

## Supporting Information

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## Table of Contents

Synthesis

Characterization

<sup>1</sup>H NMR, <sup>13</sup>C NMR, MS spectra for methyl 2-(2,5-di(thiophen-2-yl)-1H-pyrrol-1-yl)acetate

<sup>1</sup>H NMR, <sup>13</sup>C NMR, HRMS spectra for 2-(2,5-di(thiophen-2-yl)-1H-pyrrol-1-yl)acetic acid

Sample Application

## Synthesis

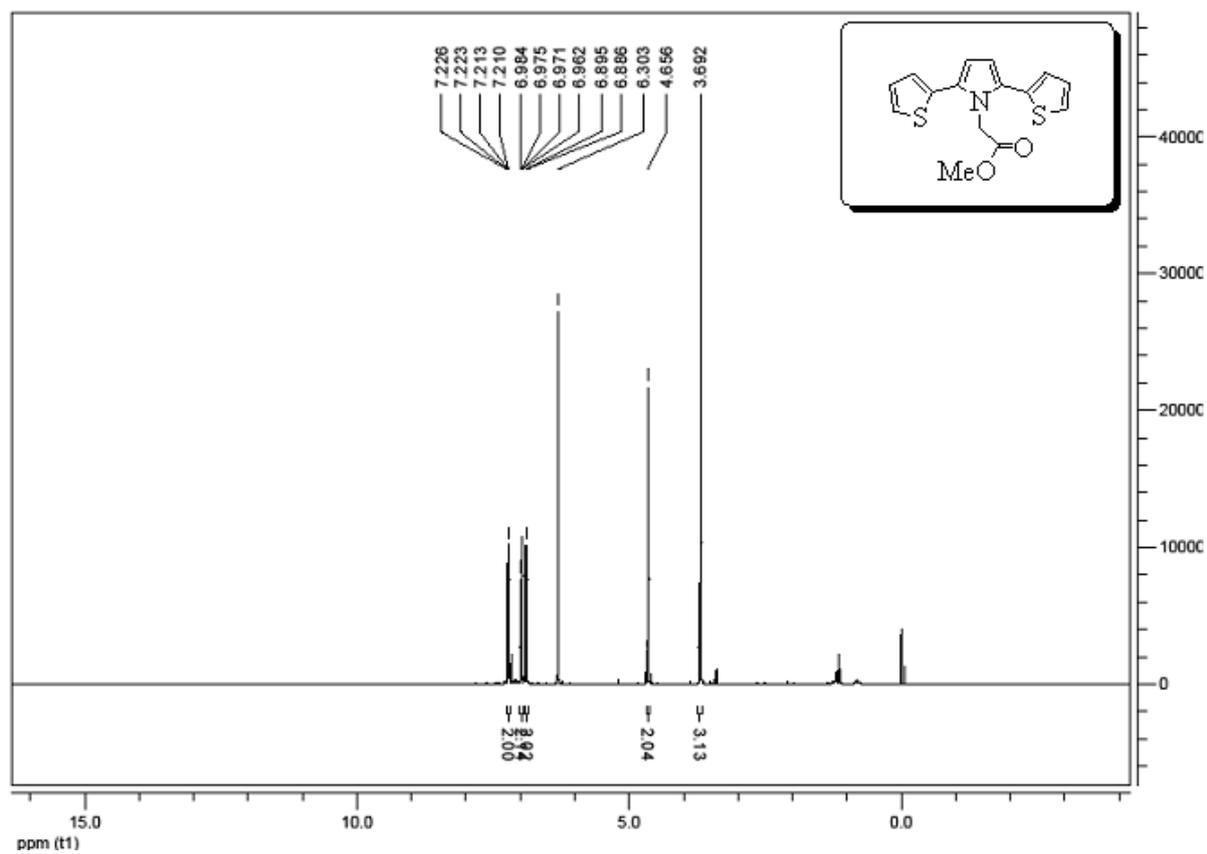
Flash column chromatography was performed using thick-walled glass columns with a flash grade silica gel (Merck Silica Gel 60). Reactions were monitored by thin layer chromatography using precoated silica gel plates (Merck Silica Gel PF-254), visualized by UV-light and polymolybdene phosphoric acid in ethanol. <sup>1</sup>H and <sup>13</sup>C NMR were recorded in CDCl<sub>3</sub> and the chemical shifts were expressed in ppm relative to CDCl<sub>3</sub> (δ 7.26 and 77.0 for <sup>1</sup>H and <sup>13</sup>C NMR, respectively) as the internal standard.

