Advanced Functional Materials Silicon-based Laser Desorption Ionization Mass Spectrometry for the Analysis of Biomolecules: A Progress Report --Manuscript Draft--

Manuscript Number:	
Full Title:	Silicon-based Laser Desorption Ionization Mass Spectrometry for the Analysis of Biomolecules: A Progress Report
Article Type:	Progress Report
Section/Category:	
Keywords:	silicon nanostructures; Surface functionalization; laser desorption ionization (LDI); mass spectrometry imaging (MSI).
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Additional Information:	
Question	Response
Please submit a plain text version of your cover letter here.	Dear Editor, Please find attached the manuscript "Silicon-based Laser Desorption Ionization Mass Spectrometry for the Analysis of Biomolecules: A Progress Report" co-authored by Stefania-Alexandra lakab, Pere Ràfols, María García-Altares, Oscar Yanes and Xavier Correig to be considered for publication as a Progress Report in the Advanced Functional Materials journal. Here, we present recent developments on silicon-based LDI approaches for mass spectrometry analysis of biomolecules. The facility of fabrication and surface functionalization of silicon-based substrates open a completely new chapter in the field of mass spectrometry imaging and in particular in the detection of low mass compounds. Since the first silicon-based substrate was introduced as desorption/ionization on silicon (Siuzdak et al, Nature, 1999), the number of papers published on designing and fabricating silicon nanostructures for LDI-MS has grown rapidly, with more than 100 papers published in the last 10 years, proving that is really an emerging research field. Our report will focus on the existing fabrication processes, functionalization techniques and practical applications of porous silicon and silicon nanostructures for LDI-MS. The report will also include a detailed revision on the current nanotechnology used to create silicon-based substrates with the essential requirements for an optimal LDI analysis by providing an overview of the ionization and desorption mechanisms in the interplay of the analyte with the silicon nanostructure and the laser. This progress report is intended to be useful for materials scientists or any experimentalist researcher familiar with nanotechnology and mass spectroscopy who needs to detect or maximise the detection of biomolecules using nanostructured silicon surfaces for targeted or untargeted analysis. All authors are aware and agree to the submission of the manuscript to the Advanced Functional Materials journal. I confirm that neither the manuscript nor any parts of its content

	Dr. Pere Ràfols Soler
Do you or any of your co-authors have a conflict of interest to declare?	No. The authors declare no conflict of interest.
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Silicon-based Laser Desorption Ionization Mass Spectrometry for the Analysis of Biomolecules: A Progress Report

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Keywords: silicon nanostructures, surface functionalization, laser desorption ionization (LDI), mass spectrometry imaging (MSI)

Abstract

Matrix-assisted laser desorption/ionization mass spectrometry (MALDI-MS) is widely used in the biomedical field for the label-free analysis of molecules such as drugs, lipids, peptides, and proteins. MALDI-MS can also perform Imaging experiments to map simultaneously the concentration of hundreds of compounds on biological tissues. However, the great potential of MALDI-MS Imaging for untargeted in-situ metabolomics has not been exploited yet, since organic matrices used in traditional MALDI-MS applications introduce excessive interferences in the low m/z range. For this reason, nanostructured materials and in particular silicon-based LDI strategies have become a strong alternative, since they provide a much weaker background. Herein, we review the recent developments in fabrication, functionalization and practical applications of silicon-based LDI-MS methods. We also report the basic requirements of silicon-based substrates for an optimal LDI analysis by providing an overview of the LDI mechanisms using silicon-based substrates instead of organic matrices. Finally, the extensive potential of silicon-based substrates is discussed, giving suggestions on topics of interest for future research.

1. Introduction

Matrix-assisted laser desorption/ionization mass spectrometry (MALDI MS) is one of the most versatile mass spectrometry analysis techniques that can detect a wide range of molecules in complex biological samples. It can be used to detect chemical compounds in liquid samples (i.e. urine, serum, etc.), but the great advantage of this technique is its ability to map the concentration of chemical compounds over an animal or vegetal surface. The molecular images obtained correlate spatially with the morphology of such tissues.^[1–4]

Traditionally, MALDI implies the use of an organic matrix deposited onto the sample for laser desorption/ionization. The requirements of a successful MALDI matrix are high optical absorption at the laser irradiation wavelength, good analyte incorporation into the matrix, available proton donor or acceptor for an efficient ionization and minimal fragmentation of analyte.^[3] Two of the most common organic matrices are α -cyano-4-hydroxycinnamic acid (CHCA) and 2,5- dihydroxybenzoic acid (DHB).

Although organic matrices ionize efficiently a wide variety of molecules, they present some important limitations:

- The use of an organic matrix hampers the analysis in the low mass range due to the presence of many background ion signals generated from the organic matrix clusters.^[5] This results in major disadvantages for the use of MALDI-MS in metabolomics.
- Inhomogeneity in the co-crystallization of the organic matrix with the analyte leads to a lack of reproducibility. The dried organic matrix crystals vary in size and shape across the surface, leading to differences in pixel-to-pixel MS signals intensities, thus making difficult quantitative MALDI analysis.^[5]
- Although a wide range of commercially organic matrices is available, its selection and deposition optimization are a manual and complex process. Each experiment may require a specific matrix which has to be optimized, and in some cases, chemical interactions between analyte and matrix may occur.^[2]

• Poor lateral resolution is frequently associated with the presence of matrix. This is mainly caused by heterogeneous co-crystallization of the matrix and analyte. Additionally, the solvent often interacts with tissues and induces delocalization of molecules.^[6] This issue was partially solved by finely controlled matrix deposition methods.^[3]

To overcome heterogeneous co-crystallization between matrix and analyte during spraycoating, organic matrices were deposited uniformly and homogeneously directly onto analytes via sublimation method.^[7] This method improves the lateral resolution, but given the long acquisition times needed in tissue imaging, the matrix may evaporate during measurement, rendering this approach hardly applicable.^[8] Comparing the two deposition methods, spraycoating results in higher analyte extraction efficiency (i.e., increased sensitivity) whereas sublimation improves lateral resolution.^[7]

In the quest to find alternatives to organic matrices deposited over the sample, scientific efforts turned to substrates that could promote the ionization and desorption of compounds without introducing exogenous material that could interfere with the detection of endogenous compounds. As a consequence, new solid-state substrate materials have been developed and used. Several reviews ^[5,9–12] have described the use of inorganic materials based on carbon, silicon, metals or metal-oxides that have been used extensively in LDI-MS applications as substitutes of the organic matrices. These materials have the main function to promote the ionization and desorption processes of the compounds. They have to be stable under vacuum, to absorb the laser irradiation and to not cause ion interferences in the low mass range of the spectra. Moreover, other properties such as low thermal conductivity, high electrical conductivity and high surface area to volume ratio may be considered for an efficient LDI process.^[13] They are synthesized as nanoparticles or nanostructured surfaces in order to enhance the absorption at UV and have a high area/volume ratio. The use of metallic

nanoparticles, especially noble metals ^[14–16], metal oxide nanoparticles ^[17], and silicon- and carbon-based substrates as nanostructured surfaces ^[11,18], are under rigorous investigation. Silicon presents a clear advantage over other alternatives as it is a versatile material with huge potential for the development of a large number of applications. Its properties such as chemical stability, high thermal conductivity, excellent biocompatibility, rich abundance, unique electronic, optical, and mechanical properties, have established silicon as a ubiquitous material widely used in biomedical applications.^[19,20] From the technological point of view, the microelectromechanical systems industry has developed advanced technologies with huge potential for silicon-based materials. These microelectronic technologies, based on dry high vacuum processes, share the advantage of being able to fabricate highly homogeneous, reliable and repetitive surfaces. Using all these resources, scientists have dedicated time and effort to use intrinsic or functionalized silicon in several forms (porous, nanostructured) as a substrate for LDI-MS applications.^[12]

Silicon-based substrates for matrix-free LDI-MS first appeared in 1998 in the form of porous silicon (pSi) as desorption ionization on silicon (DIOS) technique.^[21] This technique has further developed into nanostructure-initiator mass spectrometry (NIMS) where a fluorinated compound, called initiator, was impregnated into the pSi ^[22] in order to get stability over the surface and have higher ion desorption and ionization yield. Other silicon-based nanostructures have been used as well: silica nanoparticles, silicon nanowires and nanostructured silicon surfaces (ex. nanocone array).^[12]

In this review, we focus on describing the various kinds of silicon-based substrates that were successfully used in all types of LDI-MS experiments. The first section describes the fabrication and application of pSi substrates, mainly based on DIOS and NIMS technologies. The second section describes the fabrication and application of different types of 2D and 3D silicon nanostructures. The third section focuses on the LDI mechanism of each type of substrate in order to give a better understanding of the current technologies and ideologies. Lastly, we

discuss important current issues on the silicon substrate technologies for LDI-MS applications and provide guidelines to select the most suitable substrate for LDI-MS experiments for diverse applications.

2. Porous Silicon – DIOS MS and NIMS

DIOS was first developed by Siuzdak and his group in 1999^[21,23] where silicon was first used as a pSi substrate that traps the analytes deposited on the surface allowing the laser irradiation to vaporize and ionize them. This matrix-free method demonstrated its great value by detecting some compounds such as peptides and small drug molecules at concentrations as low as femtomole and attomole levels and also resulting in little or no fragmentation. Budimir et al.. reported using commercial DIOS chip, which was patented by Dr. Gary Siuzdak, from Mass Consortium Corporation (San Diego, CA, USA).^[24]

2.1. Fabrication of the porous silicon substrate

The basic DIOS fabrication process starts from a flat crystalline silicon etched by a simple galvanostatic procedure as described previously by Cullis et al. in 1997.^[25] Basically, n- or p-type silicon wafers can be etched in a solution of ethanol/hydrofluoric acid in the presence of current with or without illumination, depending on the silicon wafer type. The schematic diagram of the electrochemical etching set-up used for pSi fabrication can be seen in the supporting information (see **Figure S1**). This fabrication method has been adjusted by many groups by varying parameters such as pre-etching cleaning methods, current density, ethanol/HF solution concentrations, etching time and illumination (see **Table S1**). In most cases, the electrochemical etching was carried out in custom-built Teflon cells. The typical pSi structure created by the group of Guinan et al. ^[26] is illustrated in **Figure 1**. Some groups used the electrochemical etching method on silicon wafers previously patterned with standard

lithography methods ^[27,28], while others developed a new processing method by combining electrochemical etching and laser processing.^[29]

To create pSi, unconventional techniques have also been used. Silicon nanocavity arrays with low (9%) or high (92%) porosities have been obtained using an electron beam lithography system for high-resolution nanopatterning, then a reactive ion etching (RIE) protocol was used to create the pSi.^[30] Compared to electrochemically etched surfaces, this method uses dry chemistry that reduces drastically surface contamination and safety risks to the users inherent to the use of HF solutions. Another atypical method for obtaining pSi for DIOS-MS is described by Gaspari et al.^[31] In this study, the pSi was fabricated by coating silicon wafer pieces with a 500 nm thick nanoporous film of silicon oxide. Goto et al. used evaporation-induced self-assembly (EISA) method to create mesoporous organosilica films with surface open pores.^[32] The pSi surface was created using an amphiphilic block copolymer as a structure-directing surfactant template for the triphenylamine (TPA)-derived sol-solution that was spin-coated onto the Si substrate. The film was treated with ammonia vapor and heat to obtain a condensed stable siloxane network. Lastly, the surfactant was removed by heating in toluene at elevated temperatures.^[32] However, all fabrication methods for DIOS substrates are incomplete without the stabilization and/or functionalization of the surface.

2.2. Surface modification methods

Porous silicon substrates were developed mainly for the easy detection of low molecular weight molecules by removing unnecessary background signal from the organic matrices. Interestingly, porous silicon together with the organic matrices can have a synergistic effect and provide enhanced signal to detect larger molecules such as peptides and proteins. However, for using pSi substrates, stabilization (or passivation) of pSi is necessary because the freshly etched pSi surface is metastable due to the silicon-hydride terminations. Therefore, the surface energy configuration can be easily affected by neighboring energies, so stabilization processes are

applied. One of the most common stabilization processes is the oxidation of the pSi surface. More stabilization methods for pSi are represented in **Table S2**. The stabilized pSi surfaces are resistant to air and other external factors (or environments), they present good longevity with lifetimes greater than 9 months ^[33] and constant DIOS-MS activity over an extended period of time.

Functionalization of a DIOS chip consists in the modification of the pSi surface with functional groups that promote specific bonding between pSi and the analytes in order to enable the detection of certain classes of molecules which cannot be detected otherwise. Also, functionalization methods are used to improve ionization. To achieve this, the pSi surface is first derivatized. The derivatization process results in slight changes in the chemical property of the surface by adding specific derivatization agents. The most common derivatization methods are described below in detail in section 2.1.1. Derivatization. After the pSi is derivatized, functional groups can be added for specific bonding. The bonding can be designed according to hydrophobic, electrostatic, coordination bond, or Lewis acid-base interactions between functional group and analyte. For example, pSi is commonly functionalized to create interactions.^[34] antibody-antigen, receptor-enzyme or DNA-protein Common functionalization methods are described below in sections 2.1.3. and 2.1.4., and all surface modification methods are represented in Table 1.

2.1.1. Derivatization

The most frequently used derivatization method is called silylation and it combines ozone oxidation treatment with hydrosylilation.^[26,33,35,36,38] Essentially, the DIOS chips are oxidized by exposure to ozone plasma and the surface derivatization is finalized with the modification of hydroxyl groups by silylation. Silylating reagents can vary according to the application of the DIOS chip. For instance, Trauger et al. ^[35] derivatized the surface with BSTFA, MSTFA, HMDS, ODMCS, CDOS, APDMES, FHCS, or PFPPDCS that generated trimethylsilyl- (TMS), amine-, C8-, C18-, perfluoroalkyl-, and perfluorophenyl-derivatized surfaces, from which the

most effective terminal groups proved to be amino- and pentafluorophenyl-. Protein model systems (BSA and hemoglobin tryptic digests), three small drug molecules (Propafenone, verapamil, midazolam), simple neutral carbohydrates (Maltotriose, sucrose) and four amino acids (phenylalanine, alanine, isoleucine/leucine, glutamic acid) were analyzed and detected. The hydrophobic perfluorophenyl-derivatized surface is more responsive to hydrophobic molecules while the amine-derivatized surface is more responsive to hydrophilic molecules, demonstrating that silvl derivatization is a flexible approach for preparing functionalized DIOS chips for analyte-specific applications and for the selective adsorption of analytes. The groups in [36,37] used the same method to obtain derivatized DIOS chips but in this case, the chips were silanized via the addition of neat silane (pentafluorophenylpropyldimethylchlorosilane, F5PhPr). These chips served to study the distribution of bioactive compounds synthesized in the hypobranchial gland of the marine sea snail, D. orbita. Results showed a strong correlation between histological regions and the localization of both known and unknown metabolites. T. Guinan and team ^[26,33,38,39] have also used neat silane (F5PhPr) as silvlating agent to detect three illicit drugs: MA, MDMA and cocaine, with detection limits comparable to current techniques.^[33] The same group also detected exogenous and endogenous drug compounds from fingerprints ^[26], using neat BisF17 for high throughput quantitative analysis of methadone in saliva, plasma and urine ^[38] and neat F13 to detect amphetamines, opiates, benzodiazepines and tropane alkaloids at concentrations relevant to body fluid testing.^[39] Other more uncommon derivatization agents were used by Gaspari et al. and Tuomikonski et al.^[31,44] Tuomikoski et al. ^[44] chemically derivatized the pSi samples with 10-undecenoic acid and ethyl undecenoate to obtain organic monolayers covalently attached to the surface by Si-C bonds. With this substrate, they analyzed solutions of midazolam, propranolol, buprenorphine, psilocin, 1-naphthalene methylamine, and 2-naphthylacetic acid, dichloromethylene bisphosphonate and dichloromethyl phosphonate. Results strengthened the analysis method's validity, obtaining detection sensitivity at the 100-150 fmol level for the pharmaceutical compounds. Similarly,

Gaspari et al. ^[31] derivatized their pSi through pentane rinsing to selectively capture three low molecular weight peptides substance P, renin substrate tetrapeptide, and angiotensin to detect calcitonin in spiked in human plasma.

