- 1 Analysis of the flavonoid component of bioactive New Zealand mānuka
- 2 (Leptospermum scoparium) honey and the isolation, characterisation and
- 3 synthesis of an unusual pyrrole.
- 4 Running title: Flavonoids and an unusual pyrrole in New Zealand mānuka
- 5 honey.
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- 11 ABSTRACT:
- 12 The flavonoid components of New Zealand mānuka (*Leptospermum scoparium*) honey have been
- 13 quantified in a series of 31 honeys of varying non-peroxide antibacterial activity to clarify
- 14 discrepancies between previous studies reported in the literature. Total flavonoid content was
- 15 1.16 mg/100 g honey. The principal flavonoids present were pinobanksin, pinocembrin, luteolin
- and chrysin and together these represented 61% of the total flavonoid content. 1, 2-formyl-5-(2-
- methoxyphenyl)-pyrrole, which was weakly correlated with the non-peroxide antibacterial activity,
- 18 was isolated from the flavonoid fraction and separately synthesised. 1 did not display inhibitory
- activity against *S. aureus* in vitro and thus the origin of the correlation, which is still unknown, is not
- a direct contribution.

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INTRODUCTION:

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New Zealand mānuka (Leptospermum scoparium) honey is noted for its non-peroxide antibacterial activity (Allen, Molan & Reid, 1991). This activity has been attributed principally to the presence of methylglyoxal (Adams et al., 2008, Mavric, Wittman, Barth & Henle, 2008), which originates from dihydroxyacetone in the nectar of the flower (Adams et al., 2009), the reasons why some individuals of this species produce high levels of dihydroxyacetone are unknown. Because of its non-peroxide antibacterial activity, New Zealand manuka honey commands a premium price worldwide and is thus subject to fraud and substitution; to establish authenticity, markers of floral origin are urgently needed. Flavonoid profiles have been used to determine the floral origin of honeys (Tomas-Barberan, Ferreres, Garcia-Viguera & Tomas-Lorente, 1992, Tomas-Barberan, Ferreres, Garcia-Viguera & Tomas-Lorente, 1993, Anklam, 1998); flavonoids thus represent a possible biomarker for mānuka honey. Flavonoids are known for their antibacterial and other bioactivities (Cushnie & Lamb, 2005, Havsteen, 2002, D'Arcy, 2005) and there exists a possibility that one or more components of the flavonoid fraction of mānuka honey contribute directly or indirectly (by involvement in the biosynthetic pathway generating high levels of dihydroxyacetone) to the nonperoxide antibacterial activity of the honey. Weston, Brocklebank and Lu (2000) found the principal flavonoids in mānuka honey to be pinobanksin with 4.3, 6.3, 5.0 μg/100g honey respectively for high (10 samples), low (3 samples) and zero (6 samples) antibacterial activity respectively, pinocembrin with 5.1, 7.3, 4.5.0 μg/100g honey for high, low and zero antibacterial activity respectively, chrysin with 2.2, 2.7, 2.2 µg/100g honey for high, low and zero antibacterial activity respectively and galangin with 2.2, 3.4, 1.9 µg/100g honey for high, low and zero antibacterial activity respectively. This is in contrast to the work of Yao, Datta, Tomas-Barberan, Ferreres, Martos and Singanusong (2003), who found the principal flavonoids in two samples of mānuka honey to be quercetin (430 μg/100g honey), isorhamnetin (400 μg/100g honey), an unknown flavonoid (390 μg/100g honey), chrysin (380 μg/100g honey) and luteolin (380 μg/100g honey) with a total flavonoid content of 3060 µg/100g honey. There is a very marked qualitative

and quantitative difference between these two sets of results and indeed Yao et al. (2003) recommended further analysis of New Zealand mānuka honeys using a large sample set and the HPLC detection methodology that they espoused, that is detection of flavonones at 290 nm and of flavones at 340 nm using diode array detection. Pyrroles in honey can arise from Maillard reactions or from the pyrolysis of amino acids when honey is heated (Jerković, Mastelić, Marijanović, Klein & Jelić, 2007). 1H-Pyrrole has been found in Robinia pseudoacacia L., Castanea sativa L. and Salvia officinalis L (Jerković et al., 2007, Jerković, Mastelić & Marijanović, 2006). 1H-Pyrrole-2-carboxylic acid has been found in Paliurus spinachristi (Jerković, Tuberoso, Marijanović, Jelić & Kasum, 2009). 2-Acetylpyrrole has been found in abbamele, a honey-based Sardinian product (Jerković, Kasum, Marijanović & Tuberoso, 2011) and 1H-pyrrole-3,4-diacetic acid has been found in pine honey (Pinus brutia Ten) (Eraslan, Kanbur, Silici & Karabacak, 2010). Phenyl substituted pyrroles are also found in natural products, pyrrolnitrin is a tryptophan-derived, antifungal antibiotic isolated from Pseudomonas pyrrocinia (van Pée & Ligon, 2000). Both pentachloropseudilin and pentabromopseudilin, which are produced by an Actinoplanes sp. strain, are strongly active against Gram-positive bacteria. The latter is also known to inhibit a number of different enzyme systems and has high in vitro activity against leukemia and melanoma cell lines (van Pée & Ligon, 2000). This study aimed to investigate the flavonoid profiles of a large sample set of New Zealand mānuka honeys using the detection methods suggested by Yao et al. (2003) and to establish if there was any link between this profile and the afore-mentioned non-peroxide antibacterial activity, which is a feature of these honeys. During the course of this investigation an unidentified peak in the HPLC chromatogram of the flavonoid fraction was shown to be weakly correlated to antibacterial activity. The compound, 2-formyl-5-(2-methoxyphenyl)-pyrrole, 1, was subsequently

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isolated from the flavonoid fraction and fully characterised.

