortisone, estrogen and testosterone were made in understanding pee tests and their focuses were 1-40 μ g/L.The quantitative and complete characterization of steroid hormone status in body fluids and tissues is important in assessing human health because mammalian steroid hormones are an important class of metabolites with varied biochemical and physiological activities. The advantages of capillary electrophoresis are high efficiency, requiring minute amounts of sample and quantitatively consuming limited amounts of reagents. (113-115)

Amino Acid Analysis:

To investigate free amino acids all the while a low acidic pH condition was utilized to give positive charge on entire amino acids. The decision of the electrolyte and its fixation affected the goal and pinnacle state of the amino acids, and 1M formic corrosive was chosen as the ideal electrolyte. (116,117)

FOOD SAFETY CE-MS APPLICATION

In the last 5 years, and concerning food safety applications,CE-MS has primarily been used to determine the presence of various pesticides and antibiotics in foods and water. Despite the fact that there are some other interesting applications such as the analysis of biogenic amines , alkaloids , antidepressants , disinfection byproducts , products formed during the processing or the analysis of biogenic amines , alkaloids , antidepressants , disinfection byproducts , byproducts , products formed during the processing or the analysis of biogenic amines , alkaloids , antidepressants , disinfection byproducts , products formed during the processing or acrylamides , food preparation, and speciation analysis, breakdown products of chemical warfare agents.(118,119)CE-MS has also been used to investigate transgenic foods and hazardous oligopeptides.(120)

Conclusion: The hyphenated techniques are very useful in the above fileds and also in pesticide analysis, chiral chromatography and food analysis

REFERAENCES:

- 1. Wilson, I. ., & Brinkman, U. A. T. (2003). Hyphenation and hypernation. Journal of Chromatography A, 1000(1-2), 325–356.
- 2. Wolfender, J.-L., Ndjoko, K., &Hostettmann, K. (2001). The potential of LC-NMR in phytochemical analysis. Phytochemical Analysis, 12(1), 2–22.
- 3. A review of: "Diode array detection in HPLC, edited by L. Huber & S. A. George, Chromatographic Science Series, J. Cazes Editor, Volume 62, Marcel Dekker, Inc. New York, Basel, Hongkong, 1993, Journal of Liquid Chromatography, 17(18), 4057–4058.
- 4. Niessen, W. M. ., &Tinke, A. . (1995). Liquid chromatography-mass spectrometry General principles and instrumentation. Journal of Chromatography A, 703(1-2), 37–57.
- 5. Joshi RR, Gupta KR and Patil SS: Hyphenated Technique- A Boon to Analytical World. Int J Pharm Sci Res. 3(11); 4184-4191.
- Wolfender, J.-L., Terreaux, C., &Hostettmann, K. (2000). The Importance Of LC-MS And LC-NMR In The Discovery Of New Lead Compounds From Plants. Pharmaceutical Biology, 38(sup1), 41–54.
- 7. Shockcor, J. P., Unger, S. E., Wilson, I. D., Foxall, P. J. D., Nicholson, J. K., & Lindon, J. C. (1996). Combined HPLC, NMR Spectroscopy.
- 8. Niessen, W. M. A. (2003). Progress in liquid chromatography–mass spectrometry instrumentation and its impact on high-throughput screening. Journal of Chromatography A, 1000(1-2), 413–436.
- 9. Sarker, S. D., & Nahar, L. (2012). Hyphenated Techniques and Their Applications in Natural Products Analysis. Natural Products Isolation, 301–340.
- Bringmann, G., Messer, K., Wohlfarth, M., Kraus, J., Dumbuya, K., &Rückert, M. (1999). HPLC–CD On-Line Coupling in Combination with HPLC–NMR and HPLC–MS/MS for the Determination of the Full Absolute Stereostructure of New Metabolites in Plant Extracts. Analytical Chemistry, 71(14), 2678–2686.
- 11. Patel, K. N., Patel, J. K., Patel, M. P., Rajput, G. C., & Patel, H. A. (2010). Introduction to hyphenated techniques and their applications in pharmacy. Pharmaceutical Methods, 1(1), 2–13.
- 12. Chauhan, A. (2014). GC-MS Technique and its Analytical Applications in Science and Technology. Journal of Analytical & Bioanalytical Techniques, 5(6).