2.1.2. Nanostructure initiator mass spectrometry (NIMS)

DIOS demonstrated such versatility by combining nanoporous surfaces and functionalization methods that scientists were inspired to evolve this technique into a new one: nanostructureinitiator mass spectrometry (NIMS). NIMS focuses on combining the properties of nanostructured materials and fluorinated compounds. This new MS analysis technique uses compounds called 'initiator' that are trapped in nanostructured surfaces to release and ionize intact molecules adsorbed on the surface.^[22] NIMS has to be differentiated from silylated pSi since the fluorinated compounds are bound differently. In case of the silylated pSi, the fluorinated compounds are chemically bound to the surface while in case of NIMS they are physically adsorbed onto the surface. In other words, NIMS has taken advantage of the nanostructured surfaces' ability to trap liquids and it is not using any functionalization methods to activate the surface. Most research groups use pSi as nanostructured substrate and Bis17 as initiator.^[22,54] Perfluorinated siloxanes are preferred for NIMS because they are effectively trapped within the nanostructured surface and have the best performance.

The well-known manufacturing process of pSi and easy implementation of initiators has led to the successful use of NIMS for various applications. Firstly, the versatility of NIMS was demonstrated by Northen et al.^[22] The group tested various initiators for a large number of applications: direct analysis of blood and urine, characterization of peptide microarrays, detection of endogenous phospholipids of aMDA-MB-231 cancer cell line and MSI analysis of mouse embryo tissue. A protocol for preparing and applying NIMS surfaces has been published by the same group where they give detailed step by step instructions accompanied by

supplementary movies.^[22,54] Secondly, NIMS performance was compared with DIOS and (nanostructure assisted laser desorption ionization) NALDI for the detection of four drug classes: amphetamines, benzodiazepines, opiates, and tropane alkaloids.^[39] NIMS produced better LODs for oxycodone in both water and PBS. Lastly, a study on the pSi morphology showed the pore-size-dependent analyte selectivity of NIMS. This selective behavior manifests mainly through physical interactions of analyte molecules and NIMS surfaces. The sensitivity of NIMS is directly affected by the pore and analyte size. In this study, analytes ranging from m/z 175.12 to m/z 3657.92 and substrates of pore size from ~4 to 12 nm (porosity from ~7% to 70%) were analyzed.^[51]

Variations on NIMS technology quickly appeared. In the first place, NIMS was modified by use of new augmenting components such as organic matrices or metallic nanoparticles. Moening et al. used a sublimated organic matrix (DHB) together with NIMS to develop a new hybrid ionization approach called matrix-enhanced nanostructure initiator mass spectrometry (ME-NIMS). ME-NIMS improved the performance of conventional NIMS reducing the limit of detection of pentamidine by at least one order of magnitude. This improvement was also seen in the detection of lipids and small drug molecules during tissue imaging.^[48] Patti et al. ^[43] replaced the organic matrix with AgNO₃ coating of the NIMS substrate. The study focused on localizing perturbations in metabolism within pathological tissues by MSI. It demonstrated that deposition of cationization agents (AgNO₃) to the NIMS surface allows imaging of otherwise difficult sterol molecules such as cholesterol. Lastly, NIMS was modified by redesigning the nanostructured silicon. For this, the highly dangerous electrochemical etching of silicon was replaced by dry-etching using plasma. Gao et al. [50] used black silicon instead of the conventional pSi. The relationship between black silicon morphology and its NIMS sensitivity was studied using several biomolecules: spermidine, arginine, adenosine, palmitoyl carnitine, verapamil, bradykinin, and STAL-2. It was found that the black silicon pillars absorb enough initiator promoting desorption of analytes, and that large surface areas can efficiently improve

NIMS sensitivity due to the enhanced energy transfer from substrates to analytes. The same dry-etching method was used to create an integrated microfluidics-NIMS device.^[52] In this investigation, the novel NIMS substrate is compatible with droplet and digital microfluidics and can be used on-chip to assay glycoside hydrolase enzyme in vitro.

2.1.3. Functionalization with metals

Alternative studies presented the use of metals to functionalize the pSi surface. Zhou et al.^[40] used several chemical reactions to obtain phosphonate-terminated pSi which was immersed into ZrOCl₂ solution to yield the Zirconium Phosphonate-Modified Porous Silicon (ZrP-pSi) wafers. The obtained ZrP-pSi wafers were sensitive to phosphopeptide detection and their high specificity was demonstrated by the analysis of tryptic digest product of α -casein, β -casein and BSA. Latest research specifically focused on the use of silver as surface functionalizing agent for pSi. Yan et al.^[41] used different mixtures of silver nitrate in water, 4-ATP in ethanol and trifluoroacetic acid (TFA) to prepare substrates with and without well-organized 4-ATP selfassembled monolayers. The study demonstrated that the 4-ATP capped substrate was more efficient than the substrate covered by naked Ag nanoparticles exhibiting higher ionization efficiency and less fragmentation. The substrate performance was excellent, with limits of detection down to several femtomoles for TPyP, sub picomoles for oxytocin, and picomoles for PEG 400 and PEG 2300. A different approach to obtain silver covered pSi was described by Gustafsson et al.^[42] The DIOS substrates were prepared by electrochemical etching and immediately sputter-coated with a 1.4nm-thick Ag layer. For testing, this new DIOS/Ag-DIOS MSI method, distributions for fingermark compounds and tissue metabolites of 6-bromoisatin were mapped. Fingermark analysis showed a broad range of small molecule classes, including environmental contaminants (ditallow dimethyl ammonium chloride, DTDMAC), lipids (FAs, TAGs), sterols and wax esters (WEs). The imprinted murine fore-stomach tissue analysis showed phosphatidylcholine head group (PC), 6,6'-dibromoindirubin (6,6'-DBI), fatty acids

(FAs), cholesterol and Ag cluster peaks. This novel approach exhibited minimal observable depletion and has the potential to simplify the analysis of multiple compound classes by using a single analytical platform to acquire multi-tiered spatial data sets. Gold was also used to functionalize pSi by Li et al.^[46] They used electrochemical deposition method to obtain a new pSi chip functionalized with gold nanoparticles (PSi-GNPs) that allows direct serum peptide profiling with a high-quality MS signal. Using both silver and gold nanoparticles is also possible; Wang et al.^[47] created patterned nanoporous silicon chips embedded with Ag and Au NPs by chemical assisted etching for detection of thiol compounds. In this case, the selectivity of the Ag and Au NPs embedded chips towards thiol compounds was monitored in cells. Also, the effect of irinotecan, 7-ethyl-10-[4-(1-piperidino)–1-piperidino]-carbonyloxy camptothecin (CPT-11), which is a potent anticancer drug, was determined using the novel chips.

2.1.4. Other functionalization methods

For specific applications that characterize bio-systems, functionalization of pSi implies the formation of particular Si- bonds on the surface of pSi. This method was used to enhance detection of slightly larger molecules such as peptides or proteins, together with the use of an organic matrix. Several research groups ^[27,28,34] developed such analytical methods. Chen et al. ^[27] analyzed a specific protein system, NTA-Ni2+/His-tagged protein, with carboxyl (–COOH) functionalized pSi microarray. The groups of Chen et al. ^[34] and Yan et al. ^[28] have immobilized antibodies on the pSi surface by physical adsorption. These experiments resulted in the detection of biomarker B-type natriuretic peptide (BNP) with a detection limit as low as 10pg/mL BNP in human plasma ^[34] and the specific detection of angiotensin I at a 10 fmol level in diluted plasma samples (10 μ L, 1 nM).^[28] Another specific biological system characterization method was developed by Sweetman et al. ^[45] where the attachment of mammalian cells to pSi was studied. For this, NHS ester and PEG-functionalized pSi surfaces were conjugated with

adhesion mediator protein fibronectin (FN) and used for the selective immobilization of human neuroblastoma cell line SK-N-SH.

3. Silicon nanostructures: nanoparticles, nanowires, and nanostructured surfaces

Nanostructures such as nanoparticles (NPs), nanowires (NWs), nano-flowers, nano-pillars etc., have been well studied for their applications in the biomedical area. Lately, silicon and silicacontaining NPs, silicon NWs and nanostructured silicon surfaces were demonstrated to be useful substrates for MS analysis of several types of biomolecules including small drug molecules, peptides, proteins, lipids, immobilized DNA. The mass spectrometry techniques using these substrates are known as surface-assisted laser desorption/ionization mass spectrometry (SALDI-MS) and nanostructure assisted laser desorption ionization mass spectrometry (NALDI-MS).^[13] As for pSi, silicon nanostructures can be used together with organic matrices for detecting larger molecules.

In this section, several nanostructured substrates will be described based on their synthesis method, functionalization method and use in LDI-MS analysis. The fabrication methods will be divided into three categories: nanoparticles and nano-powders (3.1.1), nanowires (3.1.2) and nanostructured surfaces (3.1.3). **Table 2** shows a resume of the several substrates, according to their fabrication method and target application. The surface modification methods will be divided into three categories: derivatization using silylating reagents (3.2.1), functionalization with metals (3.2.2) and finally, other functionalization methods (3.2.3).

3.1. Fabrication methods

3.1.1. Silicon/silica nanoparticles

Silicon and silica-containing nanoparticles have emerged as a reliable alternative to organic matrices for LDI ever since pSi demonstrated its extensive value in MS applications. As a result, many groups developed a variety of silica and silicon nanostructures. Silica nanoparticles can

be easily synthesized by wet chemical methods but they are available to be purchased from several commercial distributors as well. Dupre et al. chose to prepare SiO₂ nanoparticles through the very well-known sol-gel method. The group used a solution of ammonium hydroxide in methanol and tetraethoxysilane (TEOS) as the precursor for the silica NPs. After all the necessary steps were completed, the SiO₂ NPs were dried at room temperature.^[55] Following the example of SiO₂ NPs, core-shell nanoparticles (CSNPs) have also been used as matrix-free substrates for LDI-MS experiments. These nanostructures proved to be more complex and in need of a rigorous design for each application. The synthesis of such nanoparticles was carried out mainly using sol-gel processes. For example, magnetite core-shell particles were prepared by several groups ^[56–59], where the core was a magnetite nanoparticle and the shell was a thin silica layer. Briefly, the CSNPs were synthesized by coating the magnetite particles with a thin silica layer through the sol-gel process with TEOS as the silica source. Xiong et al.^[59] designed and synthesized aptamer-immobilized magnetic mesoporous silica/Au nanocomposites (MMANs) (Figure 2) for highly selective detection of unlabeled insulin in complex biological media using MALDI-TOF MS (with CHCA matrix). Briefly, the aptamer was anchored onto the gold nanoparticles in the mesochannels of MMANs for efficient and specific enrichment of insulin. Zhu et al. approached the CSNP method differently. The magnetite core was substituted by AuNPs (between 18 and 50 nm) and the outer layer was an ultrathin silica shell (~2-4 nm). The adhesion between gold and silica was possible due to the prior functionalization of the AuNPs' surface with amino groups.^[60] However, other groups ^{[61–} ^{63]} chose to purchase the SiNPs and experimented with storage and functionalization methods.

3.1.2. Silicon nanowires

Silicon nanowires (SiNWs) used in LDI-MS experiments have been primarily synthesized using vapor-liquid-solid (VLS) growth mechanism.^[64–66] The VLS method uses a nano-sized catalyst metal (deposited onto the silicon wafer) that can rapidly adsorb gaseous precursors

from which the growth of nanowires occurs. The growth and diameter of SiNWs are directly dependent on the Au colloid particles' size. In this case, the groups that developed SiNWs have used Au nanocluster catalyzed VLS growth mechanism. Briefly, Au nanoparticles are distributed on a silicon substrate by spin-coating, then solvents and organic residues are removed and finally the growth of SiNWs is obtained during chemical vapor deposition with silane as the vapor-phase reactant. Muck et al. ^[66] combined the VLS growth method with the typical fabrication methods of patterned silicon: photolithography, wet-chemical etching, and dry plasma etching. Firstly, the silicon substrates were patterned with holes (~50 mm depth) arranged in arrays of 2.5 mm circles with a 4.5 mm pitch (**Figure 3**). Lastly, the patterned silicon served as the substrate where the SiNWs were grown at the bottom of the etched holes by the VLS method using chemical vapor deposition from silane. SiNWs were commercially available for a limited time from Bruker.^[80]

3.1.3. Nanostructured silicon surfaces

Along with the use of silicon nanoparticles and nanowires, silicon nanostructured surfaces have also been tested as LDI-MS substrates.^[67–77,79,80] In most of the studies, the nanostructured surfaces are referred to as silicon nanowire arrays. These are densely packed arrays of vertical nanopillars, also called black silicon (BSi), not to be confused with silicon nanowires. The fabrication methods of these substrates can be categorized in two: wet etching and dry etching. The most popular fabrication method for silicon nanowire arrays is chemical etching (see **Figure 4**).^[67–70] Briefly, Si wafers are cleaned using well-known methods (Piranha solution, sonicating in acetone, isopropanol, methanol, rinsing with deionized water, etc.) to obtain clean and oxide-free Si surfaces, then the wafers are etched in a solution of AgNO₃/H₂O/HF of different concentrations to form bare silicon nanopillars. This classical method was slightly modified and replaced by metal-assisted chemical etching (MACE).^[71–74] In this case, a new step is added to the process. After Si wafer cleaning, a thin layer of Au is deposited onto Si

substrates and then immersed in an $HF/H_2O_2/EtOH$ etching mixture to create nanostructured silicon on the wafer.

However, all chemical etching methods use very toxic and dangerous substances for the user. For example, the piranha solution is a mixture of H_2SO_4/H_2O_2 , which is a strong oxidant that reacts violently with organic materials and can cause severe skin burns. Another dangerous substance used is HF, which is a hazardous acid that can cause serious tissue damage if burns are not appropriately treated. All etching processes must be handled with extreme care in a wellventilated fume hood while wearing appropriate chemical safety protection: face shield, double layered nitrile gloves, etc.^[61,67,71] All these factors can reduce the interest of groups in using these methods.

Alternative solutions, consisting of variations on dry etching methods used in creating nanostructures on silicon wafers have quickly emerged.^[75–79] Basically, the silicon wafers are etched using plasma in the presence of a protective mask. In the work of Gulbakan et al.^[75] an alumina mask was used together with argon plasma with various etching parameters to form different pore depth nanowell arrays. In the work of Wang et al.^[76], the mask was a polystyrene (PS) nanosphere monolayer and the plasma was formed by a mixture of etching gases: SF₆, CHF₃, and O₂. In this case, the resulting nanostructure was a biomimetic silicon nanocone array, with cone heights ranging between 200-600 nm. Morris et al. used traditional microelectronic fabrication steps for all the process: first they patterned the silicon wafer using deep-UV-photolithography technique and after they performed silicon etching using both ICP and RIE and a mixture of etchant gases: C₄F₈, SF₆, and Ar. This resulted in well-ordered silicon nanopost surfaces (NAPA) that were useful for small molecule analysis (Figure 4).^[77,78] The fabrication method of Chen et al.^[79] is slightly unconventional compared to the above-presented techniques. The silicon wafers were exposed to repeated laser irradiations in the presence of air, SF₆ gas or deionized water. This method however generated mostly microcolumn arrays but with special

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conditions (600 laser shots at 0.13 J/cm² in water environment) submicrometer microcolumns were generated with average 800 nm height and periodicity 600 nm.

3.2. Nanostructure derivatization and functionalization methods

All dry techniques presented in this review produced stable substrates without additional modifications, contrary to the surfaces produced by chemical attack. The latter were stabilized during derivatization or functionalization. All methods are summarized in **Table 2** below.

3.2.1. Derivatization using silylating reagents

The groups in ^[61,62] chose to silvlate purchased SiO₂ nanoparticles with PFP, PHP, TMS, C10, C6, and C3, and test the silicon-nanoparticle-assisted laser desorption/ionization (SPALDI) method on pharmaceuticals, peptides, pesticides, nucleic acids, and salt derivatives. The SiNPs were oxidized with HNO₃ before silvlation, and after silvlation they were mixed with the analyte solutions and spotted on standard MALDI plates. Nanowires have also been silvlated with PFPPDCS.^[64,65] In both cases the SiNWs were first etched in HF solution to remove the oxide layer and then reoxidized with ozone, followed by the final silvlation step. Go et al. ^[64] examined the effect of laser energy, nanowire density, nanowire size, and growth orientation of the SiNWs while performing mass spectrometry experiments on peptides and small drug molecules. Luo et al. focused on studying the internal energy transfer in SALDI from SiNWs and its relationship to nanoporous silicon in DIOS and to conventional MALDI.^[65] By means of benzyl-substituted benzylpyridinium thermometer ions, the authors in ^[65] could prove that very low laser fluence is needed for LDI, due to the high thermal energy confinement achieved in SiNWs. The chemically etched substrates ^[67–69] were modified with OTS, FDTS or ODCS. These studies were mostly focused on validating the use of the new substrates by detection of various types of peptides and small molecules. Lastly, a silvlated (with F₁₃) MACE substrate was used to detect peptides (middle range 1–3 kDa) and methadone (low range <500 Da). ^[72]

The substrate allowed for the detection of methadone in saliva, blood plasma and urine from clinical samples of real methadone-treated patients.

