

Utility of the *Leptospermum scoparium* Compound Lepteridine as a Chemical Marker for Manuka Honey Authenticity

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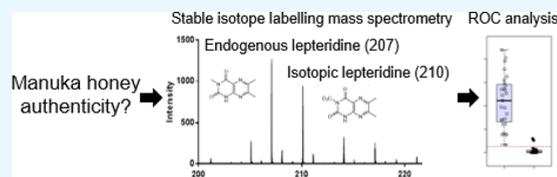


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ABSTRACT: Manuka honey is a premium food product with unique antimicrobial bioactivity. Concerns with mislabeled manuka honey require robust assays to determine authenticity. Lepteridine is a *Leptospermum*-specific fluorescent molecule with potential as an authenticity marker. We describe a mass spectrometry-based assay to measure lepteridine based on an isotopically labeled lepteridine standard. Using this assay, lepteridine concentrations in manuka honey samples strongly correlated with concentrations quantitated by either high-performance liquid chromatography-ultraviolet (HPLC-UV) or fluorescence. A derived minimum lepteridine threshold concentration was compared with the New Zealand regulatory definition for manuka honey to determine “manuka honey” authenticity on a set of commercial samples. Both methods effectively distinguished manuka honey from non-manuka honeys. The regulatory definition excludes lepteridine but otherwise includes the quantification of multiple floral markers together with pollen analysis. Our findings suggest that the quantification of lepteridine alone or in combination with leptosperin could be implemented as an effective screening method to identify manuka honey, likely to achieve an outcome similar to the regulatory definition.



1. INTRODUCTION

Manuka honey from New Zealand is a high-value natural food product and also has biomedical applications in wound healing. One bioactivity that distinguishes manuka honey from other non-*Leptospermum* honeys is a non-peroxide antibacterial activity described as unique manuka factor (UMF),¹ which has been attributed to the presence of methylglyoxal (MGO).² MGO content increases in manuka honey over time as it is derived from the nectar precursor dihydroxyacetone (DHA), which spontaneously converts into MGO during storage and controlled heating.^{3,4}

Unfortunately, because of its premium market value, non-manuka honeys are sometimes packaged and sold as manuka honey for increased commercial gain. There is therefore a need for robust assays to ensure that manuka honey is true to label. In addition, New Zealand kanuka and manuka honeys share the same geographical location and flowering season, which can make it difficult to distinguish these honey varieties.^{5,6} To address these issues, the New Zealand Ministry for Primary Industries (MPI) has developed a regulatory definition for monofloral and multifloral manuka honey using chemical fingerprinting together with pollen DNA polymerase chain reaction (PCR) analysis.^{7–10} These criteria are based on five key attributes: the presence of certain levels of 2'-methoxyacetophenone (2'-MAP), 2-methoxybenzoic acid (2-MB), 4-hydroxyphenyllactic acid (4-HPA), 3-phenyllactic acid (3-PA), and DNA from manuka pollen.

Our studies have demonstrated that two *Leptospermum* nectar-derived compounds, leptosperin and lepteridine, can also be utilized to determine manuka honey authenticity.^{11–15} Leptosperin and lepteridine are not currently included in the MPI definition yet are responsible for two unique fluorescent signatures in manuka honey, called MM1 and MM2.^{13,16} Principal component analysis on a range of manuka and non-manuka honeys show that both leptosperin and lepteridine together with the MPI-recommended compounds, 2-MB and 4-HPA, are the most significant contributors to the PC1 axis.¹¹ Receiving operating characteristic curve (ROC) analyses for leptosperin and lepteridine, which are stable in honey over time unlike 2-MB,¹³ show a strong discriminatory power for distinguishing manuka honey.

One unanswered question is how the use of leptosperin/lepteridine performs relative to the MPI definition in discerning authentic manuka honeys from commercial manuka honey products that may or may not adhere to label claims. In this study, we focus on lepteridine and hypothesize that its quantitation potentially provides a simple and high-throughput mode of measurement to determine manuka honey authentic-

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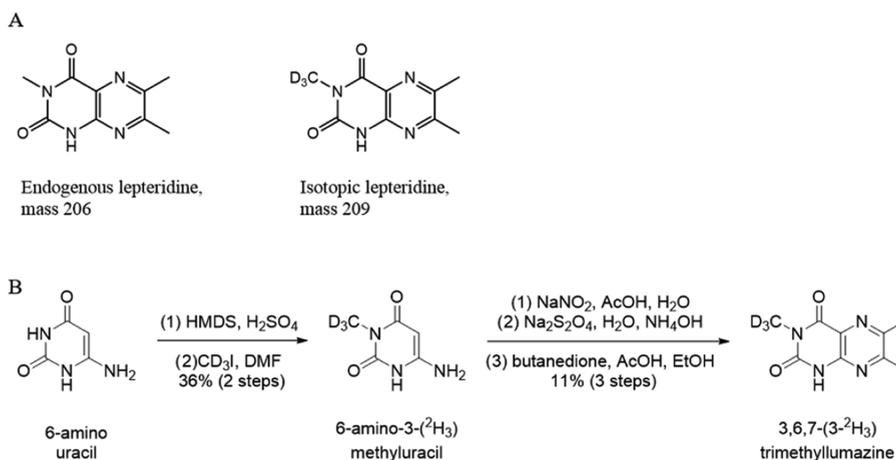


Figure 1. (A) Molecular structure of endogenous lepteridine (left) and the synthesized lepteridine isotope (right). (B) Total synthesis of isotopic lepteridine.

ity. As mass spectrometry analyses are well established in commercial honey testing laboratories, we developed a technique for determining lepteridine content in honey based on a liquid chromatography tandem mass spectrometry (LC–MS/MS) approach utilizing an isotopically labeled lepteridine internal standard. Utilizing these various analyses for lepteridine, we first determined threshold values through measurements on manuka honey samples. We then quantified MPI-specified chemical markers (the pollen DNA marker was not included) versus lepteridine-specific analyses on a set of 33 “manuka” honey samples that were purchased from the commercial market of Singapore. Previous analyses of 17 of these 33 samples showed that 3 were either not manuka in origin or were potentially adulterated.¹⁶