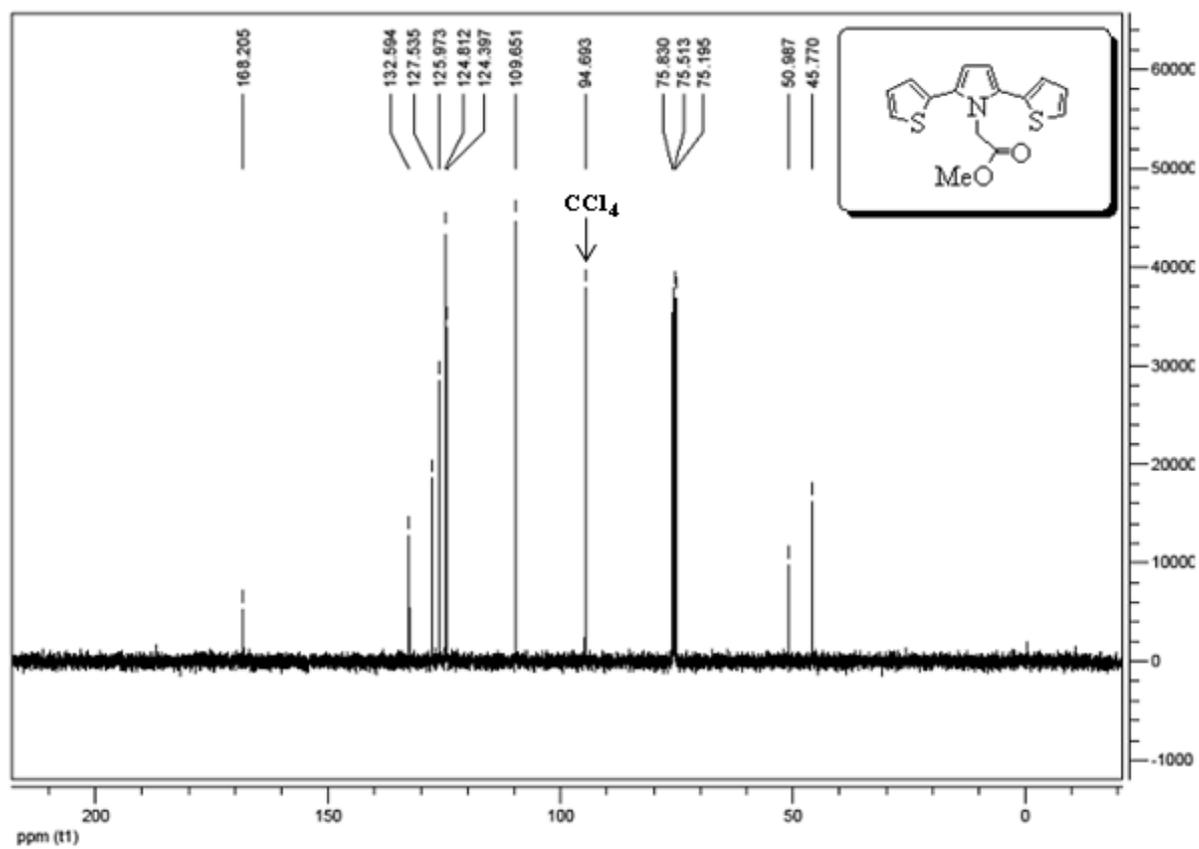
## Characterization

XPS (X-ray Photoelectron Spectroscopy) was carried out on a PHI 5000 Versa Probe ( $\Phi$  ULVAC-PHI, Inc., Japan/USA) model X-ray photoelectron spectrometer instrument with monochromatized Al K $\alpha$  radiation (1486.6 eV) as an X-ray anode at 24.9 W. The pressure inside the analyzer was maintained at  $10^{-7}$  Pa. The binding energy scale was referenced by setting the C–H peak maximum in the C 1s spectrum to 285.0 eV and the atomic composition estimated using Multipak software.