MATERIALS AND METHODS

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Materials: Thirty one manuka honey samples with non-peroxide antibacterial activities ranging from very low to 27.3 UMF™, were kindly supplied by Comvita New Zealand Ltd., Te Puke, New Zealand. A bulk sample of active mānuka honey (6 kg, Haddrell's, Cambridge, New Zealand) was used for the optimisation and testing of the extraction method. Mānuka honey (15 kg, Natures Country Gold, Hamilton, New Zealand) was used for the isolation of flavonoids. 3 was used as a model compound and 4 as an analogue for the characterisation of 1. 3, 5-(2nitrophenyl)-2-furaldehyde, was obtained from Acros Organics. 4, 2-Formyl-5-phenyl-pyrrole, was generously donated by Dr J. T. Reeves of the Department of Chemical Development, Boehringer Ingelheim Pharmaceuticals Inc., Ridgefield, CT 06877-0368, USA. Methanol was redistilled from drum grade or HPLC grade (Scharlau), Milli-Q water was obtained from a Barnstead E-pure system (18.2 MΩcm). Dried and purified diethyl ether, hexanes, tetrahydrofuran and dichloromethane were obtained from a Pure Solvent Purification System (Model: PS-SD-5) as needed and used promptly. Dimethyl sulfoxide- d_6 (99.9 atom% D containing 0.03% v/v TMS) and chloroform- d_1 (99.8 atom% D) were obtained from Sigma-Aldrich Inc. Lithium diisopropylamide (LDA), o-methoxyacetophenone, oxazole-4-carboxaldehyde, diisobutylaluminium hydride (DIBALH) and triethylamine (Et₃N) were obtained from Sigma-Aldrich Inc., methane sulphonyl chloride (MsCl) was obtained from Riedel-de Haën. Caffeic acid (≥98%), (+)-catechin hydrate (98%), chlorogenic acid (predominantly trans) (\geq 95%), chrysin, trans-cinnamic acid (99+%), p-coumaric acid (\geq 98%), (-)-epicatechin (97%), galangin (no stated purity), gallic acid (97%), luteolin (≥ 98%), myricetin (approx 85%), pinocembrin (95%), quercetin dihydrate (98%), (+)-rutin hydrate (95%) and syringic acid (98%) were obtained from Sigma-Aldrich Inc., isorhamnetin (pure), kaempferol (pure) and naringin (pure) were obtained from Indofine Chemical Company Inc.

XAD-2 Amberlite resin (pore size 9 nm, particle size 0.3-1.2 mm) was obtained from Sigma-Aldrich Inc. Sephadex LH-20 resin was obtained from Amersham Pharmacia Biotech AB and silica gel was obtained from Merck.

General Methods: Evaporation of aqueous samples under reduced pressure was accomplished using a rotary evaporator with a water bath set to 35 °C, smaller samples were reduced in volume by evaporation under a stream of dry nitrogen (35 °C). UV-Visible spectra of isolated flavonoids were recorded on a Varian Cary 100 Scan UV-Visible Spectrophotometer, sodium methoxide, sodium acetate, sodium acetate/boric acid, aluminium chloride and aluminium chloride/hydrochloric acid were variously added to measure changes in the UV spectrum (Markham & Mabry, 1975, Jurd, 1962, Mabry, Markham & Thomas, 1970) for purposes of identification.

Isolation of Flavonoid Fraction: Phenolics were extracted from samples of honey using XAD-2 resin according to the method of Ferreres, Tomas-Barberan, Gil & Tomas-Lorente (1991). Phenolic acids and flavonoids were separated using Sephadex-LH20 according to the method of Bohm (1998). Semi-preparative High Performance Liquid Chromatography (HPLC): Semi-preparative HPLC of bulk fractions of honey for isolation of flavonoids was carried out using two Waters 515 pumps, an Alltech Elite Degassing System, a Rheodyne 7725i injector fitted with a 500 μ L sample loop, and a Waters 996 photodiode array detector. A reversed phase octadecylsilane Waters Xterra Prep column (PR₁₈, 10 μ m, 7.8 mm x 300 mm) was used in this system. A binary mobile phase was used; this consisted of solvent A which was Milli-Q water acidified with acetic acid (0.075% v/v) and methanol (5% v/v) and solvent B which was methanol. The overall flow rate was 4 mL/min. The gradient used was held at 60% A for 5 min, decreasing to 50% A at 10 min and 20% A at 60 min and finally 0% A at 60.2 min.

Analysis of flavonoids and isolation of 1 by High Performance Liquid Chromatography (HPLC): Analysis of flavonoid extracts and isolation of 1 was carried out using two Waters 515 HPLC pumps with their flows combined through a static mixer (Grace Binary Large Volume Mixer SS Housing with 350 μL Mixer Cartridge). Both solvents were degassed prior to entering the pumps by passing through a Waters In-Line Degasser AF. Samples and standards were warmed (40 °C) and injected manually using a Rheodyne 7725i injector system fitted with a Rheodyne loop (5 μL for analyses, 50 µL for isolations). Separation was achieved using a Waters Symmetry Shield™ octadecylsilane HPLC column (RP18, 5 μm, 3.0 x 250 mm) with a Waters Universal Sentry™ Guard SymmetryShield™ Column. A binary gradient was used, solvent A was Milli-Q water and methanol (5% v/v), acidified with acetic acid (0.075% v/v), solvent B was methanol. The gradient was run at a constant combined flow rate of 0.3 mL/min. Detection was achieved using a Waters 996 Photodiode Array Detector (240-400 nm). Gradient method 1 (for analyses and initial isolation of 1) consisted of 70% A for 15 min, decreasing to 40% A at 20 min and 0% A at 60 min and held for 10 min. Gradient method 2 (for final purification of 1) consisted of 70% A for 5 min, decreasing to 55% A at 10 min, 20% A at 50 min and 0% A at 52 min and held for 8 min. Gas Chromatography-Mass-Spectrometry (GC-MS): GC-MS was carried out on a HP 6890 series GC fitted with a Phenomenex ZB-5 5% phenyl-methylsiloxane) column (30 m x 0.25 mm x 0.025 μm) interfaced to a HP 5973 mass selective detector. Conditions used were 120 °C (0.75 min), 50 °C/min up to 200°C, and 10°C/min up to 295°C (held 15 min). The mass spectrometer was operated in either total ion chromatogram (TIC) or a selected ion monitoring (SIM) mode using m/z201 (M^{+}), 158 and 130 ions in the case of **1**. High Resolution Mass Spectrometry (HRMS): Mass spectra were recorded in positive-ion mode on a Bruker MicrOTOF mass spectrometer with electrospray interface and MeOH as mobile phase. Assignments of major peaks were confirmed by comparison of the high-resolution isotope pattern of the ions with the theoretical pattern obtained using isotope ratios.

