

Matrix-Free Laser Desorption Ionization Mass Spectrometry as an Efficient Tool for the Rapid Detection of Opiates in Crude Extracts of Papaver somniferum

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2 rapid detection of opiates from crude extracts of *Papaver somniferum*

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7 ABSTRACT

Having a long history of traditional medicinal applications, Papaver somniferum is also 8 known as a source of various pharmacologically highly active opiates. Consequently, their 9 10 detection from plant extracts is an important analytical task and generally addressed by methods of GC and LC-MS. However, opiates do also show structural similarities to matrix 11 12 molecules used in matrix assisted laser desorption ionization and may therefore ionize upon simple laser irradiation. Following this analytical approach, the present work thoroughly 13 evaluated the direct detection of opiates by matrix free laser desorption ionization (LDI) from 14 crude extracts of P. somniferum. The method facilitated the identification of ten reported 15 16 opiates by their molecular formulae without any chromatographic pre-purification. Moreover a principal compound analysis based on LDI-MS data permitted the correct grouping of all 17 extracts according to their inherent chemistry. Overall these results highlight the promising 18 potential of LDI-MS for the swift detection of opiates from complex mixtures. 19

20 KEYWORDS: opiates detection, Papaver somniferum, LDI-MS, LC-MS, PCA

22 INTRODUCTION:

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Papaver somniferum L. (Papaveraceae), commonly known as opium poppy, is a globally 23 cultivated medicinal plant and important source of many pharmacologically active alkaloids 24 such as codeine, thebaine, noscapine papaverine and morphine.¹ Exhibiting strong effects on 25 the central nervous system, some of these opiates are used as analgesic, antitussive and 26 antispasmodic medicines.²⁻⁴ Contrary to capsules, seeds of *P. somniferum* do not contain any 27 latex or notable amounts of alkaloids and are a popular food commodity in Central and 28 Eastern Europe. However, opiate contamination of seeds caused by insect damage and bad 29 harvesting practices is a major concern recently highlighted by European food safety 30 authorities.¹ Consequently, the rapid detection and chemical identification of opiates from 31 complex plant matrices is an important issue for food safety and generally addressed by 32 methods of HPLC-DAD, LC-MS or GC-MS.⁵⁻⁹ Any of these methods represent powerful 33 34 analytical tools however, very little is known about opium alkaloids in the context of laser desorption ionization (LDI). Some of the very few reports describe isoquinolines like 3-35 aminoquinoline and 1-hydroxyisochinolin as working matrices in matrix assisted laser 36 desorption ionization (MALDI).¹⁰⁻¹¹ As many opiates actually share close structural 37 similarities with these compounds it is reasonable to assume that they should ionize upon 38 simple laser irradiation and without matrix support. With this in mind, the present study 39 thoroughly evaluated the potential of LDI-MS for the direct detection of morphinan and 40 benzylisoquinoline alkaloids from crude methanolic caplsule extracts of P. somniferum (I). 41 Moreover results were systematically compared with concurrently performed LC-MS 42 analyses (II). Eventually, a principal component analysis (PCA) verifying whether sample 43 material could be statistically differentiated based on LDI-MS data was conducted (III). 44

46 MATERIAL AND METHODS:

47 Plant Material and Extraction.

Voucher samples from five different batches of pulverized capsules of P. somniferum were 48 collected at a confidential breeding farm and provided by a private institute that wants to 49 remain anonymous. Reference specimens of each sample are kept at the institute of SONAS. 50 Approximatively forty milligrams of powdered plant material were mixed with 1550 µL 51 52 analytical grade MeOH (Carlo Erba Reagents, Val de Reuil, France) and sonicated for 20 min. The suspension was then centrifuged for 10 min at 14800 rpm and 800 µL of the supernatant 53 were recovered. Next 800 µL of fresh MeOH were added to the plant material and the 54 extraction was repeated. Combined supernatants were dried under nitrogen yielding 4-6 mg 55 dried extract per sample. Detailed information indicating exact weights of sample and plant 56 57 material is given in Table S1 (Supplementary Information).