2. RESULTS AND DISCUSSION

2.1. Synthesis of Isotopically Labeled Lepteridine.

Mass spectrometry is a powerful analytical tool in food science that has been applied in the analyses of numerous foods including milk products^{17,18} fruits,^{19,20} and beverages.²¹ Most of these methods employ an isotopically labeled standard to compensate for matrix effects between samples.²² Absolute quantification can be achieved by comparing relative ion abundance between the paired compounds. To reduce interference from background ions, quantification can be performed on specific fragment ions using tandem mass spectrometry.²³

We utilized these principles to generate isotopically labeled lepteridine with the same physicochemical properties as native lepteridine but which can be identified by mass spectrometry due to the relative atomic mass difference. The use of a heavy isotope standard to quantitate lepteridine concentration in manuka honey required the synthesis of a suitable isotopically enriched analogue of lepteridine. Replacement of the methyl group at N-3 of lepteridine with a deuteromethyl group provided a mass difference of 3 amu, which was sufficient to differentiate the two isotopes by mass spectrometry (Figure 1A). The use of iodomethane-D₃ in place of iodomethane¹⁴ facilitated the synthesis of isotopic lepteridine (Figure 1B).

2.2. Validation of Isotopic LC–MS/MS Lepteridine Quantification. Figure 2A,B shows the mass spectrum of typical manuka honey before and after supplementation of isotopic lepteridine, respectively. As expected, endogenous and

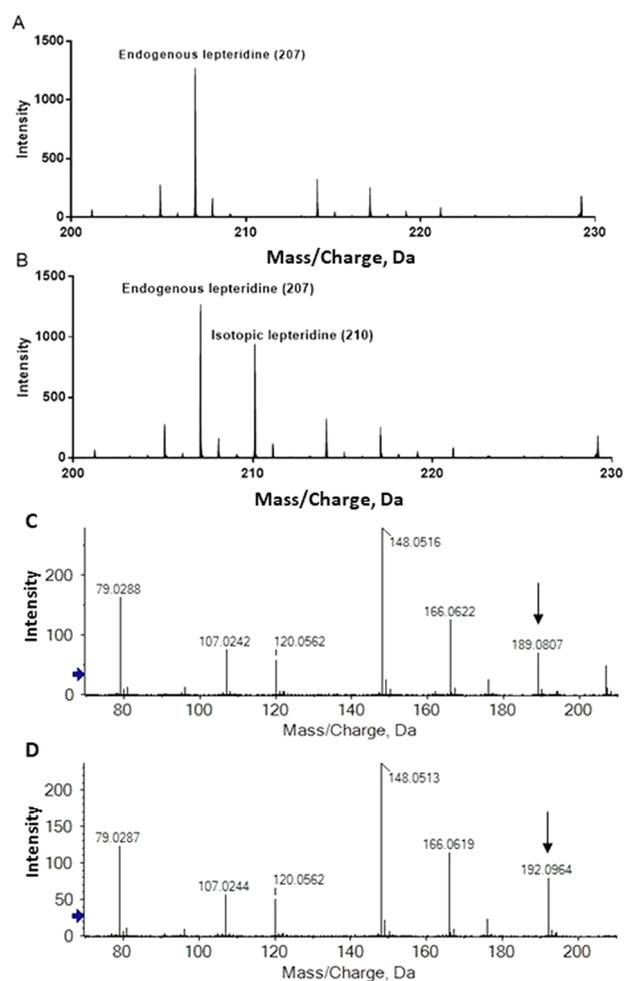


Figure 2. Mass spectrum of a typical manuka honey sample before (A) and after (B) supplementation with isotopic lepteridine. MS/MS spectrum of endogenous lepteridine (C) and isotopically labeled lepteridine (D). Black arrows denote a 3 amu shift in one of the fragment ions.

isotopically labeled lepteridine displayed almost identical MS/MS spectra and co-eluted at the same time. No significant interfering peaks were identified from endogenous compounds in manuka honey between m/z 210 and 212 (Figure 2A). The

3 amu mass difference between endogenous and isotopically spiked leperidine was also clearly discernible in the mass spectrum (Figure 2B).

Figure 2C,D shows individual MS/MS spectra for endogenous and isotopic leperidine, respectively. The two species were differentiated by a 3 amu shift in the fragment ion from m/z 189 (Figure 2C) to m/z 192 (Figure 2D). The most abundant common fragment ion observed at m/z 148.05 lacked the heavy isotopes and was employed for leperidine quantification. The high ion count relative to the noise of this fragment ion minimized background interference. The final testing concentration of manuka honey was determined at 0.2% w/v to maintain an adequate signal-to-noise ratio and minimize instrument contamination with residual honey sugars.

2.2.1. Linearity. Linearity measures the proportionality of a method's analytical response to the amount of analyte in the sample. The linearity of the LC–MS/MS method in this study was evaluated by regression statistics based on a 7-point calibration curve prepared in duplicate. The calibration concentrations of leperidine ranged from 1 to 100 ng/mL, corresponding to a 0.5–50 mg/kg concentration range in undiluted manuka honey, while the isotopically labeled leperidine was present at 20 ng/mL. The calibration curve of leperidine over this concentration range was established by weighted ($1/x^2$) linear regression analysis, where the coefficient correlation (r) from three independent analyses established over different days was determined to be ≥ 0.995 (Figure S1).

The calibration curve was linear with respect to the concentration range tested. Residual plots for each of the calibration curves did not show any obvious trend and appeared to be randomly distributed across the concentration range tested. All data points included in the calibration curves were within measurement accuracy of $\pm 15\%$ (Table S1).