3.2.2. Functionalization with metals

Metallic nanoparticles such as gold and silver have previously demonstrated their efficiency in MS experiments ^[3,81], so functionalizing nanostructures substrates with metals is a compelling opportunity for enhancing LDI substrates. Au was used by various groups as a functionalization agent ^[59,63,73]. Xiong et al.^[59] modified magnetite core-shell nanoparticles with Au NPs on the silica outer shell obtaining aptamer-immobilized magnetic mesoporous silica/Au nanocomposites (MMANs). The main use of these composite nanoparticles is for insulin detection, however, more proteins were detected as well: immunoglobulin G, human serum albumin, α 1-antitrypsin, horseradish peroxidase, lysozyme, and cytochrome C. Another group ^[63], attached purchased silica nanoparticles to a silicon wafer using polymers for adhesion. Then the SiNPs were modified with Au by immersion of the wafer into an AuNP solution. This way the AuNPs constructed the outer layer of the substrate facilitating the formation of a selfassembled monolayer (SAM) of Capture DNA-1. The final substrate was an Au and SiO2 NPsassembled hybrid porous nanostructure and was used to analyze confined DNA structures. Another method to create AuNPs on nanostructured surfaces was developed by Tsao et al.^[73] The nanostructured Si surface was oxidized with O₂ plasma and immersed in HF/HAuCl₄ solution to graft the Au nanoparticles. The resulting AuNPs-nSi chip was used as LDI-MS substrate for the analysis of glucose from standard solutions and urine. High detection sensitivity and specificity for glucose in a biological sample confirmed the potential of the substrate.

Other metals used to create functional substrates were Cu ^[57], Li ^[66], and Ag ^[70]. Copper was used for the surface grafting of magnetite core-shell nanoparticles with Cu²⁺ ions. The functional microspheres could capture small peptides from complex sample systems (human

serum and urine) with the large proteins excluded because of the porosity of mesoporous silica and the specific affinity of Cu²⁺ ions toward peptides.^[57] Muck et al. chose Li as a doping agent. In this case, the sample solutions were deposited on the silicon nanowire substrate and the ⁷LiOH solution was deposited after the sample dried. The ⁷LiOH solution was prepared in methanol–dichloromethane 1:4 at 7.5 mg mL⁻¹. Most of the compounds were detected as clean [M+⁷Li]⁺ signals.^[66] Finally, AgNPs were used to decorate silicon nanowire arrays by a pulsed laser deposition method.^[70] The applicability of AgNP-decorated SiNWs was demonstrated by the detection of unsaturated compounds (SQ and oleic acid) in a complex matrix.

3.2.3. Other functionalization methods

Some groups chose to approach the functionalization method differently. The study by Chen et al.^[56] modified the surface of magnetite core-shell nanoparticles (CSNPs) first by modification with MPS and finally by a seeded aqueous-phase radical polymerization with MMA. These CSNPs were specifically designed for the enrichment of peptides and proteins for mass spectrometric analysis. Other groups have also designed their novel substrate based on magnetite CSNPs roughly for the same purpose: selective enrichment of peptides.^[58] In this case, the microspheres were dried and calcined in nitrogen to improve the hydrophobic property of the surface. A different study developed CSNPs with AuNP core and ultra-thin silica shell.^[60] The stabilization of these Au@utSiO2 CSNPs was carried out by further growth of the silica shell using the Stöber method. Small functional molecules and small polymers were successfully detected by CSNP-based LDI-TOF-MS. In 2008 Daniels et al. ^[80] functionalized commercially available NALDI target plates from Bruker Daltonics, Billerica, MA. The surface of the silicon nanowires was modified by depositing organic layers to change the hydrophobicity of the substrate. The new target plate was used for the detection of small molecules. Unfortunately, there is no mention about the organic layer type and the NALDI plates are no longer available for purchase.

4. LDI mechanism: ionization and desorption processes

Matrix-assisted laser desorption/ionization mass spectrometry (MALDI-MS) has become an indispensable analysis technique for many research communities since its first use by Karas et al. in the 80s.^[82] The fundamental strategy of a complete MALDI-MS analysis consists of several steps: sample preparation, excitation of sample and desorption of the condensed phase, ionization of analyte molecules by the generation of charges, extraction, and detection of ions.^[83] The processes that occur between the excitation of the sample and the extraction of ions are collectively called laser desorption/ionization (LDI) mechanism. This mechanism is most commonly considered as a convolution of two processes: desorption and ion formation. These processes have been previously studied in detail by several groups.^[83–87] LDI is portrayed as a complex process that involves both optical and mechanical phenomena as well as thermodynamic and physicochemical processes of phase transition and ionization. Ionization, in particular, was described as a collection of chemical and physical pathways including gasphase photoionization, ion-molecule reactions, disproportionation, excited-state proton transfer, energy pooling, thermal ionization, and desorption of preformed ions.^[84] The LDI mechanism has been an active topic of research and was revisited by Chang et al. and Jaskolla et al. ^[86,87]. Since the first years of MALDI analysis, ionization of analytes was helped by an organic matrix that possesses some essential functions. First, it has to isolate analyte molecules by preventing analyte aggregation. Then, it has to absorb the laser energy while the disintegration of the condensed phase takes place without excessive destructive heating of the analyte molecules. Lastly, the matrix should provide an efficient ionization of analyte molecules.^[84] With all these specificities of the matrices, however, the generated spectra are far from ideal and the use of a matrix is not as suitable for untargeted MS experiments focused in small molecules. The matrix complicates the sample preparation, introduces background ions in the spectra, hampers quantitative analysis and also complicates the acquisition of MS images at high lateral resolutions. These difficulties may be reduced by replacing the organic matrix with a less

invasive component that has at least the same specific/required properties as the matrix. Therefore, by understanding the LDI mechanism of molecules adsorbed to a specific substrate, especially nanostructured silicon, one could be able to maximize ion yields, control analyte charge states and fragmentation, and gain access to analyzing new classes of compounds.

This section provides an overview of the LDI mechanisms described in LDI-MS experiments using silicon-based substrates instead of organic matrices. The specific physicochemical properties of the substrate necessary for aiding desorption and ion formation will be discussed, as well as the specific LDI processes for each substrate type.

4.1. Influence of physical properties

4.1.1. Porous silicon

For pSi, the main influencing factors of the LDI process are considered to be the physical properties of the material surface. The pSi structure has a high surface area which provides an optimal environment for the co-adsorption or entrapment of the analyte and solvent.^[23,24,29,53] High pore density and smaller pore sizes might produce better results as the increase in ion signal can be correlated with the increase of the overall surface area and analyte coverage. Furthermore, the tendency of energy localization near defects, protrusions, and edge sites makes these areas more active in LDI processes.^[23,53] In addition, analyte penetration into the pSi is critical because the pSi also manifests quantum confinement effects: large optical absorption and low thermal conductivity. These effects cause rapid heating of pSi in the presence of laser and the resulted energy may be transferred from silicon to the trapped analyte. Basically, instant heating of the pSi provides the necessary energy for analyte desorption and ionization.^[23,29]

4.1.2. Nanostructured silicon

Silicon nanostructures such as nanoparticles, nanowires, and nanostructured surfaces have specific physical properties that favor LDI processes. Although many different nanostructures have been fabricated, their properties are common for most of them: specific dimensions that

maximize laser light absorption, specific heat capacity and conductivity of the nanostructured silicon and geometric effects that promote molecule adhesion and desorption.

For silicon nanowires, the main features correlated with efficient LDI are morphology and low thermal conductivity. For example, the ionization performance of nanowires was described as strongly dependent on wire length and density.^[64] Here, SiNWs act as tiny antennas where the laser energy is efficiently absorbed. When the energy is focused on a small area it generates a field desorption effect resulting in the gas-phase generation of the deposited analyte molecules. This energy focusing effect primarily promotes desorption of molecules and results in very little surface-related background ions.^[64] The SiNW arrays showed superior laser desorption properties by requiring lower laser energy to desorb molecules.^[66] In this case, the obtained signal was enhanced due to three main reasons. Firstly, because of the increased absorption of the nanowire forest near the wavelength of the laser, then because of the large surface area of the nanostructures.^[66] The group of Vertes et al. ^[65] has committed to thoroughly study the internal energy transfer in laser desorption/ionization from silicon nanowires. They also refer to geometry and thermal properties of the SiNWs as main influencing factors on the more efficient energy transfer resulting in a more efficient desorption.

In the case of nanostructured silicon surfaces, the key physical factors that promote efficient ionization are the structure's morphology and its optical and thermal properties. For example, the dimensions of the nanostructured silicon surfaces described by several groups ^[70,72,76,77] were directly associated with ionization efficiency. In these cases the ion intensity decreased considerably with length increase, the longer structures (> 450 nm ^[72]) did not allow efficient energy transfer from the laser to the analyte. Morris et al. ^[77] correlated three main features of their nanopost arrays with optimum LDI, namely: (1) well-ordered nanopost arrays, (2) dimensions that maximize laser light absorption and subsequent resonance effects that promote analyte desorption and ionization, and (3) a highly porous surface to maximize analytical

sensitivity. Gulbakan et al. ^[75] have also suggested that these factors generate a local environment to which analytes adhere with optimal heat capacity and heat conductivity that result in effective absorption of laser light followed by effective sublimation.

Another morphological aspect that was associated with increased ionization efficiency was surface roughness. It was implied that the surface roughness has a more direct effect on the ion intensity efficiency than the silicon nanostructure pore size, depth and surface area/volume ratio.^[71]

4.2. Influence of functional group

4.2.1. Porous silicon

Another factor that affects the LDI mechanism of pSi is the chemical property of the surface. Hydrophobic or hydrophilic groups may cover the substrate surface and affect the analyte adsorption on the pSi. Hydrophobic surfaces may give intense MS signals for hydrophobic analytes from aqueous medium because the solvent is not confined to the porous area.^[23] On the contrary, hydrophilic surfaces enhance adsorption of all molecules from an aqueous medium onto the pSi and result in lower ion desorption efficiency.^[44] In the case of derivatized substrates with terminal hydride and silanol groups, deposition of the analyte leads to the adsorption and trapping of the solvent.^[53] Another example is the interaction between the analyte and metallic cations. Ag is a well-known cationization agent that produces enhanced D/I processes.^[42] LDI of analyte molecules using Ag nanoparticles is obtained through high absorption and low reflection of energy. The laser energy is quickly absorbed by the Ag nanoparticles which are rapidly heated, resulting in the vaporization and ionization of the analyte molecules.^[41] In the same study, the surface was functionalized with 4-ATP as well, which acted as a matrix. This small aromatic molecule enhanced the absorption of energy in the ultraviolet region. Gold is another well-known metal used to functionalize pSi surfaces for enhanced ionization.^[46,47] Li et al. ^[46] describe the effect on ionization of the plasmonic property of AuNPs. They assume that AuNPs may act primarily as antennae which concentrate the laser-induced field within the

nanoporous channel, which leads to an enhanced MS signal. They also postulate that plasmonic metals absorb resonant photons, and the energetic electrons formed by the SPR excitation are transferred to the semiconductor (Si), leading to the accumulation of positive charges on the metal surface.

4.2.2. Nanostructured silicon

Chemically derivatized silicon nanostructures have been designed and synthesized to improve the LDI efficiency mainly by separating targeted molecules from biological media and inducing a controlled ionization of the selected analytes. In case of the group of Liu et al. ^[57], this process consists in functionalizing the magnetic mesoporous microspheres with immobilized Cu²⁺ for a high efficiency in peptide enrichment. Another example is for the aptamer functionalized magnetic nanoparticles. These were used to enhance the MS intensity of insulin, where aptamer and magnetism facilitate immunoreactions between the aptamer and the target, improving detection sensitivity.^[59]

A more traditional chemical improvement of nanostructured silicon was reported by Morris et al.^[77] In this study, the silanol groups are used as proton source for ionization and to induce increased analyte sensitivity. They described the presence and electronegativity of fluorine as a factor that increased the acidity of the silanol group, thus providing a favorable environment for the protonation of molecules. The altered chemical composition of the silicon nanostructures also affects optical absorption. In the case of Chen et al. the absorption values of microcolumns are increased across the entire UV-visible spectrum, and significant absorption extends into the near-IR region. ^[79]

4.3. Influence of other interactions

4.3.1. Porous silicon

Proton affinity of certain analytes has been also speculated to affect ionization.^[33,44,53] The ionization efficiency is similar for analytes with comparable secondary amino groups (e.g.

structurally related MA and MDMA) while the ionization efficiency is higher for other analytes (e.g. cocaine), which contain tertiary amino groups, that have higher proton affinity.^[33] Functional groups and solvents are the primary proton sources, so the right combination between the substrate, the analyte, and the solvent could be critical for optimal DIOS efficiency. External elements, such as laser fluence and substrate storage, impact the D/I process as well. High laser fluence results in analyte fragmentations that could lead to crowded mass spectra in the low mass range. This is sheer evidence of the thermal driving force in the D/I process.

4.3.2. Nanostructured silicon

Some LDI mechanisms are not directly affected by the physicochemical properties of the substrate but by the inherent physicochemical properties of analytes such as pI (isoelectric point), hydrophobicity, hydrophilicity, number of charges and their concentration. For example, due to each molecule's specific pI and due to the sample pH, some molecules appear as negatively charged ions and cannot be detected in positive mode and vice versa. One example of negatively charged molecule is fibrinopeptide B, due to the presence of three glutamic acid and one aspartic acid residues in its sequence.^[68] In this study, the super-hydrophilic pattern of the surface was investigated as well. This specific surface promoted a better ionization efficiency of peptides because most peptides are positively charged and adsorb specifically to the SiO₂ surface. This specific adhesion property of biological samples has been attributed to amino acid residues that have different adhesion properties to inorganic interfaces such as SiO₂, Si₃N₄, and metals, essentially depending on their side chains.^[68]

Other external factors that affect the LDI mechanism are laser energy and laser plume. So far laser energy has been discussed as UV light efficiently absorbed by the silicon nanostructures. Laser energy can be varied to induce different effects on the D/I processes. Zenobi and Knochenmuss ^[84] described in great detail ion formation and the effect of laser fluency in the classical MALDI process. A similar description could be given for nanostructure assisted LDI.

However, several authors imply that the optimal laser intensity required to get good MS signals is typically much lower than that required for both standard MALDI and DIOS.^[88–90] This is assumed to be due to the optical and thermal properties of silicon nanostructures. The low laser fluencies are also beneficial in the ionization of thermally labile compounds because it may reduce the necessary energy for desorption and it may also reduce fragmentation.^[79]

The laser plume effect has been described as the presence of electrons in the laser plume due to photoelectric effects. Ions can be generated through electron impact ionization or recombination between electrons emitted from the plume and analyte surface. This way the desorbed species mix with the plume and after ion-molecule reactions protonated species are generated. At increased laser power, the elevated electron density can neutralize the protons to form hydrogen-free radicals, meanwhile, more alkali ions are released from the hot silicon surface that results in peptide-alkali adduct ion formation.^[79]

Another interesting hypothesis suggested that absorbed UV laser energy was transferred from the surface to pre-charged analytes causing desorption.^[79] However, it is unclear whether the protonation process happens in solution (before laser interaction) or it is a laser-induced proton transfer on/near the surface. Also, residual solvents retained in the cavities of the nanostructured surfaces are probable sources of protons that help to ionize analytes.^[79] Tsao et al. proposed another ionization process: glucose samples were catalyzed to negatively charged gluconic acid molecules by the on-chip AuNPs-nSi surface Au- based catalyst reactions.^[73] In this case, enhanced detection sensitivity was a result of the Au nanoparticles grafted to the nanostructured silicon surface.

5. Discussion

Silicon is one of the prime candidates for fabrication of LDI-MS substrates for three main reasons: silicon is biologically inert; the silicon technology is very advanced and flexible, and silicon nanostructures can be designed with controllable morphology and properties for LDI-

MS. It is important to mention that silicon is a material that can be easily stabilized, derivatized and functionalized.