- Jinno, K., Fujimoto, C., & Hirata, Y. (1982). An Interface for the Combination of Micro High-Performance Liquid Chromatography and Infrared Spectrometry. Applied Spectroscopy, 36(1), 67–69.
- 14. Jinno K. Infrared Detect, in Encyclopedia of Chromatography. In: Cazes J, Editor. New York, USA: Marcel Dekker; 2001
- 15. Amirav, A., Gordin, A., Poliak, M., &Fialkov, A. B. (2008). Gas chromatography-mass spectrometry with supersonic molecular beams. Journal of Mass Spectrometry, 43(2), 141–163.
- 16. Bliesner DM (2006) Validating Chromatographic Methods: A Practical Guide. John Wiley and Sons.
- 17. Fialkov, A. B., Steiner, U., Jones, L., & Amirav, A. (2006). A new type of GC–MS with advanced capabilities. International Journal of Mass Spectrometry, 251(1), 47–58.
- Alon, T., & Amirav, A. (2006). Isotope abundance analysis methods and software for improved sample identification with supersonic gas chromatography/mass spectrometry. Rapid Communications in Mass Spectrometry, 20(17), 2579–2588.
- 19. Adams, R.P. (2007) Identification of Essential Oil Components by Gas Chromatography/Mass Spectrometry. 4th Edition Allured Publishing Corporation, Carol Stream.
- Stein, S. E., & Scott, D. R. (1994). Optimization and testing of mass spectral library search algorithms for compound identification. Journal of the American Society for Mass Spectrometry, 5(9), 859–866.
- 21. Handley AJ, Adlard ER (2001) Gas chromatographic techniques and Applications. Sheffield Academic, London.
- 22. Pesticides Method Reference, 2nd ed. 2011, Thermo Fisher Scientific, Austin, TX, USA.
- 23. Leary, P. E., Kammrath, B. W., Lattman, K. J., &Beals, G. L. (2019). EXPRESS: Deploying Portable Gas Chromatography–Mass Spectrometry (GC–MS) to Military Users for the Identification of Toxic Chemical Agents in Theater. Applied Spectroscopy.
- 24. Eiceman, G. A. (2006). Instrumentation of Gas Chromatography. Encyclopedia of Analytical Chemistry.
- 25. Kitson FG, Larsen BS, McEwen CN (1996) Gas chromatography and mass spectrometry: a practical guide. Academic Press, Boston.
- Medeiros, P. M. (2018). Gas Chromatography–Mass Spectrometry (GC–MS). Encyclopedia of Geochemistry, 530–535.
- Hunter, A. P., Desbrow, C. E., & Severs, J. C. (1993). Mass spectral and GC data of drugs, poisons, pesticides, pollutants and their metabolites, Parts 1-3, 2nd edition. Edited by K. Pfleger, H. H. Maurer and A. Weber Publisher: VCH, Weinheim, Germany 1992, pp. 3220, 1290.00 DM. ISBN 3 527 26989 4. Rapid Communications in Mass Spectrometry, 7(6), 567–567.
- 28. Modern Practice of Gas Chromatography, 4th ed Edited by Robert L. Grob (Villanova University) and Eugene F. Barry (University of Massachusetts, Lowell). John Wiley & Sons.
- Zhao, B., Zhang, S., Zhou, Y., He, D., Li, Y., Ren, M., ... Fang, J. (2015). Characterization and quantification of PAH atmospheric pollution from a large petrochemical complex in Guangzhou: GC–MS/MS analysis. Microchemical Journal, 119, 140–144.
- 30. Giannelli PC, Imwinkelried EJ (1999) Drug Identification: Gas Chromatography. In Scientific Evidence, Volume 2, Charlottesville: Lexis Law Publishing.
- Ziegenhals, K., Hübschmann, H.-J., Speer, K., & Jira, W. (2008). Fast-GC/HRMS to quantify the EU priority PAH. Journal of Separation Science, 31(10), 1779–1786.