Laser Desorption Ionization Mass Spectrometry. Low-resolution (LR), LDI-MS 58 experiments were carried out on a Bruker Biflex III time of flight (TOF) mass spectrometer 59 (Bruker Daltonik, Bremen, Germany), equipped with a 337 nm pulsed nitrogen laser (model 60 VSL-337i, Laser Sciences Inc., Boston, MA, USA). Samples were analyzed in the reflectron 61 positive mode, within a mass range of 40-2000 Da. Acceleration voltage was set to 19 kV, 62 pulse ion extraction was 200 ns, and laser frequency was 5 Hz. Laser energy was individually 63 adapted to sample requirements and in the range of 80-85%. Stock solutions of extracts poppy 64 65 were prepared in MeOH at 20 mg/ml before being diluted to 10 mg/mL with a solution comprising MeOH and formic acid (Acros organics, Geel, Belgium) (99:1, v/v). Finally, 0.5 66 µL were deposited in quintuplicates on a MALDI plate (ground steel, MTP 384 Bruker 67 68 Daltonics, Bremen, Germany). Data were acquired in manual and automatic mode. The latter facilitated an unbiased operator independent data acquisition, which was essential for 69 consecutively performed PCA. In automatic mode, 300 single spectra from 30 random 70

locations (10 laser shots each) were acquired and summed for each sample deposition spot.
Spectra were processed by FlexAnalysis 2.0 (Bruker Daltonik, Bremen, Germany).

High-resolution (HR) experiments were performed on a Spiral-TOFTM MALDI TOF/TOF mass spectrometer (Jeol LTd., Tokyo, Japan) equipped with a pulsed laser operating at 349 nm. Experiments were performed in the positive spiral mode within mass range of 100-1000 Da. Acceleration voltage was set to 20 kV, laser frequency was 250 Hz and laser power was 59%. Signals exhibiting a signal to noise ratio (S/N) of less than ten were considered not detected. Exported spectra were processed by mMass 5.5.0 a freeware by Martin Strohalm.¹²

HPLC-ESI-MS. Experiments were performed on a Trap Esquire 3000+ instrument (Bruker, 80 Bremen, Germany) equipped with ESI/APCI Ion Trap as well as a UV-VIS wavelength 81 82 detector. Instrument parameters and experimental conditions were as follows: Collision gas, He; collision energy amplitude, 1.0 V; nebulizer and drying gas, N₂, 7 L/min; pressure of 83 nebulizer gas, 20 psi; dry temperature, 300 °C; flow rate, 0.5 mL/min; solvent split ratio 1:9; 84 85 scan range, m/z 20-1000. In addition absorption chromatograms were acquired at 230 and 280 nm. Chromatographic separation was performed according to a previously published 86 protocol with some modifications.⁷ Experiments were conducted on a Gemini C_{18} column 87 (3µm, 110 A, 150 x 4.6 mm, Phenomenex, Le Pecq, France) protected by a pre-column 88 security guard cartridge C_{18} (4 × 3 mm i.d.) from the same company. The mobile phase 89 contained a solution of ammonium bicarbonate (Servilab, Le Mans, France) at 20 mM 90 adjusted to pH 9.0 by ammonium hydroxide (Acros organics, Geel, Belgium) (solvent A), as 91 well as LC-MS grade MeOH (Honeywell, Seelze, Germany) comprising 5% of solvent A 92 93 (solvent B). Gradient conditions were as follows: 40% B (0-1 min), 40-100% B (1-30 min), 100% B (30-42 min). Signals exhibiting a S/N<10 in MS, as well as less than 0.03 mAU in 94 UV were considered not detected. 95

Data Processing and Principal Component Analysis. Data from LR LDI-MS experiments
were processed by FlexAnalysis 2.0 (Bruker Daltonik, Bremen, Germany) and mMass 5.5.0
(Martin Strohalm, Czech Republic)¹². Principal component analysis (PCA) was performed on
Origin Pro (OriginLab Cooperation, Northampton, MA, USA), using the Principal
Component Analysis for Spectroscopy software extension. Before conducting PCA, all LDIMS spectra were normalized to the base peak. A covariance analysis was performed and the
number of extracted components was two.