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Nuclear Magnetic Resonance Spectroscopy (NMR): 1D- and 2D- ¹H and ¹³C NMR spectra of samples were obtained using a Bruker Avance DRX-400 spectrometer. ¹H and ¹³C experiments were carried out at 400.13 and 100.62 MHz respectively using a 5 mm inverse ¹³C/¹H probe head except for the ¹³C spectrum of **1**, which was acquired using a 5 mm dual ¹³C/¹H probe head. Samples were dissolved in dimethylsulfoxide- d_6 or chloroform- d_1 for NMR analysis. Operation of the NMR spectrometer and processing of spectra were performed using Bruker Topspin 1.3 software. Measurement of UMF™: Measurement of non-peroxide antibacterial activity was carried out using the method previously described by Allen et al. (1991) and isolated fractions were mixed with clover honey as an artificial matrix for testing (Adams et al., 2008). Measurement of Bioactivity of 1: Bioactivity testing was carried out at the ithree Institute, School of Medical and Molecular Sciences, University of Technology, Sydney, Australia. Staphylococcus aureus strain NCTC 8325 was used throughout this part of the study. Growth assays were set up in cation-adjusted Mueller Hinton II Broth (CaMHB, Becton Dickinson) whereas biofilm assays were performed in tryptic soya broth (TSB, Oxoid). For a growth study, 1 was diluted in dimethyl sulfoxide (DMSO) and then further serial dilutions were made in DMSO. Each dilution was added in duplicate to a 96-well plate to a final DMSO concentration of 2%. An overnight culture was diluted to 1 x 108 cfu/mL (determined by backtitration on tryptic soy agar plates) and 150 µL was added to each well of the 96-well plates. Controls included a serial dilution of Lincomycin (to assess plate-to-plate variation), a positive control with bacteria alone in CaMHB with 2% DMSO and a negative (no bacteria) with 150 µl CaMHB containing 2% DMSO. Plates were incubated in a shaking incubator at 37 °C for 22 h and absorbance was measured at a wavelength of 595 nm using a plate reader (Biotek Synergy HT). For a biofilm assay, plates were set up as for the growth assay except that bacteria were grown in TSB. Plates were then sealed with AeraSeal (Excel Scientific) and incubated at 37 °C in a humidified

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incubator for 24 h. Plates were washed with PBS and dried at 60 $^{\circ}$ C for 1 h. Biofilm was stained by the addition of 200 μ L of 0.2% crystal violet at room temperature for 1 h. Plates were washed 3 times with water and air-dried and the amount of biofilm biomass was quantified by destaining the biofilm with 250 μ L of 30% acetic acid. Plates were incubated on a shaking platform at room temperature for 15 min and absorbance was then read at a wavelength of 635 nm.

For a disc diffusion test, an overnight culture (100 μ L) was streaked out with a sterile cotton swab on a tryptic agar plate. Discs containing 10 μ g of **1**, rifampicin (positive control) or DMSO (negative control) were placed on the lawn of bacteria. Plates were then incubated overnight at 37 °C and assessed for zones of bacterial inhibition.

3-hydroxy-1-(2-methoxyphenyl)-3-(oxazol-4-yl) propan-1-one (5): To THF (20 mL) at -84 °C was added LDA (2.5 mL, 5.5 mmol). The resultant solution of LDA was added dropwise to a solution of *o*-methoxyacetophenone (0.6 mL, 4.5 mmol). The reaction mixture was stirred at -84 °C for 30 minutes before a solution of oxazole-4-carboxaldehyde (0.5 g, 5.5 mmol) in a minimal amount of THF was added dropwise at -84 °C. The reaction mixture was stirred for 30 minutes at -84 °C, quenched with saturated aqueous NH₄Cl (20 mL) and allowed to warm to room temperature. The layers were allowed to separate, and the aqueous layer was extracted with EtOAc (2 x 20 mL). The combined organic layers were dried (MgSO₄), filtered and concentrated. The product was recrystallized from CH₂Cl₂/hexanes yielding **5** as pale yellow crystals (0.751 g, 68%), Mpt: 104 – 107 °C, HRMS: found: 270.0735, calculated for C₁₂H₁₂NO₄Na [M+Na]*: 270.0737, ¹H NMR (400.13 MHz, DMSO-*d*₆) δ ppm 3.87 (s, 2-OCH₃), 5.36 (d, J = 5.4 Hz, OH), 3.26 (dd J = 8.2, 16.2 Hz, H₃-2′), 3.37 (dd J = 4.6, 16.2 Hz, H_b-2¹), 5.06 (m, H-3¹), 7.89 (s, H-5¹), 8.26 (s, H-6¹), 7.16 (d, J = 8.6 Hz, H-3), 7.54 (t, J = 7.5 Hz, H-4), 7.03 (t, J = 7.5 Hz, H-5), 7.52 (d, J = 8.6 Hz, H-6), ¹³C NMR (100.62 MHz, DMSO-*d*₆ δ ppm 199.6 (C-1¹), 50.4 (C-2¹), 62.6 (C-3¹), 143.3 (C-4¹), 135.1 (C-5¹), 151.5 (C-6¹), 128.1 (C-1), 158.0 (C-2), 112.3 (C-3), 133.6 (C-4), 120.9 (C-5), 129.4 (C-6).

2-formyl-5-(2-methoxyphenyl)-pyrrole (1): A solution of 5 (0.23 g, 0.9 mmol) in THF (4 mL) was treated at 0 °C with Et₃N (0.4 mL) followed by dropwise addition of MsCl (0.1 mL), the reaction mixture was stirred at 0 °C for 1 h. Aqueous NaOH (2 M, 6 mL) was added and the reaction mixture heated (70 °C, 72 h). After cooling to room temperature, the reaction mixture was diluted with saturated aqueous NaHCO₃ solution, extracted with EtOAc (2 x 20 mL) and the organic layer dried (MgSO₄), filtered and concentrated. The product was purified by chromatography on a silica gel column (EtOAc – hexane, 1:9 \rightarrow 1:0), the fraction containing 1 was concentrated and further purified with preparative layer chromatography using a circular plate (220 mm diameter, 2 mm silica layer: Merck PF245) installed on a Chromatotron (Harrison Research) using 50 mL portions of hexane-diethyl ether mixtures (4:1, 2:3, 4:1) as eluent. This yielded 1 as a yellow solid (0.36 mg, 0.2%), HRMS: found: 224.0653, calculated for $C_{12}H_{11}NO_2Na$ [M+Na]⁺: 224.0682, ¹H NMR (400.13) MHz) and ¹³C NMR (100.62 MHz) see Table 4. The major product formed during the synthesis of $\mathbf{1}$ was (E)-1-(2-methoxyphenyl)-3-(oxazol-4-yl)prop-2-en-1-one, **6**, and its (*Z*)-isomer (minor) eluting in GC-MS at 6.28 min and 5.60 min respectively. Chromatography under the conditions used for 1 gave 6 as orange crystals (0.075g, 36%), Mpt: 75 - 80 °C, HRMS: found: 252.0644, calculated for $C_{13}H_{11}NO_3Na$ [M+Na][†]: 252.0631, ¹H NMR (400.13 MHz, DMSO- d_6) δ ppm 3.86 (s, 2-OCH₃), 7.42 (dd, J = 15.5, 0.8 Hz, H-2'), 7.35 (d, J = 15.5) 15.5 Hz, H-3'), 8.48 (t, J = 0.8 Hz, H-5'), 8.50 (s, H-6'), 7.19 (dd, J = 8.3, 1.0 Hz, H-3), 7.55 (td, J = 8.3, 1.8 Hz, H-4), 7.06 (td, J = 7.5, 1.0 Hz, H-5), 7.48 (dd, J = 7.5, 1.8 Hz, H-6), 13 C NMR (100.62 MHz), DMSO- $d_6\delta$ ppm 191.9 (C-1'), 131.1 (C-2'), 127.0 (C-3'), 136.3 (C-4'), 141.7 (C-5'), 153.1 (C-6'), 128.4 (C-1), 157.7 (C-2), 112.3 (C-3), 133.1 (C-4), 120.6 (C-5), 129.5 (C-6), 55.8 (2-OCH₃). The ¹H NMR spectrum of the crude reaction mixture also showed signals at 6.85 ppm (J = 12.6 Hz) and 6.99 ppm (J = 12.6 Hz) consistent with the presence of the olefinic protons of the corresponding (Z)-isomer.