2.2.2. Repeatability, Precision, and Reproducibility. Repeatability of the LC–MS/MS method was determined by supplementing leperidine-free clover honey with leperidine at three different concentrations spanning the working range in this study (3, 40, and 80 ng/mL). The analysis was performed using six replicate preparations, made independently over three different days. Precision was assessed by intraday repeatability and expressed as the percentage coefficient of variation (CV). The method showed good precision with CV ranging from 4.5 to 7.7, 5.1 to 7.4, and 2.4 to 9.0% at 3, 40, and 80 ng/mL measured over 3 days, respectively (Supporting Information Table S2). Interday repeatability was also assessed to determine the reproducibility of the method over time. Leperidine was reproducibly measured over three different days with overall CVs of 7.9, 6.4, and 5.8% at 3, 40, and 80 ng/mL, respectively (Supporting Information Table S3).

2.2.3. Sensitivity (Limit of Detection, LOD, and Limit of Quantification, LOQ). Detection sensitivity is expressed in terms of limit of detection (LOD) and limit of quantification (LOQ). The LOD of the method was calculated based on 3 times standard deviation (SD) of analyte area under curve (AUC) from 10 blank honey runs, and the LOQ was based on 10 times of this standard deviation.

The standard deviation (SD) of the AUC for the leperidine fragment ion from 10 blank clover honey samples was determined to be 6.3 counts, giving a theoretical LOD peak area of 18.8 counts ($3 \times \text{SD}$) and a theoretical LOQ peak area of 62.5 counts ($10 \times \text{SD}$). To allow a generous margin of error,

the working range for this method was set to 1–100 ng/mL (corresponding to 0.5–50 mg/kg leperidine in raw honey) for which the lowest calibration standard gave a mean AUC of 144 counts across the three batches ($n = 6$, intensity range 101–211 counts). This is more than double the required $10 \times \text{SD}$ value for the LOQ. The LOD of this method was calculated to be far lower at 0.15 ng/mL (corresponding to 0.075 mg/kg raw honey). Given that the expected minimum threshold of leperidine concentration in genuine manuka honey is around 2.5 mg/kg, this working range was deemed more than sufficient to discriminate manuka honey from non-manuka honey samples.

2.2.4. Interference. The absence of interference was demonstrated by a lack of signal for both leperidine and its isotope-labeled counterpart in nonspiked clover honey and a lack of an isotopic leperidine signal in manuka honey (Figure S2). Additionally, supplementation of the heavy isotopic leperidine into clover honey did not interfere with the leperidine signal (Figure S3).

2.2.5. Matrix Effect and Recovery. Matrix effects and recovery were assessed by comparison of the solution and honey supplemented with leperidine at 3, 40, and 80 ng/mL. Matrix recovery was 97.6–104% at these concentrations, with no significant change in background signals for the leperidine peaks in honey compared to those in the solution (Table S4).

2.3. Comparison between Leperidine Quantification Using LC–MS/MS, High-Performance Liquid Chromatography-Ultraviolet (HPLC-UV), and Fluorescence Methods. Authentic field-collected manuka honey samples harvested throughout the North Island of New Zealand were analyzed to quantitate endogenous leperidine concentrations (Table 1). Mass spectrometry analyses employing isotopically labeled leperidine as an internal standard yielded leperidine concentrations ranging between 3 and 44 mg/kg. These leperidine concentrations correlated well with leperidine concentrations quantified by either HPLC (Figure 3A, $R^2 = 0.9517$) or by fluorescence at the MM2 wavelength (Figure 3B, $R^2 = 0.8995$) on the same honey set.

Compared to quantitation by HPLC-UV,¹⁵ leperidine concentrations were reduced by approximately 24% when measured using mass spectrometry. This difference is likely due to the presence of other UV-absorbing compounds co-eluting with leperidine under the same HPLC peak. We previously identified the presence of these compound(s) by ^1H NMR spectra of leperidine fractions purified by HPLC (B.J. Daniels, unpublished observations). Ultimately, structural elucidation of leperidine was reliant on direct-phase preparative thin-layer chromatography (TLC) to remove these species,¹⁴ one of which was identified as 6,7-dimethyl-2,4(1*H*,3*H*)-pteridinedione.²⁴ This compound also fluoresces at the leperidine signature MM2 wavelength²⁴ and so will also contribute to the MM2 fluorescent signature. Based on this information, mass spectrometry is the more accurate approach for leperidine quantification in manuka honey.

2.4. Determining Manuka Honey Authenticity Using Leperidine. Using ROC analyses of field-collected manuka honey samples (Table 1, samples 1–27) against non-manuka honey samples (Table 1, samples 28–53), a minimum leperidine threshold concentration of 2.6 mg/kg was derived, as measured by mass spectrometry (Figure 4A). This value is similar to a previous minimum leperidine threshold value of 2.1 mg/kg as quantified by HPLC-UV using a different set of field-collected manuka honey samples.¹¹ This threshold