In case of silicon substrates designed for LDI-MS experiments, the wet (chemical) fabrication techniques used to create pSi slowly evolved to dry (physical) techniques. Wet techniques use highly toxic compounds (ex.: piranha solution for cleaning and HF for etching) for dangerous fabrication methods (ex.: electrochemical etching in a closed Teflon cell). Therefore, dry fabrication techniques have emerged that eliminate all dangers to the user. These techniques also improve the essential needs of a good quality substrate: fabrication process is highly repetitive, the surface is free of air or liquid contaminants, automated fabrication is possible and most importantly, no danger for the user. Most of the dry fabrication techniques are derived from the micro/nanoelectronics fabrication methods. This is another reason why dry fabrication methods are becoming more popular: silicon technology is improving day by day and together with it, silicon nanostructure fabrication methods keep advancing.

Silicon nanostructures of all types: nanoparticles, nanowires, black silicon, pSi, etc., have been stabilized and derivatized using different strategies for many purposes. There is no standard procedure for all types of experiments. Muthu et al. ^[89] suggested that the sample preparation is an art form, and, given its complexity, surface modification of any silicon nanostructure can also be considered an art form. Surfaces can be modified with a variety of agents: from metallic nanoparticles (Ag, Au, Cu, etc.) to small (amine groups, perfluorophenyls, etc.) and big (neat silane, antibodies, etc.) biomolecules (Table 1 and 2). Depending on the application of the substrate, one can design a targeted analysis method by choosing the appropriate functionalization method and/or combine various functionalization methods to reduce unwanted signal or to obtain a broader range of detection. It all depends on the users' creativity. All these surface-modified matrix-free methods can be used together with an organic matrix to expand the detection range from small to large molecules.

Although silicon nanostructures can be very repetitive and homogeneous, the reliability of the LDI-MS measurements depends also on the sample deposition method. Irreproducible sample deposition leads to irreproducible results. This is strongly related to the uneven crystallization of liquid samples on the substrate which induces signal variation between different spots of the same sample. For this reason, sample deposition was rigorously studied for silicon nanostructures. In the case of pSi, reproducible liquid sample deposition was achieved through acoustic printing.^[50,91]

As we have described above, silicon-based nanoparticles/nanowires have been used as LDI substrates. In this case, the sample analyte is usually mixed with the nanoparticle/nanowire solution in the appropriate concentrations prior to spotting on the MALDI target plate and subsequent co-crystallization.^[60,61] Although this wet chemistry deposition technique is very flexible, it suffers from the same problems as in the case of the organic matrices (i.e. heterogeneous spots).

Most of the studies mentioned in this report focus on liquid samples, however, the analysis of tissues is highly demanded as well. The first issue to be considered in an MSI experiment is the performance of the substrate-tissue-air system. In current standard MALDI analysis, the tissue is cut into ~10 μ m thick slices and each slice is mounted onto an indium tin oxide (ITO) coated glass slide, then coated with a thin layer of matrix.^[8] In this case, the thickness of the tissue does not affect the transfer of energy from the laser to the matrix and implicitly the tissue for LDI processes. In other cases, when the matrix and the ITO glass slide are replaced by a nanostructured surface on which the tissue is mounted, the thickness of the tissue matters. The laser energy has to pass the tissue and reach the nanostructured surface in order to promote the ion desorption and ionization processes. In NIMS applications, in order to facilitate the laser to get to the tissue-nanostructure interface, the tissue slices are 3-5 μ m thick, so they are much thinner than in usual MSI experiments.^[43,48,49] Even in this case, a high laser power is needed which mostly results in burning the tissue and produces harder ionization. Moreover, NIMS

MSI experiments have not been reported on Bruker MALDI-MS spectrometers, which suggests that there is some instrumental limitation with this manufacturer.

An alternative technique that avoids the effect of tissue thickness, consists of imprinting (or stamping) a tissue over the surface.^[36] The imprinting process consists in putting the tissue in contact with the nanostructured substrate and then removing it, leaving only the adhered molecules from the tissue sample onto the substrate. Although it avoids the cutting of thin tissue samples and depositing them over the surface, the imprinting processes have been reported to lack reliability because of possible smears.^[92] Within this method, new applications of MSI in the clinical practice are envisaged, consisting in the imprint of a tissue on a silicon-based substrate for needle biopsy processes, without the need to remove the biopsy tissue.^[93]

The imprinting process is compatible with the new strategies developed to create specific adhesion of molecules to the nanostructured surface through surface functionalization. The silicon-based nanoparticle methods presented in this review often use linker molecules for specific detection of analytes. This approach should be considered for nanostructured silicon surfaces as well.

Guidelines for detecting biomolecules by Si LDI-MS

All detected molecules and detection strategies used in the publications presented in this progress report are listed in **Table S3** and summarized in **Figure 5**. The bar charts in **Figure 5** show the great versatility of Si based substrates, as all types of Si substrates have been used to detect most types of biomolecules. Since there are no standard protocols for detecting all kinds of molecules simultaneously, each strategy is focused on detecting specific biomolecules. As such, we highlight the following trends in Si LDI-MS:

A. Peptides and proteins are often detected with the help of organic matrices while small molecules do not need the organic matrix to be detected. Larger biomolecules are known to be thermally labile and they need the controlled energy transfer from an organic matrix to be analyzed by the laser desorption method.

- B. Most of the pSi, SiNW and SiNP surfaces are modified, while half of the published work on nano Si does not use surface modification. This suggests that dry fabrication techniques produce stable surfaces, while wet chemistry fabrication techniques produce surfaces that need stabilization.
- C. Si LDI-MS has first appeared as DIOS (pSi), then evolved into NIMS (pSi) and lately it is appearing as nanostructured silicon LDI. The use of matrix is reflected on the evolution of Si LDI-MS as organic matrices were used together with pSi and SiNP but are less and less used together with newer nanostructured Si substrates. We envision that matrix-free strategies will be the LDI strategy of choice in the future.

6. Conclusion

The extensive variety of silicon nanostructures used for MS applications draws attention to the major prerequisites for a successful LDI-MS analysis. For this, we conclude that the basic requirements for any silicon-based substrates are: (1) to absorb UV laser irradiation without deteriorate from it, (2) to provide a surface for easy functionalization, (3) to be stable under vacuum, (4) to improve and enhance the analyte ionizability, (5) to not cause interferences or clusters with the target analyte, (6) to be easy to fabricate, (7) to allow reproducible results and finally, (8) to be cheap. Taking into consideration that silicon technology (regardless of the field of study) is constantly advancing, fabrication and functionalization of silicon-based substrates will rapidly evolve and have a great impact in the mass spectrometry imaging field.

Supporting Information

Supporting Information is available from the Wiley Online Library or from the author.

Acknowledgements

The authors want to acknowledge the financial support of the Spanish Ministry of Economy and Competitivity (MINECO) through projects TEC2015-69076-P and S. A. Iakab's predoctoral grant No. BES-2016-076483, the Spanish Ministry of Science, Innovation and Universities (MICIN) through projects RTI2018-096061-B-I00 and M. Garcia-Altares' postdoctoral grant IJCI-2017-33438, and the 2017-SGR-1119 grant awarded recently by the Agency for Management of University and Research Grants of the Generalitat de Catalunya (AGAUR).

> Received: ((will be filled in by the editorial staff)) Revised: ((will be filled in by the editorial staff)) Published online: ((will be filled in by the editorial staff))

References

- [1] D. S. Cornett, M. L. Reyzer, P. Chaurand, R. M. Caprioli, *Nat. Methods* 2007, *4*, 828.
- [2] H. N. Abdelhamid, TrAC Trends Anal. Chem. 2017, 89, 68.
- [3] L. A. McDonnell, R. M. A. Heeren, *Mass Spectrom. Rev.* 2007, 26, 606.
- [4] J. L. Norris, R. M. Caprioli, *PROTEOMICS Clin. Appl.* **2013**, *7*, 733.
- [5] C.-K. Chiang, W.-T. Chen, H.-T. Chang, *Chem. Soc. Rev.* 2011, 40, 1269.
- [6] C. Gregson, *Biosci. Horizons* **2009**, *2*, 134.
- [7] E. Gemperline, S. Rawson, L. Li, Anal. Chem. 2014, 86, 10030.
- [8] N. Ogrinc Potočnik, T. Porta, M. Becker, R. M. A. Heeren, S. R. Ellis, *Rapid Commun. Mass Spectrom.* 2015, 29, 2195.
- [9] D. S. Peterson, *Mass Spectrom. Rev.* **2007**, *26*, 19.
- [10] L. Qiao, B. Liu, H. H. Girault, *Nanomedicine* **2010**, *5*, 1641.
- [11] C. Y. Shi, C. H. Deng, Analyst 2016, 141, 2816.
- [12] K. P. Law, J. R. Larkin, Anal. Bioanal. Chem. 2011, 399, 2597.
- [13] Y. E. Silina, D. A. Volmer, Analyst 2013, 138, 7053.
- [14] J. Sekuła, J. Nizioł, W. Rode, T. Ruman, Anal. Chim. Acta 2015, 875, 61.
- [15] M. Dufresne, A. Thomas, J. Breault-Turcot, J.-F. Masson, P. Chaurand, *Anal. Chem.***2013**, 85, 3318.
- [16] H. Kawasaki, T. Ozawa, H. Hisatomi, R. Arakawa, *Rapid Commun. Mass Spectrom.***2012**, *26*, 1849.
- [17] Y. Gholipour, S. L. Giudicessi, H. Nonami, R. Erra-Balsells, *Anal. Chem.* 2010, 82, 5518.
- [18] A. Y. Lim, J. Ma, Y. C. F. Boey, Adv. Mater. 2012, 24, 4211.
- [19] Y. He, C. Fan, S. T. Lee, *Nano Today* **2010**, *5*, 282.
- [20] F. Peng, Y. Su, Y. Zhong, C. Fan, S. T. Lee, Y. He, Acc. Chem. Res. 2014, 47, 612.
- [21] J. J. Thomas, Z. Shen, J. E. Crowell, M. G. Finn, G. Siuzdak, Proc. Natl. Acad. Sci. U.

S. A. **2001**, *98*, 4932.

- [22] T. R. Northen, O. Yanes, M. T. Northen, D. Marrinucci, W. Uritboonthai, J. Apon, S. L. Golledge, A. Nordström, G. Siuzdak, *Nature* 2007, 449, 1033.
- [23] J. Wei, J. M. Buriak, G. Siuzdak, *Nature* **1999**, *399*, 243.
- [24] N. Budimir, J. C. Blais, F. Fournier, J. C. Tabet, *Rapid Commun. Mass Spectrom.*2006, 20, 680.
- [25] A. G. Cullis, L. T. Canham, P. D. J. Calcott, J. Appl. Phys. 1997, 82, 909.
- [26] T. Guinan, C. Della Vedova, H. Kobus, N. H. Voelcker, *Chem. Commun.* 2015, *51*, 6088.
- [27] L. Chen, Z.-T. Chen, J. Wang, S.-J. Xiao, Z.-H. Lu, Z.-Z. Gu, L. Kang, J. Chen, P.-H.
 Wu, Y.-C. Tang, J.-N. Liu, *Lab Chip* 2009, *9*, 756.
- [28] H. Yan, A. Ahmad-Tajudin, M. Bengtsson, S. Xiao, T. Laurell, S. Ekstr??m, Anal. Chem. 2011, 83, 4942.
- [29] J. Li, C. Lu, X. K. Hu, X. Yang, A. V. Loboda, R. H. Lipson, *Int. J. Mass Spectrom*.
 2009, 285, 137.
- [30] X. Yongsheng, T. R. Scott, K. T. Darrell, T. Jia-Yuan, H. Lin, J. Phys. Chem. C 2009, 113, 3076.
- [31] M. Gaspari, M. M. Cheng, R. Terracciano, X. Liu, A. J. Nijdam, L. Vaccari, E.
 Fabrizio, E. F. Petricoin, L. A. Liotta, G. Cuda, S. Venuta, M. Ferrari, *J. Proteome Res.*2006, *5*, 1261.
- [32] Y. Goto, N. Mizoshita, Y. Yamada, Y. Maegawa, J. Amano, S. Inagaki, *Microporous Mesoporous Mater.* 2018, 268, 125.
- [33] T. Guinan, M. Ronci, H. Kobus, N. H. Voelcker, *Talanta* **2012**, *99*, 791.
- [34] Y.-Q. Chen, F. Bi, S.-Q. Wang, S.-J. Xiao, J.-N. Liu, J. Chromatogr. B 2008, 875, 502.

[35] S. a Trauger, E. P. Go, Z. Shen, J. V Apon, B. J. Compton, E. S. P. Bouvier, M. G.Finn, G. Siuzdak, *Anal. Chem.* 2004, 76, 4484.

- [36] M. Ronci, D. Rudd, T. Guinan, K. Benkendorff, N. H. Voelcker, *Anal. Chem.* 2012, 84, 8996.
- [37] D. Rudd, M. Ronci, M. R. Johnston, T. Guinan, N. H. Voelcker, K. Benkendorff, *Sci. Rep.* 2015, *5*, 1.
- [38] T. M. Guinan, D. Neldner, P. Stockham, H. Kobus, C. B. Della Vedova, N. H. Voelcker, *Drug Test. Anal.* 2017, 9, 769.
- [39] T. Guinan, M. Ronci, R. Vasani, H. Kobus, N. H. Voelcker, *Talanta* 2015, 132, 494.
- [40] H. Zhou, S. Xu, M. Ye, S. Feng, C. Pan, X. Jiang, X. Li, G. Han, Y. Fu, H. Zou, J. Proteome Res. 2006, 5, 2431.
- [41] H. Yan, N. Xu, W.-Y. Huang, H.-M. Han, S.-J. Xiao, *Int. J. Mass Spectrom.* 2009, 281,
 1.
- [42] O. J. R. Gustafsson, T. M. Guinan, D. Rudd, H. Kobus, K. Benkendorff, N. H. Voelcker, *Rapid Commun. Mass Spectrom.* 2017, *31*, 991.
- [43] G. J. Patti, L. P. Shriver, C. A. Wassif, H. K. Woo, W. Uritboonthai, J. Apon, M. Manchester, F. D. Porter, G. Siuzdak, *Neuroscience* 2010, *170*, 858.
- [44] S. Tuomikoski, K. Huikko, K. Grigoras, P. Ostman, R. Kostiainen, M. Baumann, J. Abian, T. Kotiaho, S. Franssila, *Lab Chip* 2002, *2*, 247.
- [45] M. J. Sweetman, M. Ronci, S. R. Ghaemi, J. E. Craig, N. H. Voelcker, Adv. Funct. Mater. 2012, 22, 1158.
- [46] X. Li, J. Tan, J. Yu, J. Feng, A. Pan, S. Zheng, J. Wu, Anal. Chim. Acta 2014, 849, 27.
- [47] J. Wang, M. Jie, H. Li, L. Lin, Z. He, S. Wang, J. M. Lin, *Talanta* 2017, 168, 222.
- [48] T. N. Moening, V. L. Brown, L. He, Anal. Methods 2016, 8, 8234.
- [49] G. J. Patti, H. K. Woo, O. Yanes, L. Shriver, D. Thomas, W. Uritboonthai, J. V Apon,R. Steenwyk, M. Manchester, G. Siuzdak, *Anal. Chem.* 2009, 82, 4271.
- [50] J. Gao, M. De Raad, B. P. Bowen, R. N. Zuckermann, T. R. Northen, *Anal. Chem.***2016**, 88, 1625.