103 **RESULTS AND DISCUSSION:**

Preceding the current work, the concept of direct LDI-MS was successfully applied for the 104 analysis of complex mixtures of phenolic compounds.¹³ Following the same methodological 105 approach, poppy extracts were first analyzed by LR LDI-MS (Biflex III, Bruker). These 106 experiments suggested the presence of ten reported opiates, namely morphine (1), oripavine 107 (2), codeine (3), thebaine (4), corytuberine (5), reticuline (6) papaverine (7), norlaudanosine 108 (8) and/or its constitutional isomer codamine, as well as laudanosine (9) and noscapine (10) 109 (Figure 1 and 2).^{1, 14-15} Their molecular formulae were then confirmed by HR-LDI-MS 110 (Table 1). It should be noted that norlaudanosine and codamine are both reported constituents 111 of *P. somniferum*,¹⁴ but, as constitutional isomersn cannot be distinguished by their quasi-112 113 molecular ion signals. Overall results from LDI-MS revealed a considerable chemical diversity within the analyzed samples, suggesting that the method may provide a valuable tool 114 for discriminative analyses of complex mixtures of alkaloids. This aspect will be discussed in 115 detail later on in the manuscript. 116



Figure 1. Structures of opiates that have been confirmed by their molecular formulae. Both norlaudanosine and codamine ($C_{20}H_{25}NO_4$) are reported consituens of *P. somniferum*,¹⁴ but as constitutional isomers cannot be disdiguished by their quasi-molecular ions.



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Figure 2. High resolution LDI-MS spectrum obtained from the crude methanolic extract of sample 3. The following alkaloids were detected and confirmed by their molecular formulae: Morphine (1), oripavine (2), codeine (3), thebaine (4), corytuberine (5) reticuline (6), papaverine (7), codamine and/or norlaudanosine* (8) and laudanosine (9). Any compound was detected by their quasi-molecular ions [M+H]⁺. A comprehensive summary of all confirmed alkaloids in all samples is provided in Table 1. Further HR spectra of all extracts are provided in the Supporting Information section (Figure S5 A-E).