Table 1. Field-Collected Honey Samples

sample	honey type	geographic origin	DHA (mg/kg)	MGO (mg/kg)	lepteredine (mg/kg) ^a
1	Manuka	Northland	3830	399	43.9
2	Manuka	Northland	3430	401	13.9
3	Manuka	Northland	3570	300	29.6
4	Manuka	Northland	4210	542	24.5
5	Manuka	Northland	>4590	454	27.0
6	Manuka	Northland	3800	395	22.3
7	Manuka	Northland	2552	862	25.3
8	Manuka	Taupo	2530	219	27.3
9	Manuka	Taupo	3340	385	26.8
10	Manuka	Taupo	>4570	451	39.2
11	Manuka	Taupo	>4550	371	28.6
12	Manuka	Taupo	3840	332	31.0
13	Manuka	Taupo	2830	511	19.4
14	Manuka	Waikato	2098	755	12.0
15	Manuka	Waikato	2083	782	14.0
16	Manuka	Waikato	2085	710	14.4
17	Manuka	East Coast	1331	397	16.6
18	Manuka	East Coast	1073	482	32.5
19	Manuka	East Coast	1066	476	7.66
20	Manuka	East Coast	1081	514	7.79
21	Manuka	East Coast	2097	753	12.3
22	Manuka	Wairarapa	2080	143	3.36
23	Manuka	Wairarapa	1483	103	8.09
24	Manuka	Wairarapa	1854	143	7.69
25	Manuka	Wairarapa	957	61	40.6
26	Manuka	Wairarapa	1413	139	19.1
27	Manuka	Wairarapa	1910	125	35.7
28	Kanuka	Taupo	<20	11	<LOD
29	Kanuka	Taupo	<20	10	<LOQ
30	Kanuka	Taupo	620	155	5.23
31	Kanuka	Northland	92	36	1.55
32	Kanuka	Northland	437	104	5.94
33	Kanuka	Northland	<20	10	<LOQ
34	Clover	Southland	<20	8	<LOD
35	Clover	Southland	<20	7	<LOD
36	Clover	Taupo	114	30	0.76
37	Clover	Southland	<20	6	<LOD
38	Clover	Southland	568	50	0.65
39	Clover	Southland	<20	9	<LOD
40	Clover	Southland	<20	6	<LOD
41	Clover	Taupo	287	34	1.58
42	Clover	Southland	222	39	<LOQ
43	Clover	Southland	318	33	<LOQ
44	Tawari	Taupo	35	12	<LOQ
45	Tawari	Taupo	134	20	0.74
46	Tawari	Taupo	<20	11	<LOQ
47	Pohutukawa	Northland	175	30	1.06
48	Pohutukawa	Northland	203	37	1.19
49	Pohutukawa	Northland	<20	7	<LOD
50	Kamahi	Taupo	<20	11	<LOD
51	Kamahi	Taupo	<20	7	<LOQ
52	Kamahi	Southland	<20	5	<LOD
53	Rewarewa	Taupo	274	38	1.794

^aDHA and MGO concentrations were supplied by Comvita NZ Limited; lepteredine concentrations were quantified using the LC–MS/MS isotopic method.

lepteredine concentration corresponded to a minimum fluorescence threshold of 537 RFU with the fluorescent plate reader utilized in the laboratory (Figure 4B). The respective

areas under the curves (AUCs = 0.997 and 0.996, Figure 4A,B, left panel) support the accuracy of these tests. The cutoff points (Figure 4A,B, right panels) were determined based on the positive likelihood ratio.

Minimum lepteredine threshold concentrations derived by ROC analyses using either mass spectrometry or fluorescence were then used to determine the authenticity of 33 commercial samples purchased from Singapore in 2016 (Table 2). These honeys were all labeled as manuka in market; however, a previous analysis of 17 of these samples found three were not true to label and incorrectly marketed as manuka honey.¹⁶ Compositional analyses of these 33 honey samples are shown in Table 2. Based on the minimum lepteredine threshold derived by either mass spectrometry or fluorescence, samples 24–33 and 22–33 were identified as mislabeled, respectively. The concentrations of commonly recognized manuka honey markers are provided for reference (Table 2).

We then compared these findings for lepteredine with the chemical markers employed in the MPI definition for the determination of manuka honey⁹ based on the presence of 2'-MAP, 2-MB, 4-HPA, and 3-PA (Table 2). As all commercial samples were labeled as “manuka honey”, we only applied the monofloral MPI definition. These classifications are mostly in agreement with the lepteredine-based analyses using the derived mass spectrometry and fluorescence minimum threshold values of 2.6 mg/kg and 537 RFU, respectively. Nevertheless, there were also some differences in classification between the MPI or lepteredine-based criteria for seven of the 33 honey samples (samples 18–24). MPI-based criteria also identified honey samples 22 and 23 as mislabelled, which is consistent with the lepteredine minimum threshold based on fluorescence but not mass spectrometry.

Overall, both MPI and lepteredine-based criteria compared well across the 33 honey samples labeled as manuka with 79% (26/33) agreement using mass spectrometry quantification and 85% (28/33) using fluorescence measurement. One caveat is that DNA pollen analysis was not available for these honey samples. Consequently, the MPI authenticity criteria is based solely on the implemented chemical marker compounds. For comparison, leptosperin concentrations in these honey samples are shown in Table 2. Analyzing these honey samples based on leptosperin-based criterion alone with a minimum threshold of 94 mg/kg¹¹ classified all but four samples (30–33) as monofloral honeys, equating to a 52% agreement with the MPI-based criteria. Thus, at least on this commercial honey set, classification based solely on lepteredine best aligns with MPI-based criteria.

In summary, our findings demonstrate the application of the chemical marker, lepteredine, in manuka honey authentication as quantified by either HPLC-UV, LC–MS/MS, and fluorescence spectroscopy. Notably, outcomes based on lepteredine alone compare favorably with the MPI regulatory definition to identify mislabelled manuka honey. Potentially, the isotopic mass spectrometry method described here for lepteredine could be incorporated together with analyses of other established chemical markers using LC/MS capability, which is commonly embedded within current honey testing facilities. Mass spectrometry offers increased specificity and sensitivity over other types of analysis methods, and the isotopic method could be refined further through optimization of flow rates and column selection to decrease cycle time.