- [51] J. Gao, K. B. Louie, P. Steinke, B. P. Bowen, M. De Raad, R. N. Zuckermann, G. Siuzdak, T. R. Northen, *Anal. Chem.* 2017, 89, 6521.
- [52] J. Heinemann, K. Deng, S. C. C. Shih, J. Gao, P. D. Adams, A. K. Singh, T. R. Northen, *Lab Chip* **2017**, , 323.
- [53] Y. Chen, H. Chen, A. Aleksandrov, T. M. Orlando, J. Phys. Chem. C 2008, 112, 6953.
- [54] H. K. Woo, T. R. Northen, O. Yanes, G. Siuzdak, Nat. Protoc. 2008, 3, 1341.
- [55] M. Dupré, S. Cantel, J. O. Durand, J. Martinez, C. Enjalbal, *Anal. Chim. Acta* 2012, 741, 47.
- [56] H. Chen, C. Deng, X. Zhang, Angew. Chemie Int. Ed. 2010, 49, 607.
- [57] S. Liu, H. Chen, X. Lu, C. Deng, X. Zhang, P. Yang, Angew. Chemie Int. Ed. 2010, 49, 7557.
- [58] H. Chen, S. Liu, H. Yang, Y. Mao, C. Deng, X. Zhang, P. Yang, *Proteomics* 2010, 10, 930.
- [59] Y. Xiong, C. Deng, X. Zhang, P. Yang, ACS Appl. Mater. Interfaces 2015, 7, 8451.
- [60] X. Zhu, L. Wu, D. C. Mungra, S. Xia, J. Zhu, Analyst 2012, 137, 2454.
- [61] X. Wen, S. Dagan, V. H. Wysocki, Anal. Chem. 2007, 79, 434.
- [62] S. Dagan, Y. Hua, D. J. Boday, A. Somogyi, R. J. Wysocki, V. H. Wysocki, Int. J. Mass Spectrom. 2009, 283, 200.
- [63] M. Hong, X. Zhou, J. Li, Y. Tian, J. Zhu, Anal. Chem. 2009, 81, 8839.
- [64] E. P. Go, J. V. Apon, G. Luo, A. Saghatelian, R. H. Daniels, V. Sahi, R. Dubrow, B. F. Cravatt, A. Vertes, G. Siuzdak, *Anal. Chem.* 2005, 77, 1641.
- [65] G. Luo, Y. Chen, H. Daniels, R. Dubrow, A. Vertes, J. Phys. Chem. B 2006, 110, 13381.
- [66] A. Muck, T. Stelzner, U. Hübner, S. Christiansen, A. Svatos, Lab Chip 2010, 10, 320.
- [67] G. Piret, H. Drobecq, Y. Coffinier, O. Melnyk, R. Boukherroub, *Langmuir* 2010, 26, 1354.

- [68] F. Lapierre, G. Piret, H. Drobecq, O. Melnyk, Y. Coffinier, V. Thomy, R. Boukherroub, *Lab Chip* 2011, *11*, 1620.
- [69] M. Dupré, Y. Coffinier, R. Boukherroub, S. Cantel, J. Martinez, C. Enjalbal, J. Proteomics 2012, 75, 1973.
- [70] R. A. Picca, C. D. Calvano, M. J. Lo Faro, B. Fazio, S. Trusso, P. M. Ossi, F. Neri, C. D'Andrea, A. Irrera, N. Cioffi, *J. Mass Spectrom.* 2016, 849.
- [71] W. Y. Chen, J. T. Huang, Y. C. Cheng, C. Chien, C. W. Tsao, *Anal. Chim. Acta* 2011, 687, 97.
- [72] H. Z. Alhmoud, T. M. Guinan, R. Elnathan, H. Kobus, N. H. Voelcker, *Analyst* 2014, 139, 5999.
- [73] C. W. Tsao, Z. J. Yang, ACS Appl. Mater. Interfaces 2015, 7, 22630.
- [74] P. Y. Chen, C. Y. Hsieh, C. J. Shih, Y. J. Lin, C. W. Tsao, Y. L. Yang, J. Nat. Prod.
 2018, 81, 1527.
- [75] B. Gulbakan, D. Park, M. Kang, K. Kececi, C. R. Martin, D. H. Powell, W. Tan, *Anal. Chem.* 2010, 82, 7566.
- [76] Y. Wang, Z. Zeng, J. Li, L. Chi, X. Guo, N. Lu, J. Am. Soc. Mass Spectrom. 2013, 24, 66.
- [77] N. J. Morris, H. Anderson, B. Thibeault, A. Vertes, M. J. Powell, T. T. Razunguzwa, *RSC Adv.* 2015, *5*, 72051.
- [78] A. R. Korte, N. J. Morris, A. Vertes, Anal. Chem. 2019, 91, 3951.
- [79] Y. Chen, A. Vertes, Anal. Chem. 2006, 78, 5835.
- [80] R. Daniels, S. Dikler, E. Li, C. Stacey, J. Assoc. Lab. Autom. 2008, 13, 314.
- [81] K. Chughtai, R. M. A. Heeren, Chem. Rev. 2011, 110, 3237.
- [82] M. Karas, D. Bachmann, F. Hillenkamp, Anal. Chem. 1985, 57, 2935.
- [83] K. Dreisewerd, The desorption process in MALDI; 2003; Vol. 103.
- [84] R. Zenobi, R. Knochenmuss, *Mass Spectrom. Rev.* 1998, 17, 337.

- [85] M. Karas, R. Krüger, Chem. Rev. 2003, 103, 427.
- [86] W. C. Chang, L. C. L. Huang, Y. S. Wang, W. P. Peng, H. C. Chang, N. Y. Hsu, W.
 Bin Yang, C. H. Chen, *Anal. Chim. Acta* 2007, 582, 1.
- [87] T. W. Jaskolla, M. Karas, J. Am. Soc. Mass Spectrom. 2011, 22, 976.
- [88] J. A. Stolee, B. N. Walker, V. Zorba, R. E. Russo, A. Vertes, *Phys. Chem. Chem. Phys.* 2012, 14, 8453.
- [89] M. Muthu, S. Chun, H.-F. Wu, M. W. Duncan, J. Gopal, J. Mass Spectrom. 2018, 525.
- [90] R. A. Picca, C. D. Calvano, N. Cioffi, F. Palmisano, Nanomaterials 2017, 7, 75.
- [91] J. L. Norris, R. M. Caprioli, *Chem Rev.* **2013**, *113*, 2309.
- [92] Y. J. Lee, D. C. Perdian, Z. Song, E. S. Yeung, B. J. Nikolau, *Plant J.* **2012**, *70*, 81.
- [93] J. Kriegsmann, M. Kriegsmann, R. Casadonte, Int. J. Oncol. 2015, 46, 893.



Figure 1. SEM of a DIOS chip in (A) the top view and (B) the cross-sectional view. (C) The schematic of pSi functionalized with F5PhPr and (D) DIOS- MSI of fingerprint sweat at 200 μ m resolution. Reprinted with permission ^[26], Copyright 2015 © The Royal Society of Chemistry



Figure 2. Electron microscope images of silicon nanoparticles. Reprinted and adapted with permission ^[59]. Copyright © 2015 American Chemical Society



Figure 3. Electron microscope images of silicon nanowires. Reprinted and adapted with permission^[66], Copyright © The Royal Society of Chemistry 2010



Figure 4. Electron microscope images of silicon nanopost array^[77] (left) and Ag-functionalized silicon nanowire array^[70] (right). Reprinted and adapted with permission, Copyright © 2015 The Royal Society of Chemistry and Copyright © 2016 John Wiley & Sons, Ltd



Figure 5. Bar charts summarizing the detection strategies used in the reviewed publications presented in this work.

Surface modification methods ^{a)}	Detected Molecules and Limit of Detection (LOD if	Matrix ^{b)}	Ref
	available)		
N/A	choline (40 μ M), BSA, FHV, adenovirus penton protein, β -lactoglobulin	-	[21]
N/A	four-residue peptide (MRFA), des-arg-bradykinin, bradykinin,	-	[23]
	angiotensin, adrenocorticotropic hormone (2 pmol each);		
	caffeine, antiviral drug WIN, reserpine (1 pmol each); N-octyl		
27/4	β-D-glucopyranoside		[20]
N/A	prednisolone, dalargin, bradykinin, adrenocorticotropic	-	[29]
N/A	4-amino-1-benzylpyridinium bromide (0.5 mM) 1.2-	_	[30]
14/7 \$	dihexadecanoyl-sn-glycero-3-phosphocholine (0.01 mM)	_	
	angiotensin III (2.5 mg/mL)		
N/A	heptadecanoic acid, stearic acid, nonadecanoic acid, arachidic	-	[24]
	acid, heneicosanoic acid, behenic acid (2 nmol each)		
Silvlation with chlorosilane and	verapamil (700 ymol), BSA (500 amol), phospholipids;	-	[22]
addition of physically adsorbed	bradykinin 2-9, bradykinin 1-7 and neurotensin (1 fmol peptide		
initiator	array), phosphatidylcholine, phosphatidylethanolamine,		
	codeine (15 ng/ml)		
Silylation using various silylating	perfluorophenyl silylated pSi: midazolam (200 fmol);	-	[35]
agents (TMS, C8, NH2)	propafenone (200 fmol); verapamil (200 fmol), des-Arg9-		
	bradykinin (800 ymol)		
	amine silylated pSi: sucrose (25 pmol); maltotriose (25 pmol)		
	phenylalanine, alanine, isoleucine/leucine, glutamic acid		
Silylation with F5PhPr	tyrindoxyl sulfate; 6,6'-dibromoindigo	-	[36]
	murexine; tyrindoxyl hydrogen sulfate; tyrindoleninone; Tyrian	-	[37]
	purple		
	cholesterol; nicotine; methamphetamine; amphetamine;	-	[26]
	nonanoic acid; methadone; EDDP; codeine		
Silylation with F5PhPr, F13 and F17	MA (2.88 ng/mL); MDMA (0.66 ng/mL); cocaine (0.86	-	[33]
	ng/mL)		
Silylation with neat BisF17	methadone (14.74ng/mL in saliva; 18.84ng/mL in plasma; 19.50ng/mL in urine)	-	[38]
Silylation with neat F17and BisF17	methadone (100 ng/mL); oxycodone (2.5 ng/mL in water and	-	[39]
	10 ng/mL in PBS); flunitrazepam (100 ng/mL); MDMA (100		
	ng/mL); cocaine (100 ng/mL)		
Two levels of derivatization	phosphopeptides from α -casein, β -casein and BSA	DHB	[40]
(aminopropyl then phosphonate) and functionalization with Zr			
Two levels of functionalization (Ag	TPyP (5.2 fmol); oxytocin (0.4 pmol); PEG 400 (3 pmol); PEG	-	[41]
and 4-ATP)	2300 (30 pmol)		
Functionalization with nanometer Ag	fingermark: DTDMA C16/C16; DTDMA C16/C18; DTDMA	-	[42]
layer	C18/C18; TAG 48:1; oleic acid; stearic acid; WE 36:1;		
	Behentrimonium		
	tissue section: PC head group; oleic acid; stearic acid; 6,6'-		
	dibromoindirubin (Tyrian purple); cholesterol		
Functionalization with AgNO ₃	cholesterol and 7DHC	-	[43]
solution and addition of physically			
adsorbed initiator			
Two levels of derivatization (10	buprenorphine (100 fmol); midazolam (150 fmol); psilocin (5	-	[44]
undecenoic acid and ethyl	pmol); propranolol (4 pmol); 1-Naphthalene methylamine (6		
undecenoate)	pmol); 2-Naphthylacetic acid (5 pmol); dichloromethyl		
	phosphonate (60 pmol); dichloromethylene bisphosphonate (40		

Table 1. Functionalization methods of porous silicon substrates for detection of biomolecules

	pmol); Leu-Enkephalin-Arg (4 pmol); Des-Arg1-Bradykinin (4		
	pmol); Substance P (3 pmol)		
Derivatization by RIE etching in gas mixture O2/CF4 and toluene wash	peptide and oligosaccharide molecules	-	[32]
Two levels of derivatization	different BNP concentration solutions prepared in PBS buffer	CHCA	[34]
(undecylenic acid and NHS) and	and in human plasma (10pg/mL)		
functionalization with BNP			
antibodies			
Two levels of silylation (alkene 1 and	tryptic peptides of fibronectin from human neuroblastoma cell	CHCA	[45]
alkene 3) and functionalization with	line SK-N-SH		
FITC-BSA or FN			
Four levels of derivatization (OH,	Trx-urodilatin(1 pM)	SA	[27]
carboxyl, NHS and NTA) and			
functionalization with Ni2+			
Functionalization with angiotensin I	angiotensin I (1 nM in diluted plasma)	CHCA	[28]
antibodies			
Derivatization with pentane	Substance P, renin substrate tetrapeptide, angiotensin I (ng/ml	CHCA	[31]
	standard solutions) and calcitonin	SA	
Functionalization with	model sample consisting of HRP digest (100 mg/mL); excess	CHCA	[46]
electrochemically deposited Au	BSA; serum peptides, insulin		
Functionalization with AuNPs	GSH (10 µg/mL in standard solution), GSH (healty and	-	[47]
	Irinotecan-treated Caco-2 cells) and l-cysteine (10 µg/mL)		
Addition of BisF17 initiator coating	angiotensin III, bradykinin, and angiotensin I (0.3 μ M each),	DHB	[48]
	lipid species		
Functionalization with AgNO ₃	testosterone, vitamin D3, glucose (300 fmol), sucrose (500	-	[49]
solution and addition of initiator	fmol), maltotriose (800 amol), maltohexaose, maltoheptaose, β -		
coating	and γ -cyclodextrin; trans-androsterone, progesterone,		
	corticosterone, and prednisone (500 fmol each), cholesterol		
Addition of BisF17 initiator coating	spermidine, arginine, adenosine, palmitoylcarnitine, verapamil,	-	[50]
on black silicon*	STAL-2 (100 fmol-10 pmol), Bradykinin		
Addition of BisF17 initiator coating	arginine, palmitoylcarnitine, streptomycin, bradykinin,	-	[51]
	angiotensin, ACTH residues ("clip" 1-17, 18-39, 7-38),		
	insulin B, neurotensin		
N/A ^{c)}	dextromethorphan and CelE-CBM3a enzymatic assay	-	[52]
N/A	arginine, tryptophan, histidine, methionine, glutamine, and	-	[53]
	glycine (5 µg/mL each)		

a) N/A = information not available; b) - matrix-free method; c) not porous silicon

Table 2. Silicon nanostructures: fabrication, functionalization and applications

	Fabrication Methods	Surface modification methods ^{a)}	Analyzed Molecules	Matrix ^{b)}	Ref.
	Sol-gel synthesis (SiO ₂ NPs)	N/A	48 sequences from synthetic peptides mimicking protein digests	-	[55]
	Sol-gel synthesis	Modification with MPS and polymerization of MMA	standard peptide angiotensin II (DRVYIHPF), standard protein cytochrome C	CHCA	[56]
	(magnetite core- shell	Functionalization with Cu ²⁺ ions by surface grafting	angiotensin II, tryptic BSA digest, peptides from human serum and urine	CHCA	[57]
s	Fe ₃ O ₄ @SiO ₂ microspheres)	CSNPs dried and calcined in N ₂	angiotensin II, MYO digest and BSA digest	CHCA	[58]
io-powder		Functionalization with Au	human insulin, immunoglobulin G, human serum albumin, α 1- antitrypsin, horseradish peroxidase, β -casein, lysozyme, cytochrome C	CHCA	[59]
ticles and nar	AuNPs coated with an ultrathin silica shell	Stabilization of Au@SiO ₂ CSNPs by growth of the silica shell	aspartic acid, N-(4-hydroxyphenyl)acetamide, 2- (2-methyl-5-nitro-1H-imidazole-1-yl)ethanol, norfloxacin, amoxicillin, erythromycin, PEG1000, roxithromycin and temporin-SHf	-	[60]
Nanopar	Particles purchased from distributors	Etching with HF, oxidation with HNO ₃ and derivatization with PFP, PHP, or TMS	peptides leucine encephalin (200 µg/mL), angiotensin II (DRVYIHPF) (100 µg/mL), nucleobase adenine (400 µg/mL), propafenone (10 pmol/µL), verapamil (10 pmol/µL), trioctylamine (80 µg/mL) morphine and propafenone from spiked urine ametryn and altretamine from spiked soil	-	[61]
		Oxidation with HNO ₃ and silylation with C10, C6 or C3	2-methyl, 4-methyl, 3-methoxy, 4-methoxy and 4-chloro benzylpyridiniums	-	[62]
		Functionalization with Au by dipping in AuNP solution	surface-confined DNA: ss-DNA and ds-DNA	3-HPA	[63]
owires	Au nanocluster- catalyzed vapor- liquid-solid (VLS) growth	Oxidation with ozone and silylation with PFP	cocaine (3 μM spiked saliva), BSA and FHV tryptic digests (1 μM), and des-Arg9-bradykinin, midazolam , propafenone, and verapamil (all 1 mg/mL aqueous solution)	-	[64]
Nano	mechanism		chloride salts of seven benzyl-substituted benzylpyridinium ions (70 µM standard solution)	-	[65]
		sample deposition	glyceride		[67]
	etching (variations on	silylation with OTS, FDTS, or ODCS	peptide mixture: Des-Arg-bradykinin, angiotensin I, fibrinopeptide B and neurotensin (10 fmol/ μ L), verapamil (5 fmol/ μ L) and Sutent (10 fmol/ μ L)	-	[07]
	HF/AgNO3 aq. solution)	Silylation with OTS	peptide mixture (fmol/mL): Des-Arg-Bradykinin, angiotensin I, fibrinopeptide B and neurotensin	-	[68]
Irfaces		Functionalization with AgNPs	15 tryptic peptides and 14 Lys-N peptides linoleic acid, oleic acid, arachidonic acid,	-	[69]
tured sı	Metal-assisted	N/A	squalene, diacyigiycerol single model peptide sample des-Arg9 Bradykinin (1 pM)	-	[71]
Nanostructured surfaces	etching (using Ag or Au nanostructures and aqueous HF etching	Oxidation by ozone and silylation with F13	methadone (100 ng/mL aqueous solution), EDDP from clinical samples of blood plasma, saliva, and urine and peptide mixture: angiotensin I, angiotensin II, substance P, bombesin, ACTH clip 1–17, ACTH clip 18–39, somatostatin 28	_	[72]
	solution)	Oxidation by O ₂ plasma and AuNP grafting by immersion in HF/HAuCl4 solution	glucose (100 µM aqueous solution) (50 mM spiked urine samples)	-	[73]