molecular	calculated	observed mass	mass error	mass error	data from	Opiates detected in sam			ple	
formula	mass [M+H] ⁺	$[\mathbf{M}+\mathbf{H}]^{+} \qquad (\mathbf{D}\mathbf{a})$		(ppm)	sample	1	2	3	4	5
C ₁₇ H ₁₉ NO ₃	286.1443	286.1437	-0.0006	-2.1	1	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{18}H_{19}NO_3$	298.1443	298.1435	-0.0008	-2.7	1	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{18}H_{21}NO_3$	300.1600	300.1596	-0.0004	-1.3	1	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{19}H_{21}NO_3$	312.1600	312.1590	-0.0010	-3.2	1	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{19}H_{21}NO_4$	328.1549	328.1536	-0.0013	-4.0	1	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{19}H_{23}NO_4$	330.1705	330.1691	0.0014	-4.2	3	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{20}H_{21}NO_4$	340.1548	340.1550	0.0002	0.6	1	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{20}H_{25}NO_4$	344.1862	344.1847	-0.0015	-4.4	3	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
$C_{21}H_{27}NO_4$	358.2018	358.2001	-0.0017	-4.7	3	\checkmark	n.d.	\checkmark	\checkmark	\checkmark
C ₂₂ H ₂₃ NO ₇	414.1552	414.1536	-0.0016	-3.9	1	\checkmark	\checkmark	n.d.	n.d	n.d.
	molecular formula C ₁₇ H ₁₉ NO ₃ C ₁₈ H ₁₉ NO ₃ C ₁₈ H ₂₁ NO ₃ C ₁₉ H ₂₁ NO ₃ C ₁₉ H ₂₁ NO ₄ C ₁₉ H ₂₃ NO ₄ C ₂₀ H ₂₅ NO ₄ C ₂₁ H ₂₇ NO ₄ C ₂₂ H ₂₃ NO ₇	molecular calculated formula mass [M+H] ⁺ C ₁₇ H ₁₉ NO ₃ 286.1443 C ₁₈ H ₁₉ NO ₃ 298.1443 C ₁₈ H ₂₁ NO ₃ 300.1600 C ₁₉ H ₂₁ NO ₃ 312.1600 C ₁₉ H ₂₁ NO ₄ 328.1549 C ₁₉ H ₂₁ NO ₄ 330.1705 C ₂₀ H ₂₁ NO ₄ 340.1548 C ₂₀ H ₂₅ NO ₄ 344.1862 C ₂₁ H ₂₇ NO ₄ 358.2018 C ₂₂ H ₂₃ NO ₇ 414.1552	molecular calculated observed mass formula mass [M+H] ⁺ [M+H] ⁺ C ₁₇ H ₁₉ NO ₃ 286.1443 286.1437 C ₁₈ H ₁₉ NO ₃ 298.1443 298.1435 C ₁₈ H ₂₁ NO ₃ 298.1443 298.1435 C ₁₉ H ₂₁ NO ₃ 300.1600 300.1596 C ₁₉ H ₂₁ NO ₃ 312.1600 312.1590 C ₁₉ H ₂₁ NO ₄ 328.1549 328.1536 C ₁₉ H ₂₁ NO ₄ 330.1705 330.1691 C ₂₀ H ₂₁ NO ₄ 340.1548 340.1550 C ₂₀ H ₂₅ NO ₄ 344.1862 344.1847 C ₂₁ H ₂₇ NO ₄ 358.2018 358.2001 C ₂₂ H ₂₃ NO ₇ 414.1552 414.1536	molecular calculated observed mass mass error formula mass [M+H] ⁺ [M+H] ⁺ (Da) C ₁₇ H ₁₉ NO ₃ 286.1443 286.1437 -0.0006 C ₁₈ H ₁₉ NO ₃ 298.1443 298.1435 -0.0008 C ₁₈ H ₂₁ NO ₃ 300.1600 300.1596 -0.0004 C ₁₉ H ₂₁ NO ₃ 312.1600 312.1590 -0.0010 C ₁₉ H ₂₁ NO ₄ 328.1549 328.1536 -0.0013 C ₁₉ H ₂₁ NO ₄ 330.1705 330.1691 0.0014 C ₂₀ H ₂₁ NO ₄ 340.1548 340.1550 0.0002 C ₂₀ H ₂₁ NO ₄ 344.1862 344.1847 -0.0015 C ₂₁ H ₂₇ NO ₄ 358.2018 358.2001 -0.0017 C ₂₂ H ₂₃ NO ₇ 414.1552 414.1536 -0.0016	molecular calculated observed mass mass error mass error formula mass [M+H] ⁺ [M+H] ⁺ (Da) (ppm) C ₁₇ H ₁₉ NO ₃ 286.1443 286.1437 -0.0006 -2.1 C ₁₈ H ₁₉ NO ₃ 298.1443 298.1435 -0.0008 -2.7 C ₁₈ H ₁₉ NO ₃ 298.1443 298.1435 -0.0004 -1.3 C ₁₉ H ₂₁ NO ₃ 300.1600 300.1596 -0.0010 -3.2 C ₁₉ H ₂₁ NO ₃ 312.1600 312.1590 -0.0013 -4.0 C ₁₉ H ₂₁ NO ₄ 328.1549 328.1536 -0.0013 -4.0 C ₁₉ H ₂₃ NO ₄ 