Fluorescence quantitation of lepteredine content in honey also offers a convenient method for fieldwork. The current

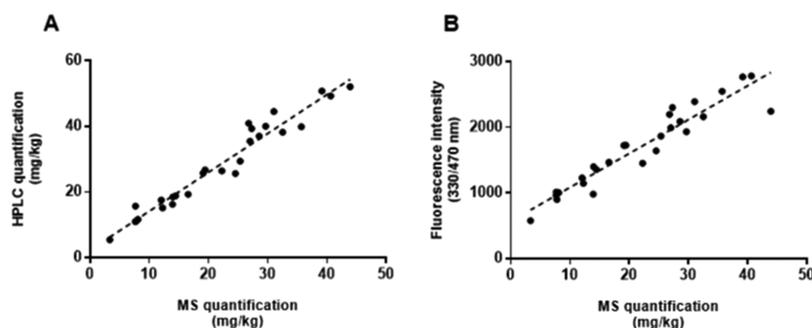


Figure 3. (A) Correlation between leperidine concentration quantified by LC–MS/MS and HPLC ($R^2 = 0.9517$). (B) Correlation between leperidine concentration quantified by LC–MS/MS and fluorescence intensity at 330–470 nm ($R^2 = 0.8995$). $n = 27$.

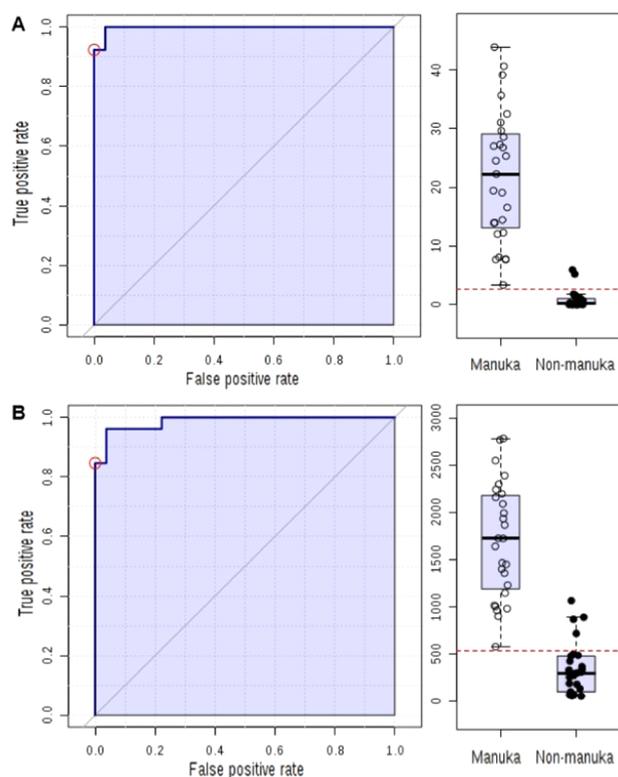


Figure 4. ROC analysis for manuka honey and non-manuka honey. (A) Leperidine concentration measured using mass spectrometry (AUC = 0.997). (B) Signature fluorescence of leperidine at ex330–em470 nm (AUC = 0.996).

MPI regulatory definition was not designed to be applied in the field; it requires quantification by mass spectrometry of four chemical markers combined with quantitative polymerase chain reaction (qPCR) analysis of pollen DNA, which also requires specialized expertise. In comparison, leperidine, which is stable over time in contrast to some other manuka markers currently in use, can be easily and rapidly quantified by fluorescence using a conventional plate reader. Here, an instrument housed on-site could be calibrated from a set of honey standards with known leperidine content to generate a fluorescence standard curve from which a minimum threshold value can be derived. In this way, the unique fluorescence signature of leperidine¹⁵ potentially, together with leptosperin,¹² offers low-cost and high-throughput screening opportunities for on-site honey and nectar analyses, particularly with the potential development of portable fluorescent devices.

3. MATERIALS AND METHODS

3.1. Honey Samples. Three honey sample collections were kindly provided by Comvita NZ Ltd. (Te Puke, New Zealand). The first comprised 27 field manuka honey samples collected from five different regions throughout New Zealand in 2009 (Table 1, samples 1–27). The floral source of these field-collected manuka honey samples was confirmed by recognized experts and beekeepers based on site analysis and flowering season. The second collection was a set of non-manuka honey samples comprising kanuka, clover, tawari, pohutukawa, kamahi, and rewarewa (Table 1, samples 28–53). These non-manuka honey samples were collected by Oritain Global Ltd. (Otago, New Zealand) in 2011, and corresponding compositional data was established by Hills Laboratory Ltd. (Hamilton, New Zealand). The third collection comprised a set of honey samples purchased from a commercial market in Singapore that were all labeled as manuka honey (Table 2). These honeys were purchased by the UMF Honey Association (UMFHA). It needs to be noted that the UMFHA samples were collected from the market before the MPI regulatory definition came into effect. All honey samples were stored at 4 °C in dark conditions prior to analysis.

3.2. Honey Sample Preparation. All honey samples were brought to room temperature and thoroughly mixed to ensure homogeneity. Manuka honey solutions (0.2% w/v) were prepared by diluting manuka honey 500-fold in 0.1% formic acid. Isotopic leperidine was spiked into each honey sample to achieve a final concentration of 20 ng/mL. All samples were thoroughly vortexed until completely dissolved in 0.1% formic acid solution.