### **<sup>1</sup>H NMR, <sup>13</sup>C NMR, MS spectra for methyl 2-(2,5-di(thiophen-2-yl)-1H-pyrrol-1-yl)acetate**

<sup>1</sup>H-NMR (CDCl<sub>3</sub>+CCl<sub>4</sub>):  $\delta$  7.22 (dd, J=1.0 Hz, J=5.1 Hz, 2H), 6.97 (dd, J=3.5 Hz, J=5.2 Hz, 2H), 6.89 (dd, J=1.0 Hz, J=3.5 Hz, 2H), 6.30 (s, 2H), 4.66 (s, 2H), 3.69 (s, 3H). <sup>13</sup>C-NMR (CDCl<sub>3</sub>+CCl<sub>4</sub>): 168.2, 132.6, 127.5, 126.0, 124.8, 124.4, 109.6, 51.0, 45.8. MS (EI) m/z (relative intensity): 303.0 (100), 304.0 (18), 305.0 (10), 244.0 (37), 230.0 (22).

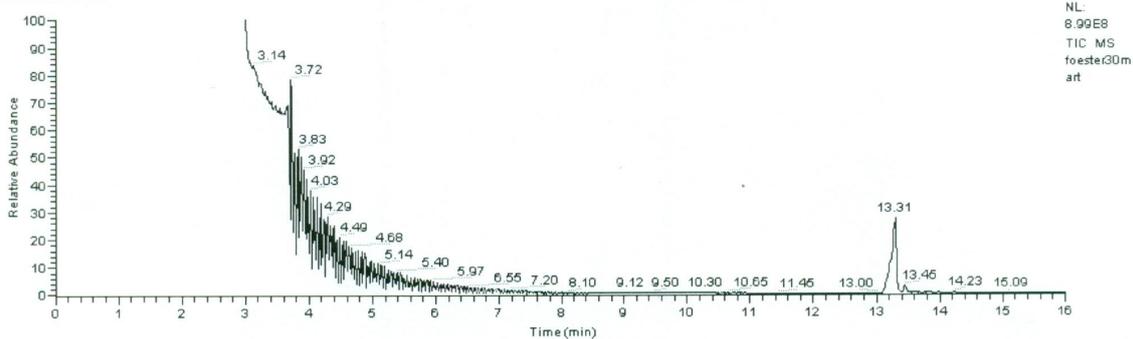




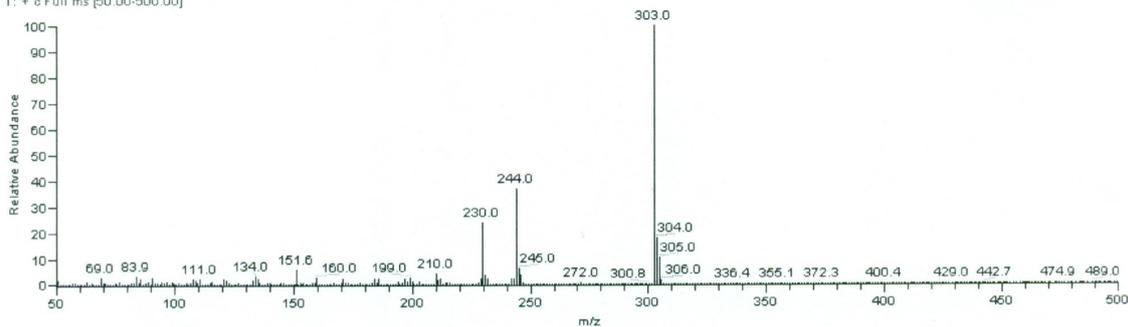
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foester