- Lakshmi HimaBindu MR, AngalaParameswari S, Gopinath C (2013) A Review on GC-MS and Method Development and Validation. International Journal of Pharmaceutical Quality Assurance 4: 42-51.
- 33. Priya V, Jananie RK, Vijayalaxmi K (2012) GC/MS Determination of Bioactive Components PleurotusOstreatus. Int Research Journal of Pharmacy 3: 150-151.
- Lanzarotta, A., Falconer, T., McCauley, H., Lorenz, L., Albright, D., Crowe, J., & Batson, J. (2016). Simultaneous Orthogonal Drug Detection Using Fully Integrated Gas Chromatography with Fourier Transform Infrared Detection and Mass Spectrometric Detection. Applied Spectroscopy, 71(5), 1050–1059.
- 35. Vissers, J. P. C., Claessens, H. A., &Cramers, C. A. (1997). Microcolumn liquid chromatography: instrumentation, detection and applications. Journal of Chromatography A, 779(1-2), 1–28.
- Somsen, G. W., Gooijer, C., Velthorst, N. H., & Brinkman, U. A. T. (1998). Coupling of column liquid chromatography and Fourier transform infrared spectrometry. Journal of Chromatography A, 811(1-2), 1–34.
- Somsen, G. W., & Visser, T. (2006). Liquid Chromatography/Infrared Spectroscopy. Encyclopedia of Analytical Chemistry.
- Baumann, W. (1987). HPLC-Detektoren: Detectors for Liquid Chromatography. Von E. S. Yeung. John Wiley & Sons. New York - Chichester - Brisbane - Toronto - Singapore 1986. 52.75 £. ISBN 0-471-84735-6. NachrichtenAusChemie, Technik Und Laboratorium, 35(10), 1055–1055.
- Wolfender, J.-L., Hostettmann, K., Abe, F., Nagao, T., Okabe, H., & Yamauchi, T. (1995). Liquid chromatography combined with thermospray and continuous-flow fast atom bombardment mass spectrometry of glycosides in crude plant extracts. Journal of Chromatography A, 712(1), 155–168.
- 40. Niessen, W.M. (1998). Advances in instrumentation in liquid chromatography-mass spectrometry and related liquid-introduction techniques. Journal of Chromatography A, 794(1-2), 407–435.
- 41. Niessen, W. M. ., &Tinke, A. . (1995). Liquid chromatography-mass spectrometry General principles and instrumentation. Journal of Chromatography A, 703(1-2), 37–57.
- 42. Dugo, P., Mondello, L., Dugo, L., Stancanelli, R., &Dugo, G. (2000). LC-MS for the identification of oxygen heterocyclic compounds in citrus essential oils. Journal of Pharmaceutical and Biomedical Analysis, 24(1), 147–154.
- 43. Wolfender, J.-L., Rodriguez, S., &Hostettmann, K. (1998). Liquid chromatography coupled to mass spectrometry and nuclear magnetic resonance spectroscopy for the screening of plant constituents. Journal of Chromatography A, 794(1-2), 299–316.
- Niessen, W. M. (1998). Advances in instrumentation in liquid chromatography-mass spectrometry and related liquid-introduction techniques. Journal of Chromatography A, 794(1-2), 407–435.
- 45. R. Willoughby, E. Sheehan, S. Mitrovich, A Global View of LC-MS, Global View Publishing, Pittsburgh, PA, 1998.
- 46. Hostettmann, K., Wolfender, J.-L., & Rodriguez, S. (1997). Rapid Detection and Subsequent Isolation of Bioactive Constituents of Crude Plant Extracts. Planta Medica, 63(01), 2–10.
- 47. Shujing Ding; Inez Schoenmakers; Kerry Jones; Albert Koulman; Ann Prentice; Dietrich A. Volmer (2010). Quantitative determination of vitamin D metabolites in plasma using UHPLC-MS/MS., 398(2), 779–789.
- 48. DanijelDjukovic; Hamid R. Baniasadi; Ravi Kc; Zane Hammoud; Daniel Raftery (2010). Targeted serum metabolite profiling of nucleosides in esophageal adenocarcinoma. , 24(20), 3057–3062.
- Wei, C.; Zhu, P.; Shah, S. J.; Blair, I. A. (2009). 15-oxo-Eicosatetraenoic Acid, a Metabolite of Macrophage 15-Hydroxyprostaglandin Dehydrogenase That Inhibits Endothelial Cell Proliferation. Molecular Pharmacology, 76(3), 516–525.