3.3. Synthesis of Isotopically Labeled Leperidine. All reactions were carried out in flame- or oven-dried glassware under a dry nitrogen atmosphere. All reagents were purchased as reagent grade and used without further purification. Dimethylformamide was degassed and dried using an LC Technical SP-1 solvent purification system. Ethanol was distilled over $Mg(OEt)_2$. Ethyl acetate, methanol, and petroleum ether were distilled prior to use. All other solvents were used as received, unless stated otherwise. Flash chromatography was carried out using 0.063–0.1 mm silica gel with the desired solvent. Thin-layer chromatography (TLC) was performed using 0.2 mm Kieselgel F254 (Merck) silica plates, and compounds were visualized using UV irradiation at 254 nm or 365 nm and/or staining with a solution of potassium permanganate and potassium carbonate in aqueous sodium hydroxide. Melting points were determined on a Kofler hot-stage apparatus. Infrared spectra were obtained using a PerkinElmer Spectrum 100 FTIR spectrometer as a

Table 2. Commercial Manuka Samples Purchased from Singapore

sample	lepterdine fluorescence (RFU)	compound concentration ^a (mg/kg)								classification ^b		
		lepterdine	4-HPA	3-PA	2-MB	2'-MAP	leptosperin	DHA	MGO	lepterdine (RFU)	lepterdine (LC/MS)	MPI
1	933	8.5	8.2	793	3.5	10.4	497	594	375	√	√	√
2	1046	8.8	4.5	910	3.0	7.8	377	711	542	√	√	√
3	1066	10.6	3.7	1164	3.1	6.2	394	1650	435	√	√	√
4	550	3.5	7.9	445	5.3	5.4	255	192	110	√	√	√
5	816	6.7	3.9	635	7.3	9.0	477	264	515	√	√	√
6	1212	14.6	4.1	1604	9.0	10.1	554	933	1110	√	√	√
7	1171	9.9	4.1	1195	7.1	16.7	411	419	915	√	√	√
8	1270	15.0	3.4	1012	6.2	13.5	615	660	688	√	√	√
9	814	6.8	5.6	509	4.4	6.1	383	262	224	√	√	√
10	1047	6.3	4.0	707	2.5	6.7	347	911	372	√	√	√
11	1390	13.6	2.9	1105	10.5	15.0	733	907	939	√	√	√
12	936	5.4	2.7	1090	18.9	10.0	234	133	682	√	√	√
13	849	5.7	5.5	989	7.9	5.3	318	712	585	√	√	√
14	1500	17.8	9.0	908	3.7	9.7	504	349	777	√	√	√
15	902	5.7	4.4	857	3.0	7.2	360	647	402	√	√	√
16	1199	7.3	6.6	932	2.3	9.5	458	1070	715	√	√	√
17	1334	18.2	6.9	969	3.2	8.6	607	855	888	√	√	√
18	754	8.6	3.3	629	2.7	4.3	278	976	313	√	√	×
19	574	3.5	2.6	338	2.1	2.2	168	160	121	√	√	×
20	539	3.4	3.8	195	3.5	2.7	152	231	103	√	√	×
21	921	3.9	6.3	379	4.5	14.6	123	36	70	√	√	×
22	355	3.0	0.9	730	3.5	3.2	157	85	128	×	√	×
23	481	3.6	2.7	1079	nd	6.4	173	295	140	×	√	×
24	469	1.5	2.2	584	5.0	6.0	168	221	241	×	×	√
25	511	1.2	3.8	402	2.8	4.6	231	165	94	×	×	×
26	458	2.4	5.7	717	1.6	2.9	156	94	117	×	×	×
27	491	1.8	3.6	411	5.1	3.4	276	354	149	×	×	×
28	438	2.5	1.9	296	3.1	3.2	157	157	124	×	×	×
29	422	2.2	2.7	257	2.9	3.4	138	50	69	×	×	×
30	359	1.1	n.d.	103	3.7	3.0	73	57	59	×	×	×
31	314	1.2	n.d.	142	1.0	2.6	75	52	59	×	×	×
32	375	<LOQ	10.4	1330	nd	3.4	63	<10	46	×	×	×
33	342	1.5	4.6	1645	nd	2.6	51	<10	33	×	×	×

^aLepterdine concentrations were quantified using the LC–MS/MS isotope dilution method; 4-HPA, 3-PA, 2-MB, 2'-MAP, and leptosperin concentrations were quantified by HPLC-UV; DHA and MGO concentrations were quantified and supplied by the UMF Honey Association (UMFHA). ^bClassification as monofloral manuka honey (√) or not monofloral manuka honey (×).

film attenuated total reflection (ATR) sampling accessory. Absorption maxima are expressed in wavenumbers (cm⁻¹).

NMR spectra were recorded as indicated on either a Bruker Avance 400 spectrometer operating at 400 MHz for ¹H nuclei and at 100 MHz for ¹³C nuclei, a Bruker DRX-400 spectrometer operating at 400 MHz for ¹H nuclei and at 100 MHz for ¹³C nuclei, a Bruker Avance AVIII-HD 500 spectrometer operating at 500 MHz for ¹H nuclei and at 125 MHz for ¹³C nuclei, or a Bruker Avance 600 spectrometer operating at 600 MHz for ¹H nuclei and at 150 MHz for ¹³C nuclei. ¹H and ¹³C chemical shifts are reported in parts per million (ppm) relative to CDCl₃ (¹H and ¹³C) or (CD₃)₂SO (¹H and ¹³C). ¹H NMR data is reported as chemical shift, relative integral, multiplicity (s, singlet; assignment). Assignments were made with the aid of correlation spectroscopy (COSY), nuclear Overhauser effect spectroscopy (NOESY), heteronuclear single quantum correlation (HSQC), and heteronuclear multiple bond correlation (HMBC) experiments where required. High-resolution mass spectra were recorded on a Bruker micrOTOF-Q II mass spectrometer with an ESI ionization source.

3.3.1. 6-Amino-3-(²H₃)methyluracil. 6-Aminouracil (1.05 g, 8.29 mmol) was suspended in hexamethyldisilazane (HMDS) (5 mL), and sulfuric acid (0.02 mL) was added. The mixture was heated at reflux for 1.5 h and then concentrated in vacuo. The residue was dissolved in dimethylformamide (6 mL), and iodomethane-*d*₃ (0.8 mL, 12.9 mmol) was added. Stirring was continued for 72 h at room temperature. The reaction was cooled to 0 °C, and sodium bicarbonate (15 mL) was carefully added. The mixture was stirred at 0 °C until no further bubbling was observed. The precipitate was filtered, washed with methanol and water, and dried to give a yellow solid (0.43 g, 36%) which was used without further purification, mp 340–345 °C (decomposition), ¹H NMR (400 MHz, DMSO-*d*₆): δ = 10.35 (1H, s, H-1), 6.15 (2H, s, NH₂), 4.56 (1H, s, H-5); ¹³C NMR (100 MHz, DMSO-*d*₆): δ = 163.2 (C-4), 153.5 (C-6), 151.1 (C-2), 74.0 (d, J = 168.1 Hz, C-4); IR (neat, cm⁻¹) = 3417, 3193, 1632, 1587, 1434, 1237, 788; HRMS (ESI): calculated for C₅H₄D₃N₃O₂Na⁺ [M + Na⁺] 167.0619; found: 167.0618.