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RESULTS AND DISCUSSION

Flavonoid profiles: The flavonoid profiles of 31 different mānuka honey samples were analysed.
XAD-2 extraction was used according to the method of Ferreres <i>et al.</i> (1991) but modified to use 5
g rather than 50 g samples, which modification was made to allow flavonoid profiling of a much
larger set of honey samples. To validate this modification multiple extractions of 50 g (3
replicates), 20 g (3 replicates) and 5 g (5 replicates) samples were performed and the total area of
the chromatogram (340 nm) in the phenolic region was measured. The results, (1.7 \pm 0.8) x 10^8 ,
(1.9 ± 1.0) x 10^8 and (2.1 ± 1.0) x 10^8 for 50, 20 and 5 g respectively, indicate that any variation due
to sample size is insignificant compared with the intra-sample size variation. Phenolic acids and
flavonoids were separated using Sephadex-LH20 (Bohm, 1998). The HPLC chromatogram of the
flavonoid fraction of a typical mānuka honey is shown in Fig.1. Flavonoids and phenolic acids were
identified by HPLC co-elution with an authentic standard and for some confirmed by isolation and
characterisation by NMR and UV spectroscopy (caffeic acid, p-coumaric acid, pinobanksin and 8-
methoxykaempferol). Several compounds were identified as flavonoids by their characteristic UV
spectra, Table 1, but isolation and identification has so far proven elusive. Unknown flavonoids 1,
2 and 5 showed only a Band II absorption for λ_{max} indicating that they are flavanones and/or
dihydroflavonols while unknown flavonoids 3, 4 and 6 showed both Band I and Band II absorptions
indicating that they are flavones and/or flavonols. Quantification of the flavonoids was achieved by
comparing their absorbance in the HPLC chromatograms to four external standards. Pinocembrin
at 290 nm was used to quantify all flavanones and dihydroflavonols, chrysin at 340 nm was used to
quantify flavones and flavonols with unsubstituted B rings and kaempferol at 340 nm was used for
those flavones and flavonols with a singly oxygenated B ring, while quercetin at 340 nm was used
for all other flavones and flavonols, Table 2. Total flavonoid content was 1.16 mg/100 g honey with
a 95% confidence interval of 0.16 mg/100 g honey and a range of 0.594-2.235 mg/100 g honey.
This value is considerably higher than the 0.0147 mg/100 g total flavonoid content reported by
Weston et al. (2000) and closer to that reported by Yao et al. (2003), that is 3.06 mg/100 g of honey

for two samples. Venugopal and Devarajan (2011) found 3.34 mg (chrysin equivalents)/100 g of honey in a single sample of mānuka honey using a chemical method.

The principal flavonoids present in the HPLC chromatograms were pinobanksin (PB, 0.27 ± 0.04 mg/100 g honey), pinocembrin (PC, 0.17 ± 0.02 mg/100 g honey), luteolin (L, 0.14 ± 0.02 mg/100 g honey) and chrysin (C, 0.13 ± 0.02 mg/100 g honey) and together these represented 61% of the total flavonoid content. Hitherto Oelschlagel, Gruner, Wang, Boettcher, Koelling-Speer and Speer (2012) have qualitatively reported the presence of luteolin in mānuka honey extracts analysed by UPLC while Daher and Gülaçar (2010) have observed the presence of pinocembrin in two mānuka honey samples examined by SPME followed by GC-MS.

In this study in addition to the four principal flavonoids five other known flavonoids quercetin (Q),

8-methoxykaempferol (8MK), isorhamnetin (IR), kaempferol (K) and galangin (G) and six unknown flavonoids 1-6 (F1-6) were detected at lower levels. In addition to the well known propolis-derived flavonoids pinocembrin, pinobanksin, chrysin and galangin, as found for example in the studies of 19 mānuka honeys by Weston, Mitchell and Allen(1999) and Weston *et al.* (2000), we detected the presence of a further five known flavonoids by comparison with standards. The identification of chrysin and luteolin as major components of the flavonoid fraction of mānuka honey is consistent with the results of Yao *et al.* (2003) which showed quercetin (13.8%), isorhamnetin (12.9%), an unknown flavanone (12.7%), chrysin (12.6%) and luteolin (12.6%) as the main components of two mānuka honey samples; isorhamnetin and quercetin were however only found at low levels in the present study. The quercetin content of the 31 mānuka honeys in the present study was found to be highly variable with levels ranging from 0.000 to 0.115 mg/100 g honey. This high variability suggests that any quercetin found in mānuka honey probably originates from floral sources (introduced by nectar or pollen) other than mānuka trees. Pinobanksin, which was the main component of the mānuka honey flavonoids in the present study, was not observed in the two mānuka honey samples analysed by Yao *et al.* (2003). In the current study a significant level (an

average of 14% of the total flavonoid content assuming a detector response similar to quercetin) of a non-flavonoid peak which eluted at 34.4 minutes, $\mathbf{1}$ (λ_{max} 342 nm), was identified in all of the flavonoid extracts.