330.1705 330.1691 0.0014 -4.2 C ₂₀ H ₂₃ NO ₄ 340.1548 340.1550 0.0002 0.6 C ₂₀ H ₂₅ NO ₄ 344.1862 344.1847 -0.0015 -4.4 C ₂₁ H ₂₇ NO ₄ 358.2018 358.2001 -0.0016 -3.9 C ₂₂ H ₂₃ NO ₇ 414.1552 414.1536 -0.0016 -3.9	molecular calculated observed mass mass error mass error data from formula mass [M+H] ⁺ [M+H] ⁺ (Da) (ppm) sample C ₁₇ H ₁₉ NO ₃ 286.1443 286.1437 -0.0006 -2.1 1 C ₁₈ H ₁₉ NO ₃ 298.1443 298.1435 -0.0008 -2.7 1 C ₁₈ H ₂₁ NO ₃ 300.1600 300.1596 -0.0004 -1.3 1 C ₁₉ H ₂₁ NO ₃ 312.1600 312.1590 -0.0010 -3.2 1 C ₁₉ H ₂₁ NO ₄ 328.1549 328.1536 -0.0013 -4.0 1 C ₁₉ H ₂₁ NO ₄ 330.1705 330.1691 0.0014 -4.2 3 C ₂₀ H ₂₁ NO ₄ 340.1548 340.1550 0.0002 0.6 1 C ₂₀ H ₂₅ NO ₄ 344.1862 344.1847 -0.0017 -4.4 3 C ₂₁ H ₂₇ NO ₄ 358.2018 358.2001 -0.0017 -4.7 3 C ₂₁ H ₂₃ NO ₇ 414.1552 414.1536 -0.0016 -3.9 1	molecularcalculatedobserved massmass errormass errordata from I formulamass [M+H] ⁺ [M+H] ⁺ (Da)(ppm)sample1 $C_{17}H_{19}NO_3$ 286.1433286.1437-0.0006-2.11 \checkmark $C_{18}H_{19}NO_3$ 298.1433298.1435-0.0008-2.71 \checkmark $C_{18}H_{21}NO_3$ 300.1600300.1596-0.0004-1.31 \checkmark $C_{19}H_{21}NO_3$ 312.1600312.1590-0.0010-3.21 \checkmark $C_{19}H_{21}NO_4$ 328.1549328.1536-0.0013-4.01 \checkmark $C_{19}H_{21}NO_4$ 330.1705330.16910.0014-4.23 \checkmark $C_{20}H_{21}NO_4$ 340.1548340.15500.00020.61 \checkmark $C_{20}H_{21}NO_4$ 344.1862344.1847-0.0015-4.43 \checkmark $C_{21}H_{27}NO_4$ 358.2018358.2001-0.0016-3.91 \checkmark	molecular calculated observed mass mass error mass error data from $Oppates oppates oppa$	molecular calculated observed mass mass error mass error data from C formula mass [M+H] ⁺ [M+H] ⁺ (Da) (ppm) sample 1 2 3 $C_{17}H_{19}NO_3$ 286.1433 286.1437 -0.0006 -2.1 1 \checkmark \checkmark \checkmark $C_{18}H_{21}NO_3$ 298.1433 298.1435 -0.0008 -2.7 1 \checkmark \checkmark \checkmark $C_{18}H_{21}NO_3$ 300.1600 300.1596 -0.0004 -1.3 1 \checkmark \checkmark \checkmark $C_{19}H_{21}NO_3$ 312.1600 312.1590 -0.0010 -3.2 1 \checkmark \checkmark \checkmark $C_{19}H_{21}NO_4$ 328.1549 328.1536 -0.0013 4.00 11 \checkmark \checkmark \checkmark $C_{19}H_{21}NO_4$ 340.1550 0.0014 4.22 3 \checkmark \checkmark $C_{20}H_{23}NO_4$ 340.1548 340.1550 0.0015 4.44 3 \checkmark \checkmark C_{2	molecular calculated observed mass mass error mass error data from Cpites detected in sample formula mass [M+H] ⁺ [M+H] ⁺ (Da) (ppm) sample 1 2 3 4 $C_{17}H_{19}NO_3$ 286.1433 286.1437 -0.0006 -2.1 1 \checkmark \checkmark \checkmark \checkmark $C_{18}H_{19}NO_3$ 298.1443 298.1435 -0.0008 -2.7 1 \checkmark \checkmark \checkmark \checkmark $C_{18}H_{21}NO_3$ 300.1600 300.1596 -0.0004 -1.3 1 \checkmark \checkmark \checkmark \checkmark $C_{19}H_{21}NO_3$ 312.1600 312.1590 -0.0010 -3.2 1 \checkmark \checkmark \checkmark \checkmark $C_{19}H_{21}NO_4$ 328.1549 328.1536 -0.0013 4.0 1 \checkmark \checkmark \checkmark $C_{19}H_{23}NO_4$ 340.1550 0.0002 0.6 1 \checkmark \checkmark \checkmark $C_{20}H_{25}NO_4$ 344.1862 344.1847