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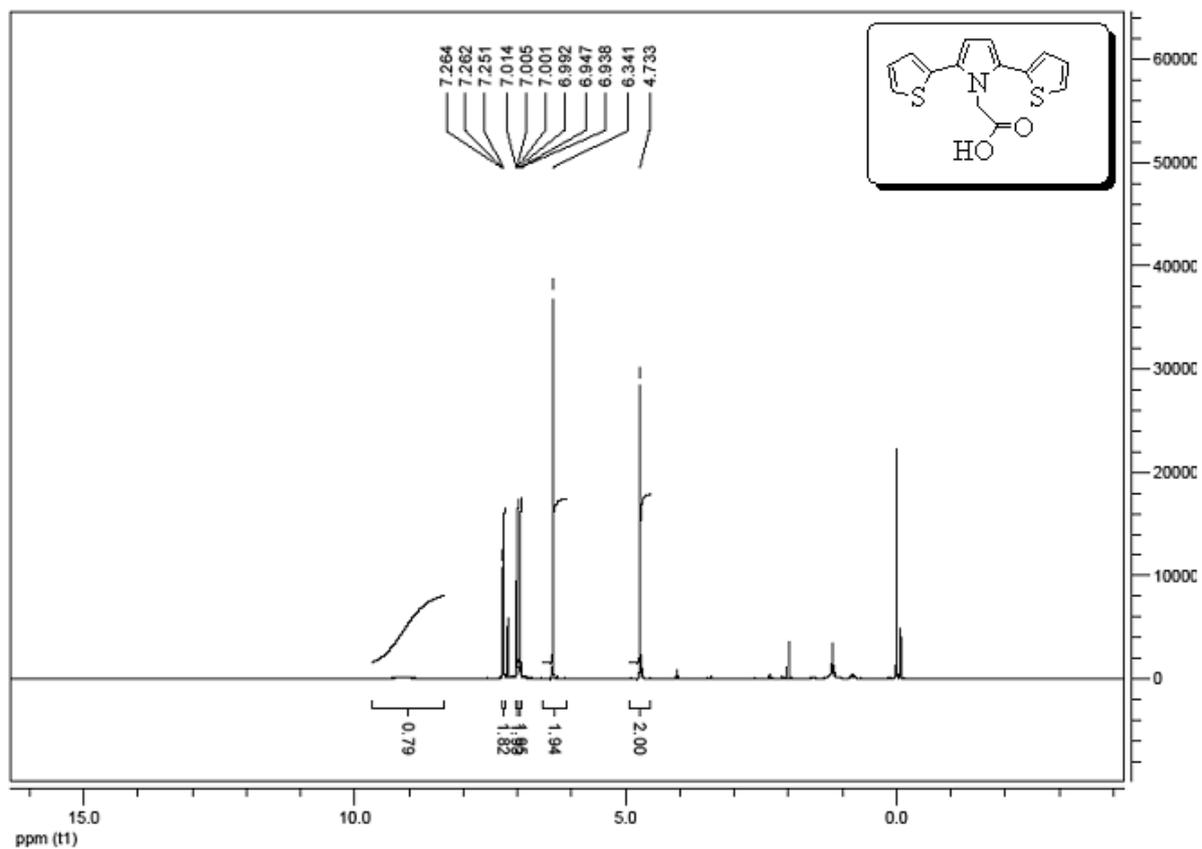