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Antibacterial activity of phenolic acid and flavonoid fractions: Fractions obtained from the XAD-2 extraction were tested for antibacterial activity to determine the contribution of the phenolic acids and flavonoids to the non-peroxide, antibacterial activity of mānuka honey. The aqueous fraction of the XAD2 extraction contained the sugars and other polar compounds including methylglyoxal, and the methanol fraction contained the phenolics including the flavonoids and phenolic acids. As a comparison, the activity of the original manuka honey and a sample of inactive clover honey (used as an artificial matrix for the methanol fractions) were tested. Extraction and testing was carried out in duplicate. The results of the activity testing are shown in Table 3. The phenolic extract, at concentrations equivalent to natural honey, made such a small contribution to the nonperoxide antibacterial activity it was not measurable by this assay, which concurred with the findings of Weston et al. (2000) and the attribution of this activity to water-soluble methylglyoxal (Adams et al., 2008, Mavric et al., 2008). At 25 times their natural concentration the phenolics did display antibacterial activity, antibacterial activity has previously been attributed to such compounds as well as other types of bioactivity (Cushnie & Lamb, 2005, Havsteen, 2002, D'Arcy, 2005). Although the flavonoid content of honey does not make a significant contribution directly to the non-peroxide antibacterial activity of honey, it is possible that there is some indirect contribution and to ascertain this, scatterplots were created of UMF™ versus peak area. The only two substances from the flavonoid profile that showed activity correlations were the non-flavonoid compound **1** ($R^2 = 0.36$) and luteolin ($R^2 = 0.23$), Fig. 2.

Isolation and characterisation of $\mathbf{1}$: The observation that $\mathbf{1}$ showed a moderate $R^2 = 0.36$ correlation with UMF levels raised the possibility that it might be a minor contributor to the bioactivity of manuka honeys and also serve as a marker compound for UMF activity, it was

therefore of interest to isolate this substance and to determine its structure and ascertain if it exhibited any significant bioactivity. 1 was isolated from the flavonoid fraction by HPLC with gradient method 1 and purified by HPLC with gradient method 2, Fig. 3. The ¹H NMR spectrum of 1 in DMSO- d_6 gave a 3 proton singlet at 3.96 ppm typical of a -OCH₃ or -COOCH₃ group, 6 proton signals in the region 6-8 ppm typical of aromatic or conjugated olefinic protons and a singlet signal at 9.52 ppm suggestive of a -CHO group. A broad peak at 11.79 ppm, which disappeared upon presaturation of the HOD peak, was attributed to an exchangeable proton. H,H-COSY revealed the presence of two separate spin systems, a 2 proton spin system comprised of mutually coupled signals (J = 3.9 Hz) centred at 6.77 and 7.04 ppm. The coupling constant exhibited by these protons, while not typical of a cis-coupled aryl proton or olefinic proton in a 7- or 8-membered ring, was typical of a cis-coupled proton in a 5-membered ring such as a furan, pyrrole or thiophene ring (27). The second 4 proton system was comprised of signals centered at 7.02, 7.14, 7.33 and 7.77 ppm and showed couplings and correlations typical of an ortho-substituted benzene. The small quantity of isolated sample meant that ¹³C NMR showed only protonated signals even after prolonged acquisition, namely OCH₃ (55.6 ppm), five or six aromatic or olefinic CH groups (114-129.5 ppm) possibly including two unresolved signals at circa 120.7 ppm and a conjugated aldehyde group (178.8 ppm). This was verified in an HSQC spectrum. An HMBC spectrum determined with a 65 msec mixing time revealed the presence of 4 quaternary carbons at 156.2, 136.5, 132.5 and 119.2 ppm. A 1D selective NOESY experiment in which the aldehyde proton at 9.52 ppm was irradiated enhanced the signal at 7.04 ppm indicating that the aldehyde group was spatially close to the cis-coupled proton spin system and distant from the aromatic protons. A similar experiment irradiating the methoxyl protons caused enhancement of the aromatic proton at 7.77 ppm indicating that the methoxyl group was a substituent on the aromatic ring. Thus it was concluded that 1 consisted of an ortho-substituted aromatic ring to which a methoxyl group was attached and a five-membered aromatic ring containing a heteroatom and to which an aldehyde group was attached. Hosoya et al. (2003) have reported the ¹H NMR spectrum of 5-(2-

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319 methoxyphenyl)-2-furaldehyde (2) in chloroform- d_1 . The reported spectrum of 2 shows similarities 320 to that of 1, however the exchangeable proton at 11.79 ppm was not observed in 2. 321 GCMS of 1 showed a base peak at m/z 201 and a much smaller peak at m/z 202. Although aldehydes 322 typically show a large [M-H]⁺, ion in this case it seemed likely that m/z 201 was the molecular ion and 323 202 was the corresponding isotope peak. The molecular mass of 2 is 202 and the substitution of N for O 324 in the five-membered ring would yield a molecular mass of 201 and be consistent with the presence of 325 an exchangeable proton in 1. 326 3 was utilised as a model compound to assist in identification of 1. The HMBC spectrum of 1 327 determined with a correlation (mixing) time of 65 msec did not include correlations which defined the 328 point at which the proposed pyrrole ring was linked to the methoxyl substituted ring. A series of HMBC 329 experiments were run using a specimen of 3 in which the mixing time was varied from 35 to 200 msec 330 and the intensity of the correlation between H-6' and C-5 was determined, this showed a maximum at 331 50 msec. Using this optimised mixing time the HMBC spectrum of 1 revealed the presence of a ³J 332 correlation between the aromatic proton at 7.77 ppm (H-6') and a quaternary carbon (C-5) at 136.5 333 ppm in the five-membered ring moiety. Other structurally significant HMBC correlations observed for 1 334 are depicted in Fig. 4. 335 GC-MS analysis of 3, under identical conditions used for 1, afforded a peak which had a retention time 336 of 11.98 min and showed a weak M⁺ ion at m/z 217 together with a base peak fragment ion at m/z 188 337 attributable to a $(M - CHO)^+$ ion. The observation that the retention time of **3** was greater than that 338 determined for 1 is consistent with the conclusion that 1 was a less polar, lower molecular weight 339 variant of 3. It was concluded that 1 was 2-formyl-5-(2-methoxyphenyl)-pyrrole. Reeves, Song, Tan,

substituent. A sample of 4 was generously donated by Dr J.T. Reeves.