	Au deposited by e-beam	natural products in microbial interactions:	-	[74]
	evaporation, immersion in	metabolites, peptides		
	1:1:1 volume ratio of			
	HF/H2O2/EtOH solution			
Reactive ion	N/A	adenosine, Pro-Leu-Gly tripeptide, and	-	[75]
etching (mask		bradykinin, fentanyl, BSA digest and standard		
and gas mixture		carnitine metabolite cocktail		
plasma)	N/A	PEG, bradykinin, Arg, and TMZ (100 µM in	CHCA	[76]
		standard solutions) and glucose from urine	DHB	
		samples from healthy and diabetic patients		
	N/A	buprenorphine, norbuprenorphine, ropivacaine,	-	[77]
		amiodarone, chlorpheniramine, fentanyl,		
		clonidine, nordiazepam, metoprolol, verapamil		
	photopatterning, RIE etching	metabolites from hepatocyte extracts or urine	-	[78]
	and cleaning using standard	aliquots		
	microelectronics steps			
Laser irradiation	N/A	angiotensin I, bovine insulin, PPG1000 and	-	[79]
(in air, SF6 gas,		PEG400 (0.5 nmol/µL)		
or DI water)				
Commercial	hydrophobic organic coating	clonidine, propranolol, quinidine, papaverine,	-	[80]
NALDI targets		verapamil, ketoconazol, prazosin, haloperidol		

^{a)} N/A= information not available; ^{b)} - no matrix used



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Supporting Information

Silicon-based Laser Desorption Ionization Mass Spectrometry for the Analysis of Biomolecules: A Progress Report

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Pre-etch cleaning process	Current	Etching solution	Time	Illumination	Teflon	Ref.
	density	(vol/vol)	(sec)		cell	
	(mA/cm^2)	(()))	(500)			
n-type	71	1:1	60-	300-W tungsten	N/A	[23]
	-	EtOH/49% HF	180	filament bulb		
p-type	37	1:1	10800	in the dark	N/A	[23]
1 51		EtOH/49% HF				
5% HF/ethanol solution and	40	1:1	300	4 ns pulses of a	Yes	[29]
then rinsed in deionized		EtOH/48% HF	prior	355 nm		
water, acetone, and			laser	frequency-tripled		
methanol			and	Nd:YAG laser		
			900	operating at a		
			with	20Hz repetition		
			laser	rate and with an		
				intensity of 4		
				W/cm ²		
N/A	5	25% EtOH/HF	60	white-light	N/A	[21]
				50 mW/cm^2		
Methanol wash 3-5 times	48	EtOH /25%HF	1800	N/A	Yes	[22]
N/A	10 - 50	1:1	N/A	300–500 W	Yes	[44]
		EtOH/50%HF		halogen lamp		
N/A	5	25% EtOH/HF	120	white light	N/A	[35]
N/A	4	2:3	100	250 W tungsten	Yes	[40]
		EtOH/49%HF		filament bulb		
Boiled in 3:1 (v/v)	100	1:3	180	N/A	N/A	[34]
concentrated H ₂ SO ₄ /30%		EtOH/40%HF				
H ₂ O ₂ for 30min and then						
rinsed extensively with						
Milli-Q water.						
Cleaned with $3:1 (v/v)$	60	1:3	600	In the dark	Yes	[41]
H ₂ SO ₄ /H ₂ O ₂ for 30 min,		EtOH/40%HF				
rinsed with copious						
amounts of water and						
absolute ethanol, and then						
immersed in water prior to						
the etching procedure						[40]
piranha solution for 30 min	300	EtOH /25%HF	1800	N/A	N/A	[43]
rinsed with nano- pure						
water, and blown dry with						
nitrogen gas			1.5.5			[27]
N/A	20	1:1 EtOH/HF	120	fiber optic light	Yes	[36]
				source*		[47]
Rinsed with methanol,	20	1:3	300	N/A	Yes	[45]
acetone and DCM		EtOH/48%HF				

Table S1. Electrochemical etching fabrication methods

Sonicated in 99.9% methanol and dried under a stream of nitrogen.	4	1:1 EtOH/HF	120	fiber optic light source*	Yes	[33]
N/A	20	1:1 EtOH/HF	120	fiber optic light source*	Yes	[26]
N/A	27	1:2 EtOH/HF	300	fiber optic light source*	Yes	[38]
N/A	20	1:1 EtOH/HF	120	fiber optic light source*	Yes	[42]
Sonicated in 99.9% methanol and dried under a stream of nitrogen.	3.2	1:1 EtOH/HF	120	fiber optic light source*	Yes	[39]
N/A	20	1:1 EtOH/HF	120	fiber optic light source*	Yes	[37]
N/A	30	1:4 EtOH/40%HF	30– 600	N/A	Yes	[46]
piranha solution for 30 min, thoroughly washed in DI H ₂ O and dried with N ₂	32	EtOH /25%HF	1800	In the dark	N/A	[48]
three solvent baths: trichloroethylene, acetone, and methanol for 15 min sequentially	2360	EtOH /24%HF	120- 4800	N/A	Yes	[51]
N/A	N/A	HF/HNO3/ H2O (1:3:5) solution	60- 240	N/A	N/A	[53]
piranha solution for 30 min and rinsed by nanopure water and dried with N ₂	300	EtOH /25%HF	1800	N/A	Yes	[49]

N/A - information not available; * - no mention of light source or type

Table S2. Stabilization methods of porous silicon substrates

Stabilization method	Ref
ozone oxidation and immersion in diluted aqueous HF solution	[21]
ozone oxidation (hydroxyl-terminated surface)	[35][36][37][33][26][38][39] [42][22]
ozone oxidation and re-etching in ethanol containing 5% HF	[40]
wash with absolute ethanol and drying with nitrogen	[41]
pSi rinsed with a mixture of ethanol–deionized water, then kept in ethanol for 5 min, and finally dried carefully under nitrogen flow.	[44]
pSi was rinsed with pure ethanol and pentane, and then dried with nitrogen	[34]
pSi washed with methanol, acetone and DCM and dried with nitrogen	[45]
Immersion in deionized water, then in ethanol and pentane, dried with nitrogen and stored in a vacuum chamber	[31]