131	Table 1 High resolution LDI-MS	performed on samples 1-5
TO T	Lasie I. Ingh lessiation DB1 his	

 \checkmark detected, n.d. not detected (S/N<10). *Constitutional isomers reported for *P. somniferum*.¹⁴

		sample	1	sample 2 sample 3			3	5	sample	4	sample 5				
compound	LC- UV	LC- MS	LDI- MS	LC- UV	LC- MS	LDI- MS	LC- UV	LC- MS	LDI- MS	LC- UV	LC- MS	LDI- MS	LC- UV	LC- MS	LDI- MS
morphine (1)	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
oripavine (2)	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	n.d.	\checkmark							
codeine (3)	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
thebaine (4)	со	\checkmark	\checkmark	со	\checkmark	\checkmark	co	\checkmark	\checkmark	co	\checkmark	\checkmark	со	\checkmark	\checkmark
corytuberine (5)	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	n.d.	\checkmark	\checkmark
reticuline (6)	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark
papaverine (7)	со	\checkmark	\checkmark	co	\checkmark	\checkmark	co	\checkmark	\checkmark	со	n.d.	\checkmark	со	n.d.	\checkmark
norlaudanosine*(8) codamine*	~	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	~	\checkmark	\checkmark
laudanosine (9)	\checkmark	\checkmark	\checkmark	n.d.	\checkmark	n.d.	\checkmark								
noscapine (10)	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	n.d.	\checkmark	\checkmark	n.d.	n.d.	n.d.	n.d.
Except for noscapine (10) (sample 3 and 4) and laudanosine (9) (sample 2), HR-LDI-MS detected any of the compounds observed by LC-UV-															

134 **Table 2.** Comparison of results from LC-UV (230 nm), LC-MS and HR-LDI-MS experiments

MS. On the other hand, papaverine (7) was exclusively detected by LDI-MS in sample 4 and 5. A comprehensive summary of all LC-UV-MS

137 chromatograms is given in the Supporting Information.

With regards to common laboratory practices, the analysis of opiates is mostly performed by 138 methods such as GC-MS, HPLC-UV and LC-UV-MS.⁵⁻⁹ One major advantage of these 139 hyphenated methods is, that complex mixtures undergo chromatographic separation before 140 entering UV or MS detectors. Consequently constituents are individually ionized, or, in case 141 of co-elution, as mixtures of very few compounds. With respect to LDI, the situation is 142 completely different, as all compounds concurrently compete in the ionization process. This 143 may induce so called "analyte suppression effects" (ASE). Initially described for the MALDI 144 process,¹⁶⁻¹⁸ ASE may probably occur in matrix free LDI too, as both methods share similar 145 ionization mechanisms. In order to address this issue and to compare LDI-MS to classical 146 analytical approaches, all samples were systematically analyzed by LC-UV-MS and results 147 are summarized in Table 2. Except for the minor alkaloid noscapine (sample 3, 4) and trace 148 amounts of laudanosine (9) in sample 2, LDI-MS detected any compound observed by LC-149 150 MS. On the other hand, solely LDI-MS permitted the detection of papaverine in samples 4 and 5. Compared to LC-UV, both LC-MS and LDI-MS exhibited higher sensitivity and were 151 not impacted by the co-elution of papaverine and thebaine observed under given 152 153 chromatographic conditions. A comprehensive summary of all LC-UV-MS experiments is provided in the Supporting Information (file: LC-UV-MS data). 154

With regards to sample processing, LDI-MS did not require any sample pre-conditioning or 155 method optimization steps and yielded instant results. While the automated acquisition of a 156 cumulative LDI spectrum was accomplished within fifty seconds, forty-two minutes were 157 required for an LC-UV-MS experiment. Moreover LDI-MS permitted the use of any solvent 158 for sample solubilization and yielded distinct fingerprint spectra with highly specific relative 159 signals intensity ratios for each sample. (Figure S3 and S4) These findings confirm results 160 previously described for phenolic compounds, which are discussed in detail in the original 161 publication.¹³ 162