Lee, Yee & Senanayake (2007) have reported the synthesis of another analogue of 1, 2-formyl-5-phenyl-

pyrrole, 4. The structure of 4 only differs from that proposed for 1 in that it lacks the aryl ring methoxy

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The ¹H and ¹³C NMR assignments for 1 are given in Table 4 and compared with those which were determined for the specimen of 4 supplied by Dr Reeves. Notable points of similarity in the ¹H spectrum include the broad signal for the exchangeable proton attached to N, the chemical shift of the aldehyde proton and the coupling constant (3.9 Hz) of the cis-protons in the pyrrole ring. When 1 was subsequently synthesised (see below) the 1 H NMR spectrum was also recorded in chloroform- d_{1} and the two pyrrole protons (H-3' and H-4') which were doublets in DMSO-d₆ were found to be a doublet of doublets in chloroform- d_1 . Homonuclear decoupling irradiating the NH proton signal reduced H-3' and H-4' to doublets with J = 3.9 Hz. GC-MS analysis of 4, under identical conditions to those used for 1 and **3** afforded a peak which had a retention time of 8.23 min and showed a M^{\dagger} ion at m/z 171, together with strong m/z 170 [M-H]⁺, 142 [M - CHO]⁺ and 115 [M - C₂H₂ON]⁺ fragment ions. Synthesis of 1: The synthesis of 1 was undertaken following the method of Reeves et al. (2007), Scheme 1. The synthesis of **5** proceeded readily with 67.5% yield but the subsequent conversion to **1** gave predominantly the by-product (E)-1-(2-methoxyphenyl)-3-(oxazol-4-yl)-prop-2-en-1-one, 6, and a lesser amount of its (Z)-isomer, which are the outcome of the dehydration of 5, together with only a small amount of 1. A significant quantity of o-methoxyacetophenone from the degradation of 5 was also present in the product mixture. Based on the mechanism proposed by Reeves et al. (2007) 6 is an intermediate in the formation of 1 from 5. It is probable that the subsequent nucleophilic attack by the N lone pair upon the carbonyl carbon is sterically hindered by the presence of the ortho methoxyl group and this prevents 1 from forming. Bioactivity of luteolin and 1: Notwithstanding the low yield (0.2%) of 1, the quantity of this compound that was isolated from the crude mixture by chromatography was sufficient for its bioactivity against S.aureus to be determined. 1 was found to be inactive at the concentrations screened (20 µg/mL-0.1325 μg/mL). On the other hand luteolin and its 4'-O-glucoside has been shown to have bactericidal activity against Staphylococcus aureus and Escherichia coli with MIC = 5.0×10^{-2} - 1.0×10^{-1}

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mg/mL), luteolin was also active against Bacillus cereus and Citrobacter freundii (MIC = 5.0×10^{-2}

mg/mL) and luteolin 3'-O-glucoside was active against *Bacillus cereus* and *Lactobacillus plantarum* with MIC = 2.5×10 -1 mg/mL and 5×10 -1 mg/mL respectively (Kumarasamy, Nahar, Byres, Delazar & Sarker, 2005). However the mean concentration at which luteolin is present in mānuka honey (0.14 ± 0.02 mg/100 g honey) shows that it would be unlikely to attain any of these minimum inhibitory concentrations in a well-diffusion assay as used in this study.

CONCLUSIONS

Analysis of the flavonoid profiles of a large sample set of mānuka honeys gave a mean total flavonoid content of 1.16 mg/100g of honey with pinobanksin, pinocembrin, luteolin and chrysin as the principal flavonoids present. Although the phenolic acids and flavonoids did not contribute directly to the non-peroxide antibacterial activity of the honey, luteolin and non-flavonoid compound 1 were weakly correlated with measured activity. 1 and the principal flavonoid which does not derive from propolis, luteolin, are potential marker compounds for mānuka monofloral honey and their correlation with non-peroxide antibacterial activity could be explained if one assumes that the activity of the honey is directly related to the proportion of mānuka in the blend.

1 was isolated and characterised by NMR and MS methods and its structure was confirmed by synthesis. 1 showed no bioactivity against S. aureus. The lack of bioactivity indicates that the moderate correlation ($R^2 = 0.36$) which 1 exhibited with non-peroxide antibacterial activity does not arise by a direct contribution from 1. At the concentration in which it is present in mānuka honey, luteolin is also unlikely to make a direct contribution to non-peroxide antibacterial activity. Given that the total phenolic fraction was inactive at the concentrations normally present in mānuka honey, there is no indication of synergy between 1 and other phenolics including luteolin.

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Captions for Tables and Figures

Fig. 1: HPLC chromatogram (gradient method 1) of flavonoid fraction of mānuka honey at 290 and 340 nm: (1) caffeic acid, (2) isoferulic acid, (3) *p*-coumaric acid, (4) pinobanksin, (5) unknown compound **1**, (6) unknown flavonoid 01, (7) luteolin, (8) unknown flavonoid 02, (9) pinocembrin, (10) unknown flavonoid 03, (11) unknown flavonoid 04, (12) unknown flavonoid 05, (13) chrysin, and (14) galangin.

Fig. 2: Scatterplot of (a) unknown compound **1** and (b) luteolin concentration *versus* UMF™ non-peroxide antibacterial activity.

Fig. 3: HPLC chromatogram (gradient method 2, 340 nm) of purified 1.

Fig. 4: HMBC correlations observed in 1.

Scheme 1: Synthesis of 1

Table 1: Retention times of phenolic acids and flavonoids found in the flavonoid fraction of mānuka honeys: pinobanksin (3,5,7-trihydroxyflavanone), unknown compound **1**, unknown flavonoid 1, quercetin (3,5,7,3',4'-pentahydroxyflavone), luteolin (5,7,3',4'-tetrahydroxyflavone), unknown flavonoid 2, 8-methoxykaempferol (3,5,7,4'-tetrahydroxy-8-methoxyflavone), pinocembrin (5,7-dihydroxyflavone), unknown flavonoid 3, isorhamnetin (3,5,7,4'-tetrahydroxy-3'-methoxyflavone), unknown flavonoid 4, kaempferol (3,5,7,4'-tetrahydroxyflavone), unknown flavonoid 5, chrysin (5,7-dihydroxyflavone), galangin (3,5,7-trihydroxyflavone), unknown flavonoid 6.