Table S3. Molecules and their respective detection strategies

I.2-dihexalecanoyi-sn-giyeerno-3 phosphocholine (DPPC)small molecule psipSiNoneno matrix n3020091.4-b-Cellotetraose-probepeptidenano SiNoneno matrix7320121.4-b-Decllotetraose-probepeptidenano SiDerivatizationno matrix7320121.5 tryptic peptidespeptidenano SiDerivatizationno matrix7320121.5 tryptic peptidespeptidenano SiDerivatizationno matrix4320022.(2-methyl-5-nitro-1H-small moleculepSiDerivatizationno matrix4320022.(2-methyl-5-nitro-1H-small moleculepSiDerivatizationno matrix4320024-anino-1-benzylpyridiniumsmall moleculepSiDerivatizationno matrix6120096.6-dihromoindigosmall moleculepSiDerivatizationno matrix6120016.6-dihromoindigosmall moleculepSiDerivatizationno matrix6120017DHCsmall moleculepSiDerivatizationno matrix612001ACTH crigidus ("clip" 1-17,peptidenano SiDerivatizationno matrix61201210HCsmall moleculepSiNoneno matrix62201220HCmanonano SiNoneno matrix752014ACTH crigidus ("clip" 1-17,peptidenano SiNoneno matrix742016 <td< th=""><th>Molecule</th><th>Class</th><th>Substrate</th><th>Surface</th><th>Matrix</th><th>Ref.</th><th>Year</th></td<>	Molecule	Class	Substrate	Surface	Matrix	Ref.	Year
1.2-dihexadecanoyi-sn-giyeeno-3- phosphocholine (DPPC) small molecule pSi None no matrix 30 2009 1.4-b-O-cellotetraose-probe peptide nano Si None no matrix 51 2012 14 Lys-N peptides peptide nano Si Derivatization no matrix 73 2012 1-Naphthalene methylamine small molecule pSi Derivatization no matrix 73 2012 1-Naphthalene methylamine small molecule pSi Derivatization no matrix 73 2012 1-Naphthalene methylamine small molecule pSi Derivatization no matrix 43 2002 2-Camethyl-S-nitro-11- small molecule pSi Derivatization no matrix 43 2002 4-anino-1-benzylpyridinium small molecule pSi Derivatization no matrix 50 2014 4-thoro benzylpyridinium small molecule pSi Perivatization no matrix 75 2014 ACTH clip 1-17 peptide nano Si Derivatization no matrix 75 2014 ACTH residue small molecule nano Si None no matrix 77 2010 ACTH residue small molecule <th></th> <th></th> <th></th> <th>modification</th> <th></th> <th></th> <th></th>				modification			
phosphocholine (DPPC)peptidenano SiNoneno matrix5120171.4.b-D-cellotetraose-probepeptidenano SiDerivatizationno matrix73201215 tryptic peptidespeptidenano SiDerivatizationno matrix73201215 tryptic peptidespeptidenano SiDerivatizationno matrix7320122-(2-methyl-5-nitro-1H-small moleculeSiNPNoneno matrix7320022-Carmethyl-5-nitro-1H-small moleculePSiDerivatizationno matrix7320022-Naphthalene methylaminusmall moleculePSiDerivatizationno matrix7320022-Naphthylacetic acidsmall moleculePSiDerivatizationno matrix7320024-anino-1-benzylpyridiniumsmall moleculePSiDerivatizationno matrix7320127DHCsmall moleculePSiFunctionalizationno matrix752014ACTH clip 1-17peptidenano SiDerivatizationno matrix752014ACTH clip 18-39peptidenano SiNoneno matrix742010ACTH clip 18-39peptidenano SiNoneno matrix742010adenosinesmall moleculenano SiNoneno matrix742010ACTH clip 18-39peptidepsiNoneno matrix742010adenosinesmall moleculenano SiNone <td>1,2-dihexadecanoyl-sn-glycero-3-</td> <td>small molecule</td> <td>pSi</td> <td>None</td> <td>no matrix</td> <td>30</td> <td>2009</td>	1,2-dihexadecanoyl-sn-glycero-3-	small molecule	pSi	None	no matrix	30	2009
1.4 b-D-cellotetraose-probepeptidenano SiNoneno matrix51201714 Lys-N peptidespeptidenano SiDerivatizationno matrix73201215 tryptic peptidespeptidenano SiDerivatizationno matrix7320121-Naphthalcen methylaminesmall moleculepSiDerivatizationno matrix4320022-(2-methyl-5-nitro-IH-small moleculepSiDerivatizationno matrix4320022-(2-methyl-5-nitro-IH-small moleculepSiDerivatizationno matrix4320022-(2-methyl-5-nitro-IH-small moleculepSiDerivatizationno matrix4320022-(3-methyl-5-nitro-IH-small moleculepSiDerivatizationno matrix4320024-chloro benzylpyridiniumsmall moleculepSiDerivatizationno matrix5120144-chloro benzylpyridiniumsmall moleculepSiFunctionalizationno matrix752014ACTH clip 1-7peptidenano SiDerivatizationno matrix752014ACTH clip 18-39peptidenano SiNoneno matrix702010adenosinesmall moleculenano SiNoneno matrix712010adenosinesmall moleculenano SiNoneno matrix722014adenosinesmall moleculepSiNoneno matrix742010adenosinesmall molec	phosphocholine (DPPC)		•				
14 1ys-N peptides peptide nano Si Derivatization no matrix 7.3 2012 15 typtic peptides peptide nano Si Derivatization no matrix 7.3 2012 1-Naphthalene methylamine small molecule pSi Derivatization no matrix 4.3 2002 2-(2-methyl-5-nitro-1H- small molecule pSi Derivatization no matrix 4.3 2002 2-Naphthylaetic acid small molecule pSi Derivatization no matrix 30 2002 4-anino-1-benzylpyridinium small molecule pSi Derivatization no matrix 35 2012 7DHC small molecule pSi Derivatization no matrix 45 2012 7DHC small molecule pSi Purctionalization no matrix 75 2014 ACTH clip 1-17 peptide nano Si Derivatization no matrix 75 2014 ACTH clip 18-39 peptide nano Si Derivatization no matrix 75 2014 ACTH clip 18-39 peptide nano Si None no matrix 77 2010 adenosine small molecule nano Si None no matrix 71 </td <td>1,4-b-D-cellotetraose-probe</td> <td>peptide</td> <td>nano Si</td> <td>None</td> <td>no matrix</td> <td>51</td> <td>2017</td>	1,4-b-D-cellotetraose-probe	peptide	nano Si	None	no matrix	51	2017
15 tryptic peptides peptide nano Si Derivatization no matrix 7.3 2012 1-Naphthalene methylamine small molecule pSi Derivatization no matrix 4.3 2002 2-(2-methyl-5-nitro-IH- small molecule pSi None no matrix 59 2012 2-Naphthylacetic acid small molecule pSi Derivatization no matrix 4.3 2002 4-amino-1-benzylpyridinium small molecule pSi None no matrix 3.5 2012 6.6 dibromoindigo small molecule pSi Derivatization no matrix 3.5 2014 ACTH clip 1-17 peptide nano Si Derivatization no matrix 7.5 2014 ACTH clip 1-3.9 peptide nano Si Derivatization no matrix 7.5 2014 ACTH residues (*clip * 1-7) peptide nano Si Derivatization no matrix 7.2 2010 adenosine small molecule nano Si None no matrix 7.2 2010 adenosine small molecule pSi	14 Lys-N peptides	peptide	nano Si	Derivatization	no matrix	73	2012
1-Naphthalene methylamine small molecule pSi Derivatization no matrix 43 2002 2-(2-methyl-5-niro-IH- imidazole-1-ylpethanol small molecule SiNP None no matrix 59 2012 2-Naphthylacetic acid small molecule pSi Derivatization no matrix 30 2002 4-amino-1-benzylpyridinium small molecule pSi None no matrix 30 2009 6,6'-dibromoindigo small molecule pSi Derivatization no matrix 35 2012 7DHC small molecule pSi Derivatization no matrix 75 2014 ACTH clip 1-37 peptide nano Si Derivatization no matrix 75 2014 ACTH clip 1-39 peptide pSi None no matrix 70 2016 adenosine small molecule nano Si None no matrix 71 2010 adenosine small molecule nano Si None no matrix 21 2001	15 tryptic peptides	peptide	nano Si	Derivatization	no matrix	73	2012
2-(2-methyl-5-nitro-1H- imidazole-1-yl)ethanol small molecule SiNP None no matrix 59 2012 2-Naphthylacetic acid small molecule pSi Derivatization no matrix 43 2002 4-anino-1-benzylpyridinium small molecule pSi Derivatization no matrix 30 2009 6.6'-dibromoindigo small molecule pSi Derivatization no matrix 32 2012 7DHC small molecule pSi Derivatization no matrix 42 2010 ACTH clip 1-17 peptide nano Si Derivatization no matrix 75 2014 ACTH residues ("clip" 1-17, peptide nano Si Derivatization no matrix 75 2017 18-39, 7-38) adenosine small molecule nano Si None no matrix 71 2010 adenosine small molecule nano Si None no matrix 21 2001 adenosine small molecule nano Si None no matrix 21 2001 adenosine small molecule pSi	1-Naphthalene methylamine	small molecule	pSi	Derivatization	no matrix	43	2002
imidazole-1-yljethanolcmpSiDerivatizationno matrix4320022-Naphthylacetic acidsmall moleculepSiDerivatizationno matrix302009bronidesmall moleculepSiNoneno matrix3520094-chloro benzylpyridinium saltssmall moleculepSiDerivatizationno matrix6120096.6'-dibromoindigosmall moleculepSiDerivatizationno matrix422010ACTH clip 1-17peptidenano SiDerivatizationno matrix752014ACTH residues ("clip" 1-17, 18-39,peptidenano SiDerivatizationno matrix752017adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculepSiNoneno matrix772010adenosinesmall moleculepSiNoneno matrix732014adenosinesmall moleculepSiNoneno matrix732019adenosinesmall moleculepSiNoneno matrix732010adenosinesmall moleculepSiNoneno matrix742004adenosine <td>2-(2-methyl-5-nitro-1H-</td> <td>small molecule</td> <td>SiNP</td> <td>None</td> <td>no matrix</td> <td>59</td> <td>2012</td>	2-(2-methyl-5-nitro-1H-	small molecule	SiNP	None	no matrix	59	2012
2-Naphthylacetic acid small molecule pSi Derivatization no matrix 43 2002 4-amino-1-benzylpyridinium small molecule pSi None no matrix 30 2009 bronide small molecule SiNP Derivatization no matrix 32 2012 4-chloro benzylpyridinium salts small molecule pSi Derivatization no matrix 35 2012 7DHC small molecule pSi Functionalization no matrix 42 2010 ACTH clip 1–17 peptide nano Si Derivatization no matrix 75 2014 ACTH residues ("clip" 1–17, 18–39, 7–38) peptide nano Si None no matrix 70 2010 adenosine small molecule nano Si None no matrix 71 2010 adenosine small molecule nano Si None no matrix 72 2010 adenosine small molecule pSi None no matrix 23 1999 Afterocorticotropic hormone peptide pSi None no matrix 34 2004 alurencorticotropic hormone small molecule SiNP Derivatization no matrix 60 <td>imidazole-1-yl)ethanol</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>	imidazole-1-yl)ethanol						
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4-chloro benzylpyridinium saltssmall moleculeSiNPDerivatizationno matrix6120096,6'-dibromoindigosmall moleculepSiDerivatizationno matrix3520127DHCsmall moleculepSiFunctionalizationno matrix422010ACTH clip 1–17peptidenano SiDerivatizationno matrix752014ACTH residues ("clip" 1–17,peptidepSiNoneno matrix50201718–39, 7–38)no matrixpeptidenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix772010adenosirus penton proteinproteinpSiNoneno matrix231999Adrenocorticotropic hormonepeptidepSiNoneno matrix342004alarinesmall moleculepSiDerivatizationno matrix342007alarinesmall moleculepSiDerivatizationno matrix342004altretaminesmall moleculepSiDerivatizationno matrix352012adinoasmall moleculepSiDerivatizationno matrix342004altretaminesmall moleculepSiDerivatizationno matrix592012anetrynsmall moleculeSiNPDerivatizationno matrix50 <td>bromide</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>	bromide						
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7DHCsmall moleculepSiFunctionalizationno matrix422010ACTH clip 1-17peptidenano SiDerivatizationno matrix752014ACTH clip 18-39peptidenano SiDerivatizationno matrix752014ACTH residues ("clip" 1-17, 18-39, 7-38)peptidepSiNoneno matrix502017adenosinesmall moleculenano SiNoneno matrix492016adenosinesmall moleculenano SiNoneno matrix772010adenovirus penton proteinproteinpSiNoneno matrix212001adenocorticotropic hormonepeptidepSiNoneno matrix231999Adrenocorticotropic HormonepeptidepSiNoneno matrix342004alarinesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix502012amphetaminesmall moleculeSiNPNoneno matrix502017amoxicillinsmall moleculeSiNPNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix502012amotrynsmall moleculeSiNPNoneno matrix502012amotrynsmall molecule	6,6'-dibromoindigo	small molecule	pSi	Derivatization	no matrix	35	2012
ACTH clip 1-17peptidenano SiDerivatizationno matrix752014ACTH clip 18-39peptidenano SiDerivatizationno matrix752014ACTH residues ("clip" 1-17, 18-39, 7-38)peptidepSiNoneno matrix502017adenosinesmall moleculenano SiNoneno matrix492016adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix212001adenocritoctropic hormonepeptidepSiNoneno matrix231999Adrenocorticotropic HormonepeptidepSiNoneno matrix242004alarinesmall moleculepSiDerivatizationno matrix342004alarinesmall moleculepSiDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amodaronesmall moleculeSiNPDerivatizationno matrix622015amodaronesmall moleculeSiNPNoneno matrix502017amodaronesmall moleculeSiNPNoneno matrix622015amodaronesmall moleculeSiNPNoneno matrix502017amodaronesmall moleculeSiNPNoneno matrix502017amodaronesmall molecule<	7DHC	small molecule	pSi	Functionalization	no matrix	42	2010
ACTH clip 18–39peptidenano SiDerivatizationno matrix752014ACTH residues ("clip" 1–17, 18–39, 7–38)peptidepSiNoneno matrix502017adenosinesmall moleculenano SiNoneno matrix492016adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix772010adenosireproteinproteinpSiNoneno matrix212001adenocorticotropic hormonepeptidepSiNoneno matrix231999Adrenocorticotropic HormonepeptidepSiNoneno matrix292009(ACTH) 1–17small moleculeSiNPDerivatizationno matrix602007alaninesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix502017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationno matrix722010angiotensin Ipeptid	ACTH clip 1–17	peptide	nano Si	Derivatization	no matrix	75	2014
ACTH residues ("clip" 1-17, 18-39, 7-38)peptidepSiNoneno matrix502017adenosinesmall moleculenano SiNoneno matrix492016adenosinesmall moleculenano SiNoneno matrix772010adenosinesmall moleculenano SiNoneno matrix212001adenovirus penton proteinproteinpSiNoneno matrix231999Adrenocorticotropic hormonepeptidepSiNoneno matrix292009(ACTH) 1-17small moleculepSiDerivatizationno matrix342004alaninesmall moleculepSiDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix502017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiFunctionalizationCHCA or SA312006angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin I <td>ACTH clip 18–39</td> <td>peptide</td> <td>nano Si</td> <td>Derivatization</td> <td>no matrix</td> <td>75</td> <td>2014</td>	ACTH clip 18–39	peptide	nano Si	Derivatization	no matrix	75	2014
18-39, 7-38)Image: Small moleculenano SiNoneno matrix492016adenosinesmall moleculenano SiNoneno matrix772010adenovirus penton proteinproteinpSiNoneno matrix212001adrenocorticotropic hormonepeptidepSiNoneno matrix231999Adrenocorticotropic hormonepeptidepSiNoneno matrix232009(ACTH) 1-17small moleculepSiDerivatizationno matrix342004alantesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amotidaronesmall moleculeSiNPDerivatizationno matrix602007amotidaronesmall moleculeSiNPNoneno matrix602007amoticillinsmall moleculeSiNPNoneno matrix602012amphetaminesmall moleculeSiNPNoneno matrix602017angiotensinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix502017amoticillinsmall moleculepSiNoneno matrix502017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSi <td>ACTH residues ("clip" 1-17,</td> <td>peptide</td> <td>pSi</td> <td>None</td> <td>no matrix</td> <td>50</td> <td>2017</td>	ACTH residues ("clip" 1-17,	peptide	pSi	None	no matrix	50	2017
adenosinesmall moleculenano SiNoneno matrix492016adenosinesmall moleculenano SiNoneno matrix772010adenovirus penton proteinproteinpSiNoneno matrix212001adrenocorticotropic hormonepeptidepSiNoneno matrix231999AdrenocorticotropicHormonepeptidepSiNoneno matrix292009(ACTH) 1–17nonoNoneno matrix342004alaninesmall moleculepSiDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amodaronesmall moleculeSiNPDerivatizationno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix262015amokicillinsmall moleculeSiNPNoneno matrix262015angiotensinpeptidepSiNoneno matrix262017angiotensin IpeptidepSiNoneno matrix231999angiotensin IpeptidepSiNoneno matrix202017angiotensin IpeptidepSiPerivatizationno matrix202017angiotensin IpeptidepSiNoneno matrix202017angiotensin Ipeptidenano SiDerivatizationno matrix	18-39, 7-38)						
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adenovirus penton proteinproteinpSiNoneno matrix212001adrenocorticotropic hormonepeptidepSiNoneno matrix231999AdrenocorticotropicHormonepeptidepSiNoneno matrix292009(ACTH) 1–17small moleculepSiDerivatizationno matrix342004alarinesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculeSiNPNoneno matrix262015angiotensinpeptidepSiNoneno matrix231999angiotensin IpeptidepSiNoneno matrix262017angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptiden	adenosine	small molecule	nano Si	None	no matrix	77	2010
adrenocorticotropic hormone Adrenocorticotropic (ACTH) 1–17peptidepSiNoneno matrix231999Adrenocorticotropic (ACTH) 1–17Formone small moleculepSiNoneno matrix292009alaninesmall moleculepSiDerivatizationno matrix342004altretaminesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPDerivatizationno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix262015angiotensinpeptidepSiNoneno matrix502017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or312006angiotensin Ipeptidenano SiDerivatizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762017angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix <t< td=""><td>adenovirus penton protein</td><td>protein</td><td>pSi</td><td>None</td><td>no matrix</td><td>21</td><td>2001</td></t<>	adenovirus penton protein	protein	pSi	None	no matrix	21	2001
Adrenocorticotropic (ACTH) 1–17Hormone PpeptidepSiNoneno matrix292009alaninesmall moleculepSiDerivatizationno matrix342004altretaminesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculenano SiNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix592012angiotensinsmall moleculepSiNoneno matrix231999angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiEructionalizationCHCA or SA312006angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiNoneno matrix702006	adrenocorticotropic hormone	peptide	pSi	None	no matrix	23	1999
(ACTH) 1–17Image: small moleculepSiDerivatizationno matrix342004altretaminesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeSiNPDerivatizationno matrix602007amoxicillinsmall moleculeNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix262015angiotensinpeptidepSiNoneno matrix262017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationno matrix502017angiotensin IpeptidepSiFunctionalizationCHCA or SA312006angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiN	Adrenocorticotropic Hormone	peptide	pSi	None	no matrix	29	2009
alaninesmall moleculepSiDerivatizationno matrix342004altretaminesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculeNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculeSiNPNoneno matrix262015angiotensinpeptidepSiDerivatizationno matrix231999angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762016angiotensin Ipeptidenano SiN	(ACTH) 1–17						
altretaminesmall moleculeSiNPDerivatizationno matrix602007ametrynsmall moleculeSiNPDerivatizationno matrix602007amiddaronesmall moleculenano SiNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculeSiNPNoneno matrix262015angiotensinpeptidepSiDerivatizationno matrix231999angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiNoneno matrix702006angiotensin Ipeptidenano SiNoneNoneNoneNoneangiotensin Ipeptidenano SiNoneDHB47 </td <td>alanine</td> <td>small molecule</td> <td>pSi</td> <td>Derivatization</td> <td>no matrix</td> <td>34</td> <td>2004</td>	alanine	small molecule	pSi	Derivatization	no matrix	34	2004
ametrynsmall moleculeSiNPDerivatizationno matrix602007amiodaronesmall moleculenano SiNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix262015angiotensinpeptidepSiNoneno matrix231999angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or SA312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiNoneno matrix702006	altretamine	small molecule	SiNP	Derivatization	no matrix	60	2007
amiodaronesmall moleculenano SiNoneno matrix652015amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix262015angiotensinpeptidepSiNoneno matrix231999angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or312006angiotensin IpeptidepSiDerivatizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepsiNoneDHB472016angiotensin IpeptidepsiNoneno matrix702006	ametryn	small molecule	SiNP	Derivatization	no matrix	60	2007
amoxicillinsmall moleculeSiNPNoneno matrix592012amphetaminesmall moleculepSiDerivatizationno matrix262015angiotensinpeptidepSiNoneno matrix231999angiotensinsmall moleculepSiNoneno matrix502017angiotensin IpeptidepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or SA312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	amiodarone	small molecule	nano Si	None	no matrix	65	2015
amphetaminesmall moleculepSiDerivatizationno matrix262015angiotensinpeptidepSiNoneno matrix231999angiotensinsmall moleculepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or SA312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiNoneDHB472016angiotensin Ipeptidenano SiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	amoxicillin	small molecule	SiNP	None	no matrix	59	2012
angiotensinpeptidepSiNoneno matrix231999angiotensinsmall moleculepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or SA312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiNoneno matrix702006	amphetamine	small molecule	pSi	Derivatization	no matrix	26	2015
angiotensinsmall moleculepSiNoneno matrix502017angiotensin IpeptidepSiDerivatizationCHCA or SA312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin IpeptidepSiNoneno matrix702006	angiotensin	peptide	pSi	None	no matrix	23	1999
angiotensin IpeptidepSiDerivatizationCHCA or SA312006angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	angiotensin	small molecule	pSi	None	no matrix	50	2017
Image: constraint of the stateImage: constraint of the stateIm	angiotensin I	peptide	pSi	Derivatization	CHCA or	31	2006
angiotensin IpeptidepSiFunctionalizationCHCA282011angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepsiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006					SA		
angiotensin Ipeptidenano SiDerivatizationno matrix722010angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	angiotensin I	peptide	pSi	Functionalization	CHCA	28	2011
angiotensin Ipeptidenano SiDerivatizationno matrix752014angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	angiotensin I	peptide	nano Si	Derivatization	no matrix	72	2010
angiotensin Ipeptidenano SiDerivatizationno matrix762011angiotensin IpeptidepSiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	angiotensin I	peptide	nano Si	Derivatization	no matrix	75	2014
angiotensin IpeptidepSiNoneDHB472016angiotensin Ipeptidenano SiNoneno matrix702006	angiotensin I	peptide	nano Si	Derivatization	no matrix	76	2011
angiotensin I peptide nano Si None no matrix 70 2006	angiotensin I	peptide	pSi	None	DHB	47	2016
	angiotensin I	peptide	nano Si	None	no matrix	70	2006

angiotensin II	peptide	SiNP	Functionalization	CHCA	56	2010
angiotensin II	peptide	nano Si	Derivatization	no matrix	75	2014
Angiotensin II	peptide	SiNP	None	CHCA	58	2010
angiotensin II (DRVYIHPF)	peptide	SiNP	Derivatization and	CHCA	55	2010
			functionalization			
angiotensin II (DRVYIHPF)	peptide	SiNP	Derivatization	no matrix	60	2007
angiotensin III	peptide	pSi	None	no matrix	30	2009
angiotensin III	peptide	pSi	None	DHB	47	2016
antiviral drug WIN	small molecule	pSi	None	no matrix	23	1999
Arachidic acid	small molecule	pSi	None	no matrix	24	2006
arachidonic acid	lipid	nano Si	Functionalization	no matrix	64	2016
arginine	small molecule	nano Si	None	no matrix	49	2016
arginine	small molecule	pSi	None	no matrix	50	2017
arginine	small molecule	pSi	None	no matrix	52	2008
arginine acids	small molecule	nano Si	None	CHCA	69	2013
Aspartic acid	small molecule	SiNP	None	no matrix	59	2012
Behenic acid	small molecule	pSi	None	no matrix	24	2006
Behentrimonium	small molecule	pSi	Functionalization	no matrix	41	2017
benzylpyridinium ions	small molecule	SiNW	Derivatization	no matrix	67	2006
beta-cyclodextrin	peptide	pSi	Functionalization	no matrix	48	2009
beta-lactoglobulin	protein	pSi	None	no matrix	21	2001
BNP	peptide	pSi	Derivatization and	CHCA	33	2008
			functionalization			
bombesin	peptide	nano Si	Derivatization	no matrix	75	2014
bovine insulin	peptide	nano Si	None	no matrix	70	2006
bradykinin	peptide	pSi	None	no matrix	23	1999
Bradykinin	peptide	pSi	None	no matrix	29	2009
bradykinin	peptide	pSi	None	DHB	47	2016
Bradykinin	peptide	nano Si	None	no matrix	49	2016
bradykinin	small molecule	pSi	None	no matrix	50	2017
bradykinin	peptide	nano Si	None	CHCA	69	2013
bradykinin	peptide	nano Si	None	no matrix	77	2010
bradykinin 1-7	peptide	pSi	Derivatization	no matrix	22	2007
bradykinin 2-9	peptide	pSi	Derivatization	no matrix	22	2007
BSA	peptide	pSi	Derivatization	no matrix	22	2007
BSA	peptide	pSi	None	no matrix	21	2001
BSA	peptide	SiNW	Derivatization	no matrix	66	2005
BSA digest	peptide	SiNP	None	CHCA	58	2010
BSA digest	peptide	nano Si	None	no matrix	77	2010
Buprenorphine	small molecule	pSi	Derivatization	no matrix	43	2002
buprenorphine	small molecule	nano Si	None	no matrix	65	2015
butyrylcarnitine (C4)	small molecule	nano Si	None	no matrix	77	2010
caffeine	small molecule	pSi	None	no matrix	23	1999