Concluding the experimental approach, a PCA based on LDI-MS data was performed. Results 163 from this experiment are summarized in Figure 3 and show the correct grouping of all 164 samples according to their chemical composition. As outlined in the Material and Methods 165 section, samples were deposited in quintuplicates, so each data point represent one cumulative 166 spectrum of 300 laser shots randomly acquired from one sample deposition spot. All data 167 points of all samples were located within the 95% ellipse of confidence, confirming the 168 robustness of the method. This is once more in line with previous observations made for 169 phenolic compounds.¹³ Data points of sample 1 and 2 are located rather closely in the PCA 170 with some overlay of their ellipses of confidence, suggesting a similar chemical composition. 171 This also confirms the visual impression observed when looking at single spectra of these 172 samples (Figure S1 and S2 Supporting information). 173



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Figure 3. Principal component analysis of poppy samples 1-5 based on their LR LDI-MS spectra. Samples were analyzed in quintuplicates and each data point represent one cumulative spectrum of 300 laser shots that were randomly acquired from ten different locations of one sample spot.

For the time being very little is known about the ionization of opiates by laser desorption. Some reports describe the MALDI detection of morphine from bodily fluids, protein conjugates and pharmaceutical preparations,¹⁹⁻²¹ as well as from a mono-component solution ²² and TLC surfaces.²³ Like LDI, MALDI yields instant results and may even provide increased sensitivity. On the other hand MALDI is inevitably linked to the formation of matrix ions in the low mass region, which may easily superpose analyte signals and prevent correct spectra interpretation. In addition, redundant matrix ions do severely impair statistical 187 separation by PCA. Bypassing the problem of matrix interference, surface assisted LDI 188 (SALDI) is commonly used for detecting small molecules. The method replaces the matrix by 189 nanotextured materials such as silicon or various specifically structured metal surfaces in 190 order to assist the ionization process. With respect to opiates, one report describes the use of 191 carbon nano-coatings for inducing LDI of papaverine.²⁴ However present results have shown 192 that opiates do sufficiently ionize by simple laser irradiation and without the support of any 193 nano-textured material or surface.

In summary the present work has discussed the inherent LDI properties of morphinan and 194 benzylisoquinoline alkaloids as well as their simultaneous detection from crude extracts of 195 P. somniferum. Permitting the detection of ten opiates by their molecular formulae, results 196 from LDI-MS were comparable with those obtained by LC-MS. Moreover, all poppy samples 197 exhibited highly specific "fingerprint" spectra allowing their unambiguous identification as 198 well as their discrimination by PCA. Facilitating an automated and rapid qualitative analysis 199 of large sample batches, LDI-MS may provide an interesting analytical tool for industrial 200 application. 201

ABBREVIATIONS:

- 204 APCI Atmospheric Pressure Chemical Ionization
- 205 ASE Analyte Suppression Effects
- 206 ESI Electro Spray Ionization
- 207 HPLC High Performance Liquid Chromatography
- 208 LC-MS Liquid Chromatography coupled with Mass Spectrometry
- 209 LDI-MS Laser Desorption Ionization coupled with Mass Spectrometry
- 210 MALDI Matrix Assisted Laser Desorption Ionisation
- 211 PCA Principle Component Analysis
- 212 TOF Time of Flight

213 UV Ultra Violet (used for absorption chromatograms in combination with HPLC)

214 ACKNOWLEDGEMENT:

215 SUPPORTING INFORMATION:

Low (Figures S1-S3) and high resolution (Figure S5 A-E) LDI-MS spectra, as well as a detailed summary of extraction yields (Table S1) are provided for samples 1-5. Further a detailed analysis of the repeatability of LDI-MS experiments is given (Figure S3, S4). A complete summary of all LC-UV-MS experiments conducted on samples 1-5 is provided as well. Finally, original, high resolution LDI-MS spectrum files of samples 1-5 were exported. These files can be opened and analyzed with mMass (freeware).

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