Table 2: Quantitation of flavonoids in mānuka honeys (UMF = non-peroxide antibacterial activity, mean of 12 determinations, other abbreviations as for Table 1)

Table 3: Non-peroxide antibacterial activity of fractions from XAD-2 separation of mānuka honey

Table 4: NMR signals assignments for **1** and **4** (δ ppm in DMSO- d_6)

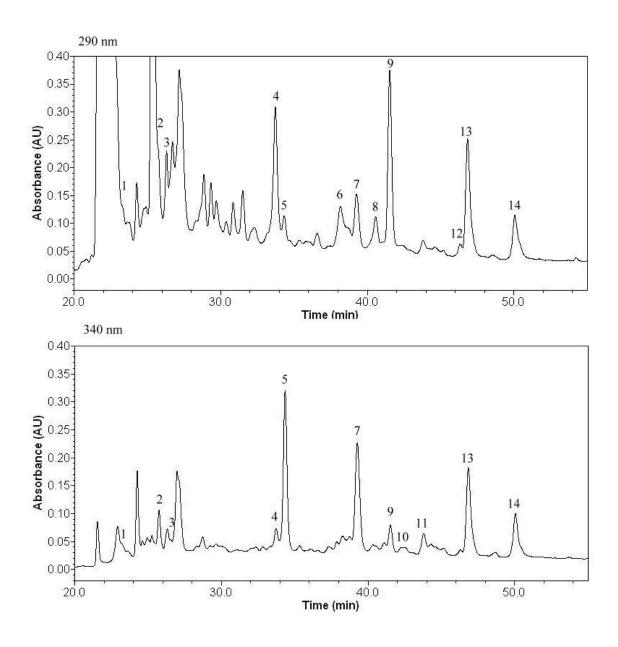
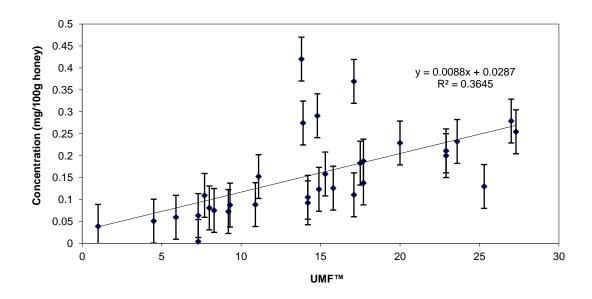
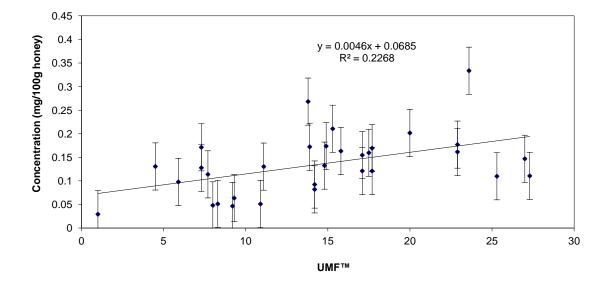


Figure 2

(a)



(b)



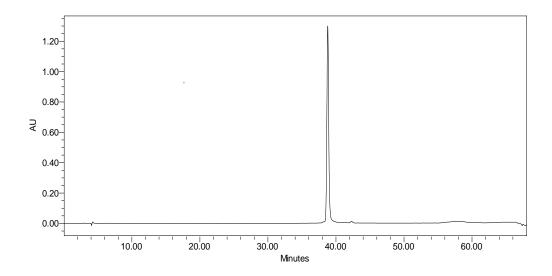


Figure 3

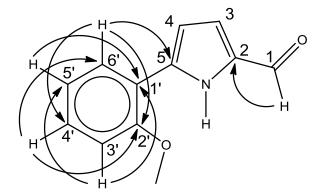


Figure 4

Scheme 1

Table 1

Name	Identification ^a	Quantification ^b	R _t (min) ^c	λ _{max} (nm) ^d
Caffeic acid	ı	е	23.3	325
Isoferulic acid	I	e	35.7	324
<i>p</i> -Coumaric acid	1	e	26.5	310
Pinobanksin (PB)	I	Р	33.9	292
1	I	Q	34.4	342
Unknown flavonoid 1(F1)	NA	Р	38.3	286
Quercetin (Q)	S	Q	38.7	367, 256
Luteolin (L)	S	Q	39.3	350, 255
Unknown flavonoid 2 (F2)	NA	Р	40.4	292
8-Methoxykaempferol (8MK)	I	K	41.2	375, 272
Pinocembrin (PC)	S	Р	41.5	290
Unknown flavonoid 3 (F3)	NA	Q	42.4	360, 258
Unknown flavonoid 4 (F4)	NA	K	43.7	361, 253
Isorhamnetin (IR)	S	K	44.3	373, 255
Kaempferol (K)	S	K	45.5	364, 255
Unknown flavonoid 5 (F5)	NA	Р	46.4	291
Chrysin (C)	S	С	46.7	314, 269
Galangin (G)	S	С	50.0	360, 266
Unknown flavonoid 6 (F6)	NA	С	50.6	311, 269

^a Method used to identify the compound: S = authentic standard, I = isolation and characterisation and NA = not identified.

^b Quantification standard used: Q = quercetin, K = kaempferol, P = pinocembrin and C = chrysin.

^c HPLC retention time.

 $^{^{\}rm d}$ Peak maxima in the UV absorbance spectrum (240 – 400 nm).

^e Phenolic Acids were not quantified as their presence in this fraction represents only a portion of their total presence in the honey.