3.3.2. 6-Amino-3-(²H₃)methyl-5-nitrosouracil. 6-Amino-3-(²H₃)methyluracil (0.40 g, 2.76 mmol) was suspended in water

(5 mL). The suspension was heated at reflux for 2.5 h and then cooled to room temperature. Acetic acid (1.68 g, 28.0 mmol) was added. A solution of sodium nitrite (0.46 g, 6.70 mmol) in water (4 mL) was then added dropwise over 5 min, during which time the pale-yellow suspension became gray. The mixture was stirred for 5 min before the precipitate was filtered, washed with methanol and water, and then dried to give a gray solid compound (0.39 g, 81%) which was used without further purification, mp > 350 °C. ^1H NMR (400 MHz, CDCl_3) δ = 11.37 (1H br, s, H-1), 7.98 (2H br, s, NH_2); ^{13}C NMR (100 MHz, $\text{DMSO}-d_6$) δ = 161.4 (C-4), 149.4 (C-2), 144.5 (C-6), 139.7 (C-5); IR (neat, cm^{-1}) = 3207, 3012, 1719, 1649, 1517, 1430, 1251, 1045, 768; HRMS (ESI): calculated for $\text{C}_5\text{H}_3\text{D}_3\text{N}_4\text{O}_3\text{Na}^+$ [$\text{M} + \text{Na}^+$] 196.0520; found: 196.0527.

3.3.3. 5,6-Diamino-3-($^2\text{H}_3$)methyluracil. 6-Amino-3-($^2\text{H}_3$)-methyl-5-nitrosouracil (0.12 g, 0.71 mmol) was suspended in a mixture of ammonium hydroxide (1.5 mL, 28–30%) and water (1.75 mL). The suspension was heated to 70 °C, and sodium dithionite (0.51 g, 2.93 mmol) was added portion-wise over 25 min until the red solution became pale-yellow. The mixture was stirred at 70 °C for 1 h before being concentrated in vacuo. The crude solid was then continuously extracted with refluxing ethanol (70 mL) for 3 h, and the extract was concentrated in vacuo to give a yellow solid (0.04 g, 37%), mp 222–225 °C (decomposition), ^1H NMR (400 MHz, CDCl_3) δ = 10.69 (1H br, s, H-1), 6.93 (2H br, s, NH_2), 3.33 (2H br, s, NH_2); ^{13}C NMR (100 MHz, $\text{DMSO}-d_6$) δ = 164.2 (C-4), 156.9 (C-6), 153.5 (C-2), 141.3 (C-5); IR (neat, cm^{-1}) = 3330, 2919, 1701, 1595, 1459, 1171, 962; HRMS (ESI): calculated for $\text{C}_5\text{H}_3\text{D}_3\text{N}_4\text{O}_2\text{Na}^+$ [$\text{M} + \text{Na}^+$] 160.0908; found: 160.0918.

3.3.4. 3,6,7-(3- $^2\text{H}_3$)Trimethylumazine. 5,6-Diamino-3-($^2\text{H}_3$)methyluracil (0.04 g, 0.26 mmol) was suspended in ethanol (2 mL). 2,3-Butandione (0.03 g, 0.33 mmol) and acetic acid (0.07 g, 1.22 mmol) were added. The mixture was heated at reflux for 24 h before being cooled to room temperature and concentrated in vacuo. The crude product was purified by flash chromatography (petroleum ether–EtOAc 1:4) to give a colorless solid (0.02 g, 36%). An analytical sample was recrystallized from a mixture of chloroform and ethanol (1:1), mp 274–277 °C. ^1H NMR (400 MHz, CDCl_3) δ = 9.53 (1H, s, H-1), 2.66 (3H, s, C7- CH_3), 2.65 (3H, s, C6- CH_3); ^{13}C NMR (100 MHz, CDCl_3) δ = 161.1 (C-4), 158.9 (C-6), 150.6 (C-7),* 150.4 (C-2),* 145.0 (C-8a), 123.7 (C-4a), 22.8 (C6- CH_3), 21.9 (C7- CH_3) (*assignments are interchangeable); IR (neat, cm^{-1}) = 2920, 1724, 1662, 1561, 1353, 1274, 940; HRMS (ESI): calculated for $\text{C}_9\text{H}_7\text{D}_3\text{N}_4\text{O}_2\text{Na}^+$ [$\text{M} + \text{Na}^+$] 232.0888; found: 232.0888.

3.4. LC–MS/MS Quantification of Leptericidine. HPLC-grade acetonitrile and formic acid were purchased from Merck. Water was purified using a Barnstead Nanopure Diamond laboratory water system. A 10 μL injection was made of each sample directly onto a 0.3 \times 100 mm Zorbax 300SB- C18 column (Agilent, Santa Clara, CA) at 12 $\mu\text{L}/\text{min}$ for 6 min. The HPLC gradient between buffer A (0.1% formic acid in water) and buffer B (0.1% formic acid in acetonitrile) was formed at 6 $\mu\text{L}/\text{min}$ as follows: 10% B for the first 3 min, increasing to 25% B by 18 min, increasing to 97% B by 21 min, held at 97% until 24 min, back to 10% B at 25.5 min, and held there until 30 min. The LC effluent was directed into the ion spray source of a QSTAR XL hybrid quadrupole time-of-flight mass spectrometer (Applied Biosystems, Foster City, CA)

scanning from 150 to 800 m/z . Two precursor ions at m/z 207 and 210 corresponding to endogenous leperidine and the heavy leperidine standard, respectively, were selected for further MS/MS analyses. The respective fragment ions were monitored between m/z 70 and 210 with a collision energy of 35 V. The mass spectrometer and HPLC system were under the control of the Analyst QS 2.0 software package (Applied Biosystems, Waltham, MA).