- Adachi, J., Hashiguchi, K., Nagano, M., Sato, M., Sato, A., Fukamizu, K., ... Tomonaga, T. (2016). Improved Proteome and Phosphoproteome Analysis on a Cation Exchanger by a Combined Acid and Salt Gradient. Analytical Chemistry, 88(16), 7899–7903.
- 51. Masuda, T., Tomita, M., &Ishihama, Y. (2008). Phase Transfer Surfactant-Aided Trypsin Digestion for Membrane Proteome Analysis. Journal of Proteome Research, 7(2), 731–740.
- Chen, Q., Yan, G., Gao, M., & Zhang, X. (2015). Ultrasensitive Proteome Profiling for 100 Living Cells by Direct Cell Injection, Online Digestion and Nano-LC-MS/MS Analysis. Analytical Chemistry, 87(13), 6674–6680.
- Kawashima, Y., Watanabe, E., Umeyama, T., Nakajima, D., Hattori, M., Honda, K., & Ohara, O. (2019). Optimization of Data-Independent Acquisition Mass Spectrometry for Deep and Highly Sensitive Proteomic Analysis. International Journal of Molecular Sciences, 20(23), 593
- Yan, B., Collins, N., Wheatley, J., Irving, M., Leopold, K., Chan, C., ... Zhao, J. (2004). High-Throughput Purification of Combinatorial Libraries I: A High-Throughput Purification System Using an Accelerated Retention Window Approach. Journal of Combinatorial Chemistry, 6(2), 255–261.
- 55. Blom, K. F. (2002). Two-Pump at-Column-Dilution Configuration for Preparative Liquid Chromatography–Mass Spectrometry. Journal of Combinatorial Chemistry, 4(4), 295–301.
- Blom, K. F., Glass, B., Sparks, R., & Combs, A. P. (2004). Preparative LC–MS Purification: Improved Compound-Specific Method Optimization. Journal of Combinatorial Chemistry, 6(6), 874–883.
- 57. Maestroni, B., Besil, N., Bojorge, A., Gérez, N., Pérez-Parada, A., Cannavan, A., ... Cesio, M. V. (2020). Optimization and validation of a single method for the determination of pesticide residues in Peumusboldus Molina leaves using GC-MSD, GC-MS/MS and LC-MS/MS. Journal of Applied Research on Medicinal and Aromatic Plants, 100254.
- 58. Yi-Yung Hsia, David; Berman, Julian L.; Slatis, Herman M. (1964). Screening Newborn Infants for Phenylketonuria. JAMA, 188(3),
- 59. Chace, D H; Hillman, S L; Millington, D S; Kahler, S G; Roe, C R; Naylor, E W (1995). Rapid diagnosis of maple syrup urine disease in blood spots from newborns by tandem mass spectrometry. Clinical Chemistry, 41(1), 62–68.
- 60. D. S. Millington; N. Kodo; D. L. Norwood; C. R. Roe (1990). Tandem mass spectrometry: A new method for acylcarnitine profiling with potential for neonatal screening for inborn errors of metabolism.
- 61. Chace, Donald H; Hillman, Steven L; Van Hove, Johan L K; Naylor, Edwin W (1997). Rapid diagnosis of MCAD deficiency: quantitative analysis of octanoylcarnitine and other acylcarnitines in newborn blood spots by tandem mass spectrometry. Clinical Chemistry, 43(11), 2106–2113.
- 62. Richard Koch; Friedrich Trefz; Susan Waisbren (2010). Psychosocial issues and outcomes in maternal PKU., 99(supp-S), 0
- 63. Michael Vogeser; Christoph Seger (2008). A decade of HPLC–MS/MS in the routine clinical laboratory Goals for further developments. , 41(9), 649–662.
- 64. Taylor, Paul J. (2004). Therapeutic Drug Monitoring of Immunosuppressant Drugs by High-Performance Liquid Chromatography–Mass Spectrometry. Therapeutic Drug Monitoring, 26(2), 215–219.
- Brandhorst, G. (2006). Quantification by Liquid Chromatography Tandem Mass Spectrometry of Mycophenolic Acid and Its Phenol and Acyl Glucuronide Metabolites. Clinical Chemistry, 52(10), 1962–1964.
- 66. Franck Saint-Marcoux; François-LudovicSauvage; Pierre Marquet (2007). Current role of LC-MS in therapeutic drug monitoring. , 388(7), 1327–1349.
- 67. Beuhler, R. J.; Flanigan, E.; Greene, L. J.; Friedman, L. (1974). Proton transfer mass spectrometry of peptides. Rapid heating technique for underivatized peptides containing arginine. Journal of the American Chemical Society, 96(12), 3990–3999.