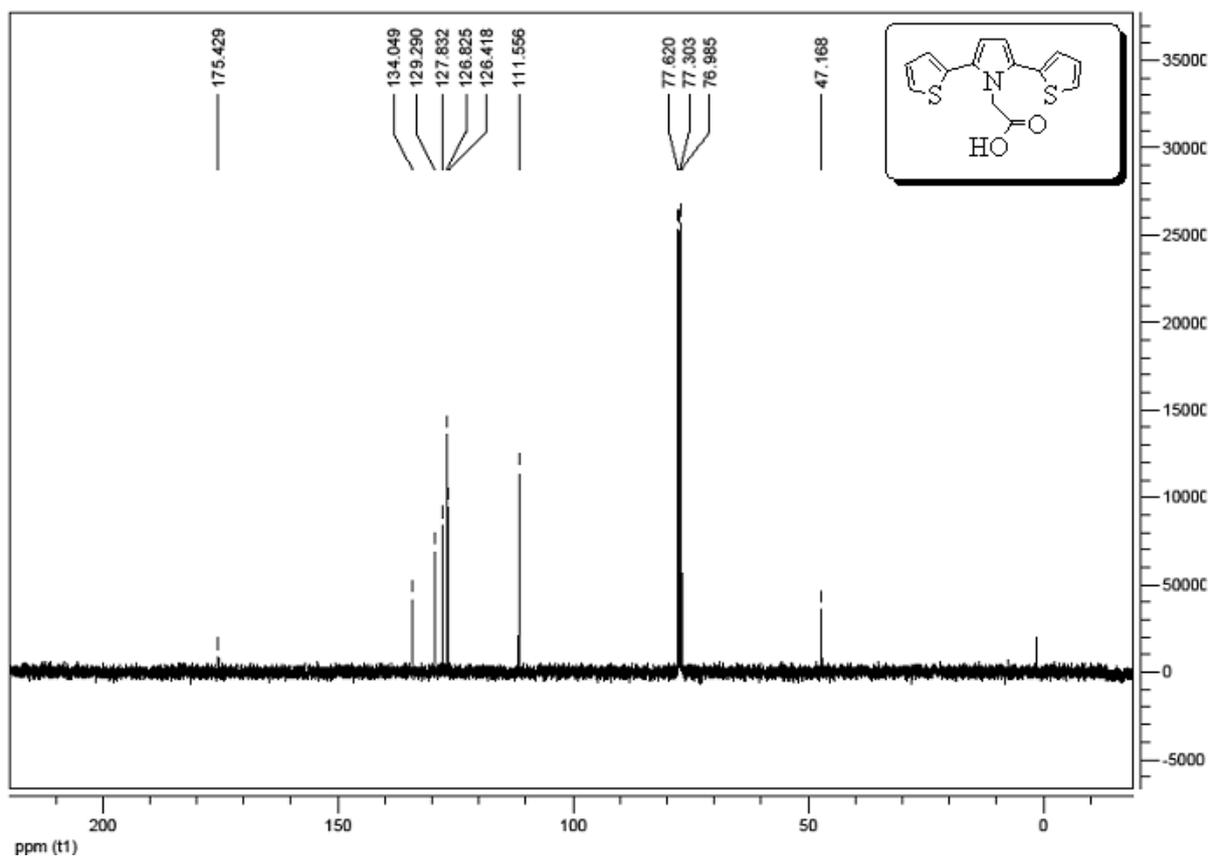
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### **<sup>1</sup>H NMR, <sup>13</sup>C NMR, HRMS spectra for 2-(2,5-di(thiophen-2-yl)-1H-pyrrol-1-yl)acetic acid**

<sup>1</sup>H-NMR (CDCl<sub>3</sub>): δ 9.1 (s, 1H), 7.26 (dd, J=1.0 Hz, J=5.2 Hz, 2H), 7.00 (dd, J=3.6 Hz, J=5.1 Hz, 2H), 6.94 (dd, J=1.0 Hz, J=3.5 Hz, 2H), 6.34 (s, 2H), 4.73 (s, 2H). <sup>13</sup>C-NMR (CDCl<sub>3</sub>): 175.4, 134.0, 129.3, 127.8, 126.8, 126.4, 111.6, 47.2. MS (EI) m/z (relative intensity): 289.0. HRMS: Calculated [M]<sup>+</sup> 289.0231, Measured [M]<sup>+</sup> 289.0229.





### Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 100.0  
Element prediction: Off  
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Odd and Even Electron Ions  
8 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)

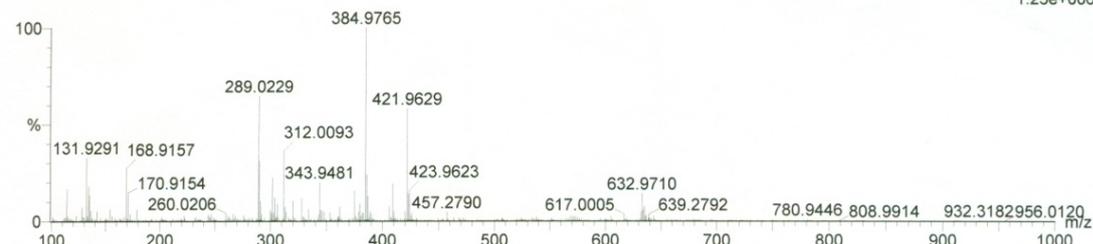
Elements Used:

C: 0-30 H: 0-30 N: 1-1 O: 2-2 S: 0-2

5339 CIHANGER TANYELI

20100922\_FND2\_03 282 (0.983) Cm (1:284)

1: TOF MS ES+  
1.25e+006



Minimum: -1.5  
Maximum: 100.0 10.0 100.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	i-FIT (Norm)	Formula
289.0229	289.0231	-0.2	-0.7	10.0	911.4	0.0	C14 H11 N O2 S2

### Sample Application

*G. oxydans* DSMZ 2343 was obtained from the German Collection of Microorganisms and Cell Cultures (Braunschweig, Germany) and was sub-cultured in a medium containing (in g/L): glucose, 100; yeast extract, 10; calcium carbonate, 20; agar, 20. Cells were cultivated in shake flasks on a shaker (175 rpm) at 28 °C, in a medium containing (in g/L): yeast extract, 5; carbon source: glucose, 5.<sup>1</sup> For sample application, 1 mL of cultivation medium was collected hourly for 5 hours. After centrifugation of the fermentation medium, the samples were applied to the biosensor and the glucose content of the samples was calculated in reference to HPLC method.

HPLC with a refractive index detector (RID) controlled by a HP-Chemstation (Agilent, Karlsruhe, Germany) was used as the reference method for independent analysis of the glucose content. HPLC column (GL Sciences Inc. Inertsil NH<sub>2</sub> 5.0 μm (4.6 I.D x 250 mm), Japan) was used for the chromatographic separation at 30 °C. Injection volume was 20 μL.

The mobile phase was H<sub>2</sub>SO<sub>4</sub> (5 mM).<sup>2</sup> The flow rate was 0.6 mL/min. Initially a standard calibration curve for glucose was plotted (0.1-50 mM for glucose with the equation of

$y=303971.24x$ ;  $R^2=0.999$ ). After dilution with the mobile phase and a centrifugation step the samples were applied to the column and then glucose concentrations were calculated using the calibration plot.

- 1 J. Tkac, I. Vostiar, L. Gorton, P. Gemeiner and E. Sturdik, *Biosens. Bioelectron.* 2003, **18**, 1125-1134.
- 2 A. Plaga, J. Stumpfel and H. P. Fiedler, *Appl. Microbiol. Biot.*, 1989, **32**, 45-49.