3.5. Isotopic LC–MS/MS Method Validation. Validation of the isotopically labeled leperidine-based LC–MS/MS method was performed based on the Eurachem international guidelines.²⁵ Several validation parameters were assessed including linearity, repeatability, detection sensitivity, interference, matrix effect, and recovery. Quantification was achieved by measuring the area under curve (AUC) ratio of endogenous leperidine to isotopically labeled leperidine (known concentration of 20 ng/mL in solution) for a common, selected fragment ion (m/z 148.05). Quantitation was based on the area under curve ratio of leperidine fragment ion peak area to isotopically labeled leperidine fragment ion peak area, where the amount of isotopic leperidine is known (20 ng/mL). The method validation procedure was carried out to cater for a working range of 0.5–50 mg/kg, covering the typical concentrations of endogenous leperidine present in New Zealand manuka honey.¹¹

Linearity was established by regression statistics using calibration standards prepared in clover honey that lack measurable levels of endogenous leperidine. Repeatability of the method was determined by three independent runs on different days using freshly prepared quality control (QC) samples analyzed in six replicates. Ten blank clover honey samples were selected for sensitivity determination. The limit of detection (LOD) and limit of quantitation (LOQ) were determined based on 3- and 10-time standard deviation (SD) of ten blank honey runs, respectively. Matrix effects and recovery were assessed by spiking a non-manuka honey with known levels of leperidine spanning the working range (3, 40, and 80 mg/kg) in triplicate and comparing to the corresponding responses in the honey-free solution (0.1% formic acid).

3.6. HPLC–UV Quantification. HPLC quantification was performed using the method described previously.¹¹ A Thermo Scientific Dionex Ultimate 3000 Standard system (Thermo Fisher Scientific Inc., Bremen, Germany) was employed, coupled to a quaternary pump (LPG-3400SD), an Ultimate analytical autosampler (WPS-3000TSL), and a diode array detector (DAD-3000). Data acquisition, peak integration, and calibrations were performed with Dionex Chromeleon 7.2 chromatography data system software. Separations were carried out on a Hypersil GOLD column (150 \times 2.1 mm; 3 μm) (Dionex, NZ) connected to a Guard column (10 \times 2.1 mm; 3 μm Hypersil), using a binary mobile phase consisted of 0.1% aqueous formic acid (solvent A) and acidified methanol containing 0.1% formic acid (solvent B). The temperature of the column compartment (TCC-3000SD) was set at 32 °C.

All honey samples were diluted to a final concentration of 0.1 g/mL, followed by centrifugation at 14 500 rpm for 5 min. A total volume of 100 μL was then loaded onto a 96-well microplate (Greiner Bio-One, polystyrene, conical bottom) for each diluted honey sample. The injection volume was 5 μL for all samples. Elution was performed with the following gradient: 0–2 min, 2% B; 2–5 min, 5% B; 5–15 min, 25% B; 15–25 min, 50% B; and 25–31 min, 100% B, followed by washing

(held 3 min, 100% B) and reconditioning (held 10 min, 2% B). The flow rate and temperature were set to 0.16 mL/min at 32 °C. Samples were monitored according to the UV absorbance at four different wavelengths of 250, 265, 280, and 330 nm. Compound concentrations were quantified against external calibration curves of respective chemical standards based on the peak area under curve (AUC).

3.7. Fluorescence Spectroscopy. Fluorescence measurements were performed using a Gemini EM dual-scanning microplate spectrofluorometer (Molecular Devices Inc. San Jose, CA) coupled with SoftMax Pro software. Honey dilutions were freshly prepared at 2% w/v, and a 100 μ L aliquot was loaded into each microplate well (OptiplateTM-384, black). The spectrofluorometer was set to top-read with activated autocalibration and automatic photomultiplier tube (PMT) sensitivity.

3.8. Data Analysis. Extracted ion chromatograms for the fragment ion of m/z 148.05 \pm 0.1 Da from leperidine and the heavy isotope-labeled standard were created and integrated using MultiQuant v3.0 (Sciex). Statistical data analyses were performed using Graphpad Prism software (version 6.01). All correlations were determined by regression analysis. Differences between group means were determined by one-tailed Student's *t*-test. ROC analyses were performed using MetaboAnalyst 3.0 (<http://www.metaboanalyst.ca>).

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsomega.0c00486>.

Data related to validation of the leperidine isotopic LC–MS/MS method; three independently prepared calibration curves for leperidine (1–100 ng/mL) based on weighted linear regression of peak area ratios (leperidine/isotopically labeled leperidine) for the selected common fragment ion m/z 148.05 (Figure S1); representative extracted ion chromatograms for the common fragment ion m/z 148.05 from (A) unspiked clover honey and (B) unspiked manuka honey showing the leperidine and heavy isotope-labeled leperidine channels (Figure S2); accuracy of isotopically labeled leperidine-based mass spectrometry measurement as determined by the percentage of theoretical concentration and correlation coefficient (Table S1); precision (intraday repeatability) of leperidine quantification expressed as % CV (Table S2); reproducibility (interday precision) of LC–MS/MS leperidine quantitation determined as % CV (Table S3); and recovery of leperidine calculated based on difference between amount recovered from spiked honey and from honey-free solution (Table S4) (PDF)

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Notes

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