- 68. R.P.W. Scott; C.G. Scott; M. Munroe; J. Hess Jr. (1974). Interface for on-line liquid chromatography—mass spectroscopy analysis., 99(none), 395–405.
- Al-Asmari, A. I.; Anderson, R. A.; Appelblad, P. (2010). Direct Determination of Ethyl Glucuronide and Ethyl Sulfate in Postmortem Urine Specimens Using Hydrophilic Interaction Liquid Chromatography-Electrospray Ionization-Tandem Mass Spectrometry. Journal of Analytical Toxicology, 34(5), 261–272.
- 70. FORSHUFVUD, STEN; SMITH, HAMILTON; WASSÉN, ANDERS (1961). Arsenic Content of Napoleon I's Hair Probably taken Immediately after his Death. , 192(4798), 103–105.
- Nishikawa, M., &Tsuchihashi, H. (1998). Applications of Lc/Ms in Forensic Chemistry. Journal of Toxicology: Toxin Reviews, 17(1), 13–26.
- Nishikawa, M., Nakajima, K., Tatsuno, M., Kasuya, F., Igarashi, K., Fukui, M., &Tsuchihashi, H. (1994). The analysis of cocaine and its metabolites by liquid chromatography/atmospheric pressure chemical ionization-mass spectrometry (LC/APCI-MS). Forensic Science International, 66(3), 149–158.
- 73. Marais, F. (2003). On-line LC NMR and Related Techniques Edited by Klaus Albert. John Wiley & Sons: New York. 2002. 290 pp. Organic Process Research & Development, 7(4), 609–609.
- 74. Corcoran, O., &Spraul, M. (2003). LC–NMR–MS in drug discovery. Drug Discovery Today, 8(14), 624–631.
- 75. Albert K. On-line LC-NMR and Related Techniques. London: Wiley; 2002.
- Sidelmann, U. G., Bjørnsdottir, I., Shockcor, J. P., Hansen, S. H., Lindon, J. C., & Nicholson, J. K. (2001). Directly coupled HPLC-NMR and HPLC-MS approaches for the rapid characterisation of drug metabolites in urine: application to the human metabolism of naproxen. Journal of Pharmaceutical and Biomedical Analysis, 24(4), 569–579.
- 77. Albert K: On-line use of NMR detection in separation chemistry. J Chromatogr A 1995; 703: 123-47.
- 78. Wolfender JL, Ndjoko K, Hostettmann K: LC/NMR in natural products chemistry. Curr Org Chem 1998; 2: 575-96.
- 79. N J.C. Bailey, P.D. Stanley, S.T. Hadfifield, J.C. lindon, J.K. Nicholson, Rapid Commun. Mass Spectrom. 14 (2000) 679
- 80. T h. Hankemeier, E. Hooijschuur, R.J.J. Vreuls, U.A.Th.Brinkman, T.Visser, J. High Resolut. Chromatogr. 21 (1998) 341.
- Wolfender, J.-L., Terreaux, C., &Hostettmann, K. (2000). The Importance Of LC-MS And LC-NMR In The Discovery Of New Lead Compounds From Plants. Pharmaceutical Biology, 38(sup1), 41–54.
- Dunayevskiy, Y. M., Vouros, P., Wintner, E. A., Shipps, G. W., Carell, T., &Rebek, J. (1996). Application of capillary electrophoresis-electrospray ionization mass spectometry in the determination of molecular diversity. Proceedings of the National Academy of Sciences, 93(12), 6152–6157.
- 83. Logar, J. K., Malej, A., & Franko, M. (2003). Hyphenated high performance liquid chromatography-thermal lens spectrometry technique as a tool for investigations of xanthophyll cycle pigments in different taxonomic groups of marine phytoplankton. Review of Scientific Instruments, 74(1), 776–778.
- Janda, K. D. (1994). Tagged versus untagged libraries: methods for the generation and screening of combinatorial chemical libraries. Proceedings of the National Academy of Sciences, 91(23), 10779–10785.
- Gallop, M. A., Barrett, R. W., Dower, W. J., Fodor, S. P. A., & Gordon, E. M. (1994). Applications of Combinatorial Technologies to Drug Discovery. 1. Background and Peptide Combinatorial Libraries. Journal of Medicinal Chemistry, 37(9), 1233–1251.