Concentration of flavonoids (mg/100g of	honev)
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	Concentration of havonoids (mg/roog of honey)																	
Sample	UMF	РВ	1	F1	Q	L	F2	8MK	PC	F3	F4	IRM	K	F5	С	G	F6	Total
1	7.7	0.343	0.109	0.058	0.000	0.114	0.037	0.013	0.200	0.013	0.016	0.020	0.017	0.006	0.102	0.027	0.009	1.083
2	11.1	0.313	0.152	0.065	0.020	0.130	0.036	0.017	0.213	0.013	0.029	0.026	0.019	0.009	0.118	0.052	0.016	1.229
3	15.8	0.365	0.126	0.071	0.031	0.163	0.040	0.025	0.206	0.023	0.046	0.030	0.038	0.014	0.207	0.083	0.014	1.482
4	17.1	0.392	0.240	0.062	0.115	0.138	0.042	0.004	0.231	0.018	0.092	0.034	0.146	0.012	0.442	0.051	0.016	2.036
5	22.9	0.224	0.205	0.043	0.040	0.169	0.028	0.008	0.128	0.023	0.036	0.030	0.072	0.005	0.246	0.020	0.007	1.285
6	27	0.200	0.279	0.039	0.016	0.147	0.015	0.010	0.112	0.013	0.016	0.016	0.012	0.006	0.081	0.021	0.008	0.990
7	13.8	0.585	0.420	0.094	0.037	0.268	0.040	0.036	0.337	0.030	0.044	0.042	0.027	0.016	0.168	0.072	0.020	2.235
8	17.7	0.297	0.188	0.039	0.049	0.121	0.023	0.020	0.157	0.018	0.024	0.024	0.072	0.004	0.151	0.033	0.011	1.231
9	20	0.166	0.229	0.047	0.006	0.202	0.027	0.010	0.129	0.015	0.020	0.017	0.013	0.008	0.081	0.038	0.013	1.021
10	27.3	0.144	0.254	0.058	0.002	0.111	0.022	0.014	0.137	0.010	0.016	0.016	0.006	0.012	0.086	0.023	0.011	0.923
11	25.3	0.062	0.130	0.027	0.013	0.110	0.015	0.005	0.070	0.007	0.018	0.009	0.007	0.004	0.075	0.034	0.008	0.594
12	23.6	0.262	0.232	0.075	0.014	0.334	0.062	0.019	0.227	0.020	0.040	0.025	0.022	0.020	0.168	0.078	0.024	1.622
13	17.5	0.279	0.183	0.068	0.020	0.160	0.033	0.009	0.146	0.023	0.032	0.028	0.021	0.006	0.103	0.043	0.008	1.162
14	17.7	0.240	0.138	0.082	0.043	0.170	0.043	0.016	0.127	0.020	0.042	0.017	0.016	0.003	0.070	0.030	0.005	1.062
15	13.9	0.310	0.274	0.066	0.051	0.172	0.045	0.016	0.197	0.018	0.037	0.031	0.019	0.010	0.104	0.053	0.013	1.417
16	14.8	0.254	0.291	0.064	0.000	0.133	0.047	0.011	0.191	0.015	0.029	0.023	0.016	0.009	0.090	0.042	0.010	1.225
17	14.9	0.197	0.123	0.048	0.010	0.174	0.058	0.000	0.142	0.007	0.025	0.015	0.012	0.009	0.082	0.047	0.011	0.960
18	10.9	0.141	0.088	0.060	0.012	0.051	0.026	0.005	0.110	0.006	0.017	0.009	0.006	0.003	0.043	0.016	0.006	0.599
19	15.3	0.125	0.158	0.067	0.035	0.211	0.035	0.015	0.102	0.012	0.022	0.013	0.016	0.003	0.052	0.021	0.005	0.894
20	14.2	0.273	0.105	0.112	0.000	0.082	0.063	0.000	0.176	0.000	0.012	0.013	0.000	0.003	0.038	0.017	0.005	0.899
21	14.2	0.336	0.092	0.129	0.000	0.092	0.098	0.009	0.312	0.017	0.017	0.025	0.000	0.010	0.099	0.054	0.011	1.302
22	1	0.340	0.039	0.034	0.017	0.029	0.018	0.083	0.239	0.010	0.017	0.017	0.011	0.003	0.051	0.016	0.004	0.927
23	9.2	0.291	0.073	0.081	0.000	0.047	0.029	0.004	0.186	0.000	0.010	0.014	0.000	0.006	0.047	0.017	0.008	0.813
24	8.3	0.235	0.075	0.075	0.000	0.051	0.017	0.000	0.164	0.000	0.008	0.014	0.000	0.007	0.046	0.015	0.005	0.711
25	9.3	0.211	0.087	0.052	0.000	0.064	0.010	0.002	0.129	0.006	0.008	0.011	0.002	0.003	0.032	0.011	0.004	0.631
26	7.3	0.325	0.004	0.056	0.017	0.128	0.023	0.007	0.185	0.014	0.018	0.017	0.016	0.004	0.064	0.022	0.004	0.904
27	7.3	0.276	0.063	0.054	0.000	0.171	0.050	0.011	0.248	0.011	0.026	0.024	0.021	0.016	0.133	0.054	0.018	1.178
28	8	0.284	0.081	0.059	0.000	0.048	0.026	0.007	0.177	0.005	0.012	0.015	0.031	0.005	0.060	0.022	0.005	0.837
29	5.9	0.269	0.059	0.057	0.000	0.098	0.057	0.008	0.181	0.007	0.009	0.015	0.004	0.007	0.048	0.018	0.003	0.841
30	4.5	0.309	0.051	0.044	0.000	0.131	0.061	0.003	0.018	0.004	0.012	0.021	0.008	0.005	0.063	0.020	0.004	0.754
31	0	0.264	0.064	0.049	0.000	0.144	0.045	0.005	0.198	0.007	0.016	0.019	0.021	0.004	0.061	0.024	0.005	0.926
Mean		0.273	0.157	0.061	0.024	0.136	0.038	0.012	0.174	0.013	0.029	0.022	0.032	0.008	0.131	0.035	0.010	1.155
S.D.		0.095	0.096	0.021	0.043	0.062	0.018	0.014	0.064	0.008	0.029	0.009	0.051	0.005	0.132	0.020	0.005	0.461
95% C.I.		0.034	0.034	0.008	0.015	0.022	0.006	0.005	0.023	0.003	0.010	0.003	0.018	0.002	0.047	0.007	0.002	0.164
CV (%)		34.9	61.3	34.7	175.8	45.7	47.4	120.7	36.6	60.8	99.5	43.3	161.5	60.3	100.4	56.4	56.8	39.9
% of total		23.6	13.6	5.3	2.1	11.8	3.3	1.0	15.1	1.2	2.5	1.9	2.7	0.7	11.4	3.0	0.8	

Table 2

Table 3

Non-Peroxide Antibacterial Activity					
(% Phenol Equivalents)					
7 ª					
table activity ^a					
3^{a}					
table activity ^b					
)					

^aMean and 95% confidence interval from 48 determinations.

^bMean and 95% confidence interval from 32 determinations.

Table 4

	1		4ª	
	¹³ C	¹H	¹³ C	¹H
1-NH		11.79 (br s)	12.39 (l	or s)
2	132.5		133.8	
3	120.5	7.04 ^b (d, <i>J</i> = 3.9 Hz)	121.9	7.08 (dd, <i>J</i> = 3.9, 2.3 Hz)
4	111.4	6.77 (d, <i>J</i> = 3.9 Hz)	108.9	6.79 (dd, <i>J</i> = 3.9, 2.3 Hz)
5	136.5		139.5	
2- <u>CH</u> O	178.8	9.52 s	178.8	9.50 s
1'	119.2		130.9	
2'	156.2		125.5	7.89 (d, <i>J</i> = 8.0 Hz)
3'	112.1	7.14 (dd, <i>J</i> = 8.5, 1.2 Hz)	128.7	7.42 (t, <i>J</i> = 8.0 Hz)
4'	129.4	7.33 (ddd, <i>J</i> = 8.5, 7.3, 1.7 Hz)	127.9	7.32 (t, <i>J</i> = 8.0 Hz)
5'	120.7	7.02 ^b (~td, <i>J</i> = 8.3, 1.2 Hz)	128.7	7.42 (t, <i>J</i> = 8.0 Hz)
6'	128.4	7.77 (dd, <i>J</i> = 7.8, 1.7 Hz)	125.5	7.89 (d, <i>J</i> = 8.0 Hz)
2'-OCH ₃	55.6	3.96 s		

^asample kindly supplied by Dr Reeves; ^bsignals partly overlapped