

Supplementary Information

Transition metal cations catalyze $^{16}\text{O}/^{18}\text{O}$ exchange of catechol motifs with H_2^{18}O

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Table of Contents

Materials and methods.....	S2
HCD fragmentation spectra of unlabeled and doubly ^{18}O labeled catechol derivatives (Fig. S1)	S4
Simulated complexation of metal cations with catechol as function of pH (Fig. S2)	S5
$^{16}\text{O}/^{18}\text{O}$ exchange of catechol with H_2^{18}O in presence of MnCl_2 (Fig. S3)	S6
Comparison of $^{16}\text{O}/^{18}\text{O}$ exchange in presence or absence of air (Fig. S4)	S6
RP-UHPLC-PDA-ESI-MS chromatograms and spectra (Fig. S5)	S7

Materials and methods

Materials

Catechol (≥ 99 wt%), 4-chlorocatechol (≥ 97 wt%), 4-methylcatechol (≥ 95 wt%), 4-nitrocatechol (≥ 97 wt%), dopamine hydrochloride (≥ 98 wt%). Copper(II) chloride (≥ 97 wt%), aluminum(III) chloride hexahydrate (≥ 99 wt%), hydrochloric acid (HCl, 37 wt%), sodium hydroxide (NaOH, ≥ 98 wt%) and ethanol (EtOH) were obtained from Merck (St. Louis, MO, USA). 3,4-Dihydroxybenzoic acid (≥ 98 wt%) was purchased from TCI Europe NV (Zwijndrecht, Belgium). 4-Methoxycatechol (≥ 95 wt%) was purchased from Toronto Research Chemicals (North York, Canada). Iron(III) chloride (≥ 97 wt%), zinc(II) chloride (≥ 98 wt%), and 3,4-dihydroxybenzaldehyde (≥ 95 wt%) were purchased from Fisher Scientific (San Jose, CA, USA). H_2^{18}O (GMP grade, $^{18}\text{O} \geq 97\%$) was purchased from ABX advanced biochemical compounds-Biomedizinische Forschungsreagenzien GmbH (Radeberg, Germany). ULC-MS grade acetonitrile (ACN) and water, both containing 0.1 vol.% formic acid (FA) were purchased from Biosolve (Valkenswaard, the Netherlands). Water for other purposes than UHPLC was prepared using a Milli-Q water purification system (Merck Millipore, Billerica, MA, USA).

General experimental procedure

High resolution RP-UHPLC-PDA-ESI-MS of catechol and derivatives

Stock solutions of catechol and derivatives in acetone were added to an Eppendorf and the acetone was evaporated under nitrogen flow. Water (H_2^{18}O) and metal chloride solution (10 mM in H_2^{18}O) were added to obtain a final concentration of 1.0 mM of metal (*i.e.*, FeCl_3 , CuCl_2 , AlCl_3 , and ZnCl_2) and ligand for the initial screening with catechol and 3,4-DHBA and 0.1 mM of FeCl_3 and catechol and derivatives (*i.e.*, 4-chlorocatechol, 4-methylcatechol, 3,4-DHBA, dopamine, 4-nitrocatechol, 3,4-DHBZ, and 4-methoxycatechol) for the continuation experiments. The pH was adjusted to 3 or 7 for the initial screening and to 3 for the continuation experiments using 0.5 M NaOH or HCl prepared in H_2^{18}O . Samples were incubated at 37 °C at 1000 rpm using an Eppendorf Thermomixer® F1.5 (Eppendorf, Hamburg, Germany). For UHPLC-MS analysis, 80 μL aliquots of the reaction mixture were taken after 2 and 24 h for the initial screening and 0.5, 1, 4, and 24 h for the continuation experiments, and centrifuged (5 min, $15,000 \times g$) prior to RP-UHPLC-PDA-ESI-MS analysis.