- Smith, R. D., Wahl, J. H., Goodlett, D. R., &Hofstadler, S. A. (1993). Capillary electrophoresis/mass spectrometry. Analytical Chemistry, 65(13), 574A–584A.
- 87. Yamashita, M., & Fenn, J. B. (1984). Negative ion production with the electrospray ion source. The Journal of Physical Chemistry, 88(20), 4671–4675.
- 88. Bakry, R., Huck, C. W., Najam-ul-Haq, M., Rainer, M., & Bonn, G. K. (2007). Recent advances in capillary electrophoresis 30(2), 192-201.
- Zhong, X., Zhang, Z., Jiang, S., & Li, L. (2013). Recent advances in coupling capillary electrophoresis-based separation techniques to ESI and MALDI-MS. ELECTROPHORESIS, 35(9), 1214–1225.
- Kaiser T, Kamal H, Rank A, Kolb HJ, Holler E, Ganser A, Hertenstein B, Mischak H, Weissinger EM. Proteomics applied to the clinical follow-up of patients after allogeneic hematopoietic stem cell transplantation. Blood. 2004;104:340–349.
- Mischak, H., Coon, J. J., Novak, J., Weissinger, E. M., Schanstra, J. P., &Dominiczak, A. F. (2009). Capillary electrophoresis-mass spectrometry as a powerful tool in biomarker discovery and clinical diagnosis: An update of recent developments. Mass Spectrometry Reviews, 28(5), 703–724.
- Y., L., J., C., & S.-Y., L. (2001). Biotechnological production of pyruvic acid. Applied Microbiology and Biotechnology, 57(4), 451–459.
- Soga, T., Ueno, Y., Naraoka, H., Ohashi, Y., Tomita, M., &Nishioka, T. (2002). Simultaneous Determination of Anionic Intermediates for Bacillussubtilis Metabolic Pathways by Capillary Electrophoresis Electrospray Ionization Mass Spectrometry. Analytical Chemistry, 74(10), 2233–2239.
- Itoh, A., Ohashi, Y., Soga, T., Mori, H., Nishioka, T., & Tomita, M. (2004). Application of capillary electrophoresis-mass spectrometry to syntheticin vitro glycolysis studies. ELECTROPHORESIS, 25(13), 1996–2002.
- Soga, T., &Imaizumi, M. (2001). Capillary electrophoresis method for the analysis of inorganic anions, organic acids, amino acids, nucleotides, carbohydrates and other anionic compounds. ELECTROPHORESIS, 22(16), 3418–3425.
- 96. Roessner, C. A., & Scott, A. I. (1996). Achieving natural product synthesis and diversity via catalytic networking ex vivo. Chemistry & Biology, 3(5), 325–330.
- Doneanu, C., Xenopoulos, A., Fadgen, K., Murphy, J., Skilton, S. J., Prentice, H., Chen, W. (2012). Analysis of host-cell proteins in biotherapeutic proteins by comprehensive online two-dimensional liquid chromatography/mass spectrometry. mAbs, 4(1), 24–44.
- Gahoual, R., Beck, A., Leize-Wagner, E., & François, Y.-N. (2016). Cutting-edge capillary electrophoresis characterization of monoclonal antibodies and related products. Journal of Chromatography B, 1032, 61–78.
- 99. Price, P. (1991). Standard definitions of terms relating to mass spectrometry. Journal of the American Society for Mass Spectrometry, 2(4), 336–348.
- 100.P. Gill, L. Fereday, N. Morling, P.M. Schneider, The evolution of DNA databasesrecommendations for new European STR loci, Forensic Sci. Int. 156 (2006) 242–244
- 101.Budowle, B., Smith, J., Moretti, T., DiZinno, J. A., DNA Typing Protocols: Molecular Biology and Forensic Analysis, Eaton Publishing, Natick, MA 2000, pp. 157–188.
- 102.Moretti, T. R., Baumstark, A. L., Defenbaugh, D. A., Keys, K. M., Smerick, J. B., Budowle, B., J. Forensic Sci. 2001, 46, 647–660
- 103.Murray, V., Nguyen, T. V., & Chen, J. K. (2012). The Use of Automated Sequencing Techniques to Investigate the Sequence Selectivity of DNA-Damaging Agents. Chemical Biology & Drug Design, 80(1), 1–8.
- 104.Puchalska, M., Orli?ska, M., Ackacha, M. A., Po-Pawlak, K., & Jarosz, M. (2003). Identification of anthraquinone coloring matters in natural red dyes by electrospray mass spectrometry coupled to capillary electrophoresis. Journal of Mass Spectrometry, 38(12), 1252–1258.
- 105.Derksen, G. C. ., Niederländer, H. A. ., & van Beek, T. A. (2002). Analysis of anthraquinones in Rubia tinctorum L. by liquid chromatography coupled with diode-array UV and mass spectrometric detection. Journal of Chromatography A, 978(1-2), 119–127.