calcitonin	peptide	pSi	Derivatization	CHCA or	31	2006
				SA		
carnitine (C1)	small molecule	nano Si	None	no matrix	77	2010
cellobiose-probe	peptide	nano Si	None	no matrix	51	2017
cellotriose-probe	peptide	nano Si	None	no matrix	51	2017
chlorpheniramine	small molecule	nano Si	None	no matrix	65	2015
Cholesterol	lipid	pSi	Functionalization	no matrix	41	2017
Cholesterol	small molecule	pSi	Functionalization	no matrix	42	2010
cholesterol	small molecule	pSi	Functionalization	no matrix	48	2009
cholesterol	small molecule	pSi	Functionalization	no matrix	48	2009
cholesterol	small molecule	pSi	Functionalization	no matrix	48	2009
cholesterol	lipid	pSi	Derivatization	no matrix	26	2015
choline	small molecule	pSi	None	no matrix	21	2001
clonidine	small molecule	nano Si	Derivatization	no matrix	68	2008
clonidine	small molecule	nano Si	None	no matrix	65	2015
Cocaine	small molecule	pSi	Derivatization	no matrix	32	2012
Cocaine	small molecule	pSi	Derivatization	no matrix	38	2015
Cocaine	small molecule	SiNW	Derivatization	no matrix	66	2005
codeine	small molecule	pSi	Derivatization	no matrix	22	2007
Codeine	small molecule	pSi	Derivatization	no matrix	26	2015
corticosterone	small molecule	pSi	Functionalization	no matrix	48	2009
cytochrome C	protein	SiNP	Derivatization and	CHCA	55	2010
			functionalization			
Dalargin	small molecule	pSi	None	no matrix	29	2009
decanoylcarnitine (C6)	small molecule	nano Si	None	no matrix	77	2010
Des-Arg1-Bradykinin	peptide	pSi	Derivatization	no matrix	43	2002
des-Arg9 Bradykinin	peptide	nano Si	None	no matrix	74	2011
des-Arg9-bradykinin	peptide	pSi	Derivatization	no matrix	34	2004
des-Arg9-bradykinin	peptide	SiNW	Derivatization	no matrix	66	2005
Des-Arg-bradykinin	peptide	nano Si	Derivatization	no matrix	72	2010
Des-Arg-Bradykinin	peptide	nano Si	Derivatization	no matrix	76	2011
des-arg-bradykinin	peptide	pSi	None	no matrix	23	1999
dextromethorphan	small molecule	nano Si	None	no matrix	51	2017
diacylglycerol	lipid	nano Si	Functionalization	no matrix	64	2016
dibutylphosphoric acid	small molecule	SiNP	Derivatization	no matrix	60	2007
Dichloromethyl phosphonate	small molecule	pSi	Derivatization	no matrix	43	2002
Dichloromethylene	small molecule	pSi	Derivatization	no matrix	43	2002
bisphosphonate						
diglycerides	lipid	SiNW	Functionalization	no matrix	63	2010
ds-DNA	oligonucleotides	SiNP	Functionalization	3-HPA	62	2009
DTDMA C16/C16	small molecule	pSi	Functionalization	no matrix	41	2017
DTDMA C16/C18	small molecule	pSi	Functionalization	no matrix	41	2017
DTDMA C18/C18	small molecule	pSi	Functionalization	no matrix	41	2017
EDDP	small molecule	pSi	Derivatization	no matrix	26	2015

EDDP	small molecule	nano Si	Derivatization	no matrix	75	2014
erythromycin	small molecule	SiNP	None	no matrix	59	2012
fatty acids and glyceride	small molecule	SiNW	Functionalization	no matrix	63	2010
fentanyl	small molecule	nano Si	None	no matrix	65	2015
fentanyl	small molecule	nano Si	None	no matrix	77	2010
FHV	protein	pSi	None	no matrix	21	2001
FHV tryptic digests	protein	SiNW	Derivatization	no matrix	66	2005
fibrinopeptide B	peptide	nano Si	Derivatization	no matrix	72	2010
fibrinopeptide B	peptide	nano Si	Derivatization	no matrix	76	2011
Flunitrazepam	small molecule	pSi	Derivatization	no matrix	38	2015
four-residue peptide (MRFA)	peptide	pSi	None	no matrix	23	1999
glucose	small molecule	pSi	Functionalization	no matrix	48	2009
glucose	small molecule	pSi	Functionalization	no matrix	48	2009
glucose	small molecule	nano Si	None	no matrix	69	2013
glucose	small molecule	nano Si	Derivatization and	no matrix	71	2015
			functionalization			
glucose	small molecule	nano Si	Derivatization and	no matrix	71	2015
			functionalization			
glucose-probe	peptide	nano Si	None	no matrix	51	2017
glutamic acid	small molecule	pSi	Derivatization	no matrix	34	2004
glutamine	small molecule	pSi	None	no matrix	52	2008
glycine	small molecule	pSi	None	no matrix	52	2008
GSH	small molecule	pSi	Functionalization	no matrix	46	2017
GSH	small molecule	pSi	Functionalization	no matrix	46	2017
haloperidol	small molecule	nano Si	Derivatization	no matrix	68	2008
Heneicosanoic acid	small molecule	pSi	None	no matrix	24	2006
Heptadecanoic acid	small molecule	pSi	None	no matrix	24	2006
histidine	small molecule	pSi	None	no matrix	52	2008
insulin	peptide	pSi	Functionalization	CHCA	45	2014
insulin	peptide	SiNP	Functionalization	CHCA	57	2015
insulin B	peptide	pSi	None	no matrix	50	2017
isoleucine/leucine	small molecule	pSi	Derivatization	no matrix	34	2004
ketoconazol	small molecule	nano Si	Derivatization	no matrix	68	2008
L-cysteine	small molecule	pSi	Functionalization	no matrix	46	2017
Leu-Enkephalin-Arg	peptide	pSi	Derivatization	no matrix	43	2002
linoleic acid	lipid	nano Si	Functionalization	no matrix	64	2016
lipid species	lipid	pSi	None	DHB	47	2016
maltoheptaose	peptide	pSi	Functionalization	no matrix	48	2009
maltohexaose	peptide	pSi	Functionalization	no matrix	48	2009
maltotriose	small molecule	pSi	Functionalization	no matrix	48	2009
maltotriose	small molecule	pSi	Derivatization	no matrix	34	2004
MDMA	small molecule	pSi	Derivatization	no matrix	32	2012
MDMA	small molecule	pSi	Derivatization	no matrix	38	2015
methadone	small molecule	pSi	Derivatization	no matrix	26	2015

Methadone	small molecule	pSi	Derivatization	no matrix	37	2017
Methadone	small molecule	pSi	Derivatization	no matrix	37	2017
Methadone	small molecule	pSi	Derivatization	no matrix	37	2017
Methadone	small molecule	pSi	Derivatization	no matrix	38	2015
methadone	small molecule	nano Si	Derivatization	no matrix	75	2014
methamphetamine	small molecule	pSi	Derivatization	no matrix	26	2015
methamphetamine	small molecule	pSi	Derivatization	no matrix	32	2012
methionine	small molecule	pSi	None	no matrix	52	2008
metoprolol	small molecule	nano Si	None	no matrix	65	2015
Midazolam	small molecule	pSi	Derivatization	no matrix	43	2002
Midazolam	small molecule	pSi	Derivatization	no matrix	34	2004
midazolam	small molecule	SiNW	Derivatization	no matrix	66	2005
model sample of HRP digest and	peptide	pSi	Functionalization	CHCA	45	2014
BSA						
morphine	small molecule	SiNP	Derivatization	no matrix	60	2007
Murexine	small molecule	pSi	Derivatization	no matrix	36	2015
MYO digest	peptide	SiNP	None	CHCA	58	2010
myristoylcarnitine (C9)	small molecule	nano Si	None	no matrix	77	2010
N-(4-hydroxyphenyl)acetamide	small molecule	SiNP	None	no matrix	59	2012
neurotensin	peptide	pSi	Derivatization	no matrix	22	2007
neurotensin	peptide	nano Si	Derivatization	no matrix	72	2010
neurotensin	peptide	nano Si	Derivatization	no matrix	76	2011
neurotensin	peptide	pSi	None	no matrix	50	2017
nicotine	small molecule	pSi	Derivatization	no matrix	26	2015
N-octyl beta-D-glucopyranoside	peptide	pSi	None	no matrix	23	1999
Nonadecanoic acid	small molecule	pSi	None	no matrix	24	2006
nonanoic acid	lipid	pSi	Derivatization	no matrix	26	2015
norbuprenorphine	small molecule	nano Si	None	no matrix	65	2015
nordiazepam	small molecule	nano Si	None	no matrix	65	2015
norfloxacin	small molecule	SiNP	None	no matrix	59	2012
nucleobase adenine	peptide	SiNP	Derivatization	no matrix	60	2007
octanoylcarnitine (C5)	small molecule	nano Si	None	no matrix	77	2010
oleic acid	lipid	pSi	Functionalization	no matrix	41	2017
oleic acid	lipid	nano Si	Functionalization	no matrix	64	2016
Oxycodone	small molecule	pSi	Derivatization	no matrix	38	2015
Oxycodone	small molecule	pSi	Derivatization	no matrix	38	2015
Oxytocin	peptide	pSi	Functionalization	no matrix	40	2009
palmitoylcarnitine	small molecule	nano Si	None	no matrix	49	2016
palmitoylcarnitine	small molecule	pSi	None	no matrix	50	2017
palmitoylcarnitine (C10)	small molecule	nano Si	None	no matrix	77	2010
papaverine	small molecule	nano Si	Derivatization	no matrix	68	2008
PC head group	lipid	pSi	Functionalization	no matrix	41	2017
PEG	polymer	nano Si	None	DHB	69	2013
PEG 2300	polymer	pSi	Functionalization	no matrix	40	2009

PEG 400	polymer	pSi	Functionalization	no matrix	40	2009
PEG1000	polymer	SiNP	None	no matrix	59	2012
PEG400	polymer	nano Si	None	no matrix	70	2006
peptide mixture	peptide	SiNP	None	no matrix	54	2012
peptides from human serum and	peptide	SiNP	Functionalization	CHCA	56	2010
urine						
peptides leucine enkephalin	peptide	SiNP	Derivatization	no matrix	60	2007
Phenylalanine	small molecule	pSi	Derivatization	no matrix	34	2004
phosphatidylcholine	lipid	pSi	Derivatization	no matrix	22	2007
phosphatidylethanolamine	lipid	pSi	Derivatization	no matrix	22	2007
phospholipids	lipid	pSi	Derivatization	no matrix	22	2007
phosphopeptides	peptide	pSi	Derivatization and	DHB	39	2006
			functionalization			
PPG1000	polymer	nano Si	None	no matrix	70	2006
prazosin	small molecule	nano Si	Derivatization	no matrix	68	2008
Prednisolone	small molecule	pSi	None	no matrix	29	2009
prednisone	small molecule	pSi	Functionalization	no matrix	48	2009
progesterone	small molecule	pSi	Functionalization	no matrix	48	2009
Pro-Leu-Gly tripeptide	peptide	nano Si	None	no matrix	77	2010
Propafenone	small molecule	pSi	Derivatization	no matrix	34	2004
propafenone	small molecule	SiNP	Derivatization	no matrix	60	2007
propafenone	small molecule	SiNP	Derivatization	no matrix	60	2007
propafenone	small molecule	SiNW	Derivatization	no matrix	66	2005
propranolol	small molecule	nano Si	Derivatization	no matrix	68	2008
Propranolol	small molecule	pSi	Derivatization	no matrix	43	2002
Psilocin	small molecule	pSi	Derivatization	no matrix	43	2002
quinidine	small molecule	nano Si	Derivatization	no matrix	68	2008
renin substrate tetrapeptide	peptide	pSi	Derivatization	CHCA or	31	2006
				SA		
reserpine	small molecule	pSi	None	no matrix	23	1999
ropivacaine	small molecule	nano Si	None	no matrix	65	2015
roxithromycin	small molecule	SiNP	None	no matrix	59	2012
serum peptides	peptide	pSi	Functionalization	CHCA	45	2014
somatostatin 28	peptide	nano Si	Derivatization	no matrix	75	2014
spermidine	small molecule	nano Si	None	no matrix	49	2016
squalene	lipid	nano Si	Functionalization	no matrix	64	2016
ss-DNA	oligonucleotides	SiNP	Functionalization	3-HPA	62	2009
STAL-2 (hexapeptide, SFLLRN-	peptide	nano Si	None	no matrix	49	2016
NH2)						
stearic acid	lipid	pSi	Functionalization	no matrix	41	2017
Stearic acid	small molecule	pSi	None	no matrix	24	2006
stearoylcarnitine (C11)	small molecule	nano Si	None	no matrix	77	2010
streptomycin	small molecule	pSi	None	no matrix	50	2017

substance P	peptide	pSi	Derivatization	CHCA or	31	2006
				SA		
substance P	peptide	nano Si	Derivatization	no matrix	75	2014
Substance P	peptide	pSi	Derivatization	no matrix	43	2002
sucrose	small molecule	pSi	Functionalization	no matrix	48	2009
sucrose	small molecule	pSi	Functionalization	no matrix	48	2009
sucrose	small molecule	pSi	Derivatization	no matrix	34	2004
Sutent	small molecule	nano Si	Derivatization	no matrix	72	2010
TAG 48:1	lipid	pSi	Functionalization	no matrix	41	2017
temporin-SHf	peptide	SiNP	None	no matrix	59	2012
testosterone	small molecule	pSi	Functionalization	no matrix	48	2009
TMZ	small molecule	nano Si	None	no matrix	69	2013
ТРуР	small molecule	pSi	Functionalization	no matrix	40	2009
trans-androsterone	small molecule	pSi	Functionalization	no matrix	48	2009
trioctylamine	small molecule	SiNP	Derivatization	no matrix	60	2007
tripalmitin	lipid	SiNW	Functionalization	no matrix	63	2010
tristearin	lipid	SiNW	Functionalization	no matrix	63	2010
Trx-urodilatin	protein	pSi	Derivatization and	SA	27	2009
			functionalization			
tryptic BSA digest	peptide	SiNP	Functionalization	CHCA	56	2010
tryptic peptide of protein	peptide	pSi	Derivatization and	CHCA	44	2012
fibronectin			functionalization			
tryptophan	small molecule	pSi	None	no matrix	52	2008
Tyrian purple	small molecule	pSi	Functionalization	no matrix	41	2017
Tyrian purple	small molecule	pSi	Derivatization	no matrix	36	2015
tyrindoleninone	small molecule	pSi	Derivatization	no matrix	36	2015
tyrindoxyl hydrogen sulfate	small molecule	pSi	Derivatization	no matrix	36	2015
Tyrindoxyl sulfate	small molecule	pSi	Derivatization	no matrix	35	2012
verapamil	small molecule	nano Si	Derivatization	no matrix	68	2008
verapamil	small molecule	pSi	Derivatization	no matrix	22	2007
verapamil	small molecule	pSi	Derivatization	no matrix	34	2004
verapamil	small molecule	SiNP	Derivatization	no matrix	60	2007
verapamil	small molecule	SiNW	Derivatization	no matrix	66	2005
verapamil	small molecule	nano Si	None	no matrix	49	2016
verapamil	small molecule	nano Si	None	no matrix	65	2015
verapamil	small molecule	nano Si	Derivatization	no matrix	72	2010
vitamin D3	small molecule	pSi	Functionalization	no matrix	48	2009
WE 36:1	small molecule	pSi	Functionalization	no matrix	41	2017
γ-cyclodextrin	peptide	pSi	Functionalization	no matrix	48	2009