RP-UHPLC-PDA-ESI-MSⁿ analysis was performed using a Thermo Vanquish UHPLC system (Thermo Scientific, San Jose, CA, USA), equipped with a pump, degasser, and autosampler, coupled to a PDA detector and a Thermo Q Exactive Focus hybrid quadrupole-orbitrap mass spectrometer. Samples (1 μL) were injected onto an Acquity UPLC BEH C18 column (150 \times 2.1 mm, particle size 1.7 μm) with a VanGuard guard column (5 \times 2.1 mm) of the same material (Waters, Milford, MA, USA). The flow rate was 400 $\mu\text{L min}^{-1}$ and the column temperature was 45 °C. Water (A) and ACN (B), both acidified with 0.1% FA, were used as eluents. The following solvent gradient was used: 0-1.09 min at 1% B (isocratic), 1.09-6.18 min from 1 to 15% B (linear gradient), 6.18-7.27 min from 15 to 100% B (linear

gradient), 7.27-12.72 min at 100% B (isocratic), 12.72-13.81 min from 100 to 1% B (linear gradient) and 13.81-19.27 min at 1% B (isocratic). The PDA detector was set to measure the wavelength range of 190–680 nm. Mass spectrometric data were collected over the m/z range of 100–1,000 in negative and positive ionization mode by using source voltages of 2.5 and 3.5 kV, respectively. Nitrogen was used as a sheath gas (50 arbitrary units) and auxiliary gas (13 arbitrary units). For both modes, the S-lens RF level was set at 50 %, the ion transfer tube temperature was 263 °C and the source heater temperature 425 °C. Data acquisition and processing were performed using Xcalibur (version 4.1, Thermo Scientific). Recoveries of catechol and derivatives were calculated based on integrals of the chromatographic UV₂₈₀ peaks and comparison to an external calibration curve of the corresponding authentic standard (0.0125-0.1 mM, in duplicate, $R^2 > 0.99$). Labeling yields were determined as the average distribution of the accurate masses corresponding to unlabeled, singly labeled and doubly labeled compound under the chromatographic peak. The labeling yield was corrected for labeling percentage of the used H₂¹⁸O (97%), by dividing the uncorrected percentage of doubly ¹⁸O labeled catechols by 0.941 (the theoretical percentage expected after complete ¹⁸O labeling of the two catechol hydroxyl groups; 0.97²).

Time-resolved ESI-MS of catechol and derivatives in H₂¹⁸O in presence of FeCl₃

Solutions (400 μL) of catechol and derivatives at 0.2 mM were prepared in H₂¹⁸O and placed in a glass vial, the pH of the solution was adjusted to 3 by 0.5 M HCl in H₂¹⁸O. The solution was stirred at 37 °C for 5 minutes after which it was directly infused into the ESI-IT-MS source via a capillary by applying a slight overpressure of N₂ (approximately 0.25 bar). When the MS signal was stable, the measurement was started and after 0.5 min measurement of the ligand, a solution (400 μL) of 0.2 mM FeCl₃ in H₂¹⁸O (pH 3) was added to the glass vial to obtain final concentrations of 0.1 mM catecholate and iron. The ESI-IT-MS measurements were acquired using an LTQ Velos Pro linear ion trap mass spectrometer (Thermo Scientific) equipped with a heated electrospray ionization (ESI) probe. Data were collected for 60 min over the m/z range of 100-600 in negative or positive ionization mode by using source voltages of 4.0 and 3.5 kV in positive ionization (PI) and negative ionization (NI) mode, respectively. The S-lens RF level was set at 67.9% in PI and 69.5% in NI, the ion transfer tube and the source heater temperatures were 275 and 45 °C, respectively. Nitrogen was used as a sheath gas (50 arbitrary units) and auxiliary gas (13 arbitrary units). Data were processed using Xcalibur 4.1 (Thermo Scientific).

HCD fragmentation spectra of unlabeled and doubly ^{18}O labeled catechol derivatives