- 106. Ackacha, M. A., Połeć-Pawlak, K., & Jarosz, M. (2003). Identification of anthraquinone coloring matters in natural red dyestuffs by high performance liquid chromatography with ultraviolet and electrospray mass spectrometric detection. Journal of Separation Science, 26(11), 1028–1034.
- 107.Boatto, G., Nieddu, M., Dessì, G., Manconi, P., &Cerri, R. (2007). Determination of four thio phenethylamine designer drugs (2C-T-series) in human plasma by capillary electrophoresis with mass spectrometry detection. Journal of Chromatography A, 1159(1-2), 198–202.
- 108.Lin, X., Cai, Y., Yan, J., Zhang, L., Wu, D., & Li, H. (2014). Determination of Gabapentin in Human Plasma and Urine by Capillary Electrophoresis with Laser-Induced Fluorescence Detection. Journal of Chromatographic Science, 53(6), 986–992.
- 109.Rice, A. S. C., &Maton, S. (2001). Gabapentin in postherpetic neuralgia: a randomised, double blind, placebo controlled study. Pain, 94(2), 215–224.
- 110. Hjertén, S. (1967). Free zone electrophoresis. Chromatographic Reviews, 9(2), 122-219.
- 111. Van den Hoop, M. A. G. ., Cleven, R. F. M. ., van Staden, J. J., &Neele, J. (1996). Analysis of fluoride in rain water comparison of capillary electrophoresis with ion chromatography and ion-selective electrode potentiometry. Journal of Chromatography A, 739(1-2), 241–248.
- 112.Karthikeyan, S., & Hirata, S. (2003). Arsenic Speciation in Environmental Samples. Analytical Letters, 36(11), 2355–2366.
- 113.Capellino, S., Riepl, B., Rauch, L., Angele, P., Cutolo, M., & Straub, R. H. (2006). Quantitative determination of steroid hormone receptor positive cells in the synovium of patients with rheumatoid arthritis and osteoarthritis: is there a link to inflammation Annals of the Rheumatic Diseases, 66(1), 53–58.
- 114.Brown, G. A., Vukovich, M. D., & King, D. S. (2004). Urinary Excretion of Steroid Metabolites after Chronic Androstenedione Ingestion. The Journal of Clinical Endocrinology & Metabolism, 89(12), 6235–6238.
- 115.Sirén, H., Seppänen-Laakso, T., &Orešič, M. (2008). Capillary electrophoresis with UV detection and mass spectrometry in method development for profiling metabolites of steroid hormone metabolism. Journal of Chromatography B, 871(2), 375–382.
- 116.Smith, J. T. (1999). Recent advancements in amino acid analysis using capillary electrophoresis. Electrophoresis, 20(15-16), 3078–3083.
- 117.Khatri, K., Klein, J. A., Haserick, J. R., Leon, D. R., Costello, C. E., McComb, M. E., &Zaia, J. (2017). Microfluidic Capillary Electrophoresis–Mass Spectrometry for Analysis of Monosaccharides, Oligosaccharides, and Glycopeptides. Analytical Chemistry, 89(12), 6645–6655.
- 118.Schmitt-Kopplin, P., &Frommberger, M. (2003). Capillary electrophoresis– mass spectrometry: 15 years of developments and applications. ELECTROPHORESIS, 24(2223), 3837–3867.
- 119.Hernández-Borges, J., Neusüß, C., Cifuentes, A., &Pelzing, M. (2004). On-line capillary electrophoresis-mass spectrometry for the analysis of biomolecules. ELECTROPHORESIS, 25(14), 2257–2281.
- 120. Van Biesen, G., Bottaro, C. S., Electrophoresis 2006, 27, 4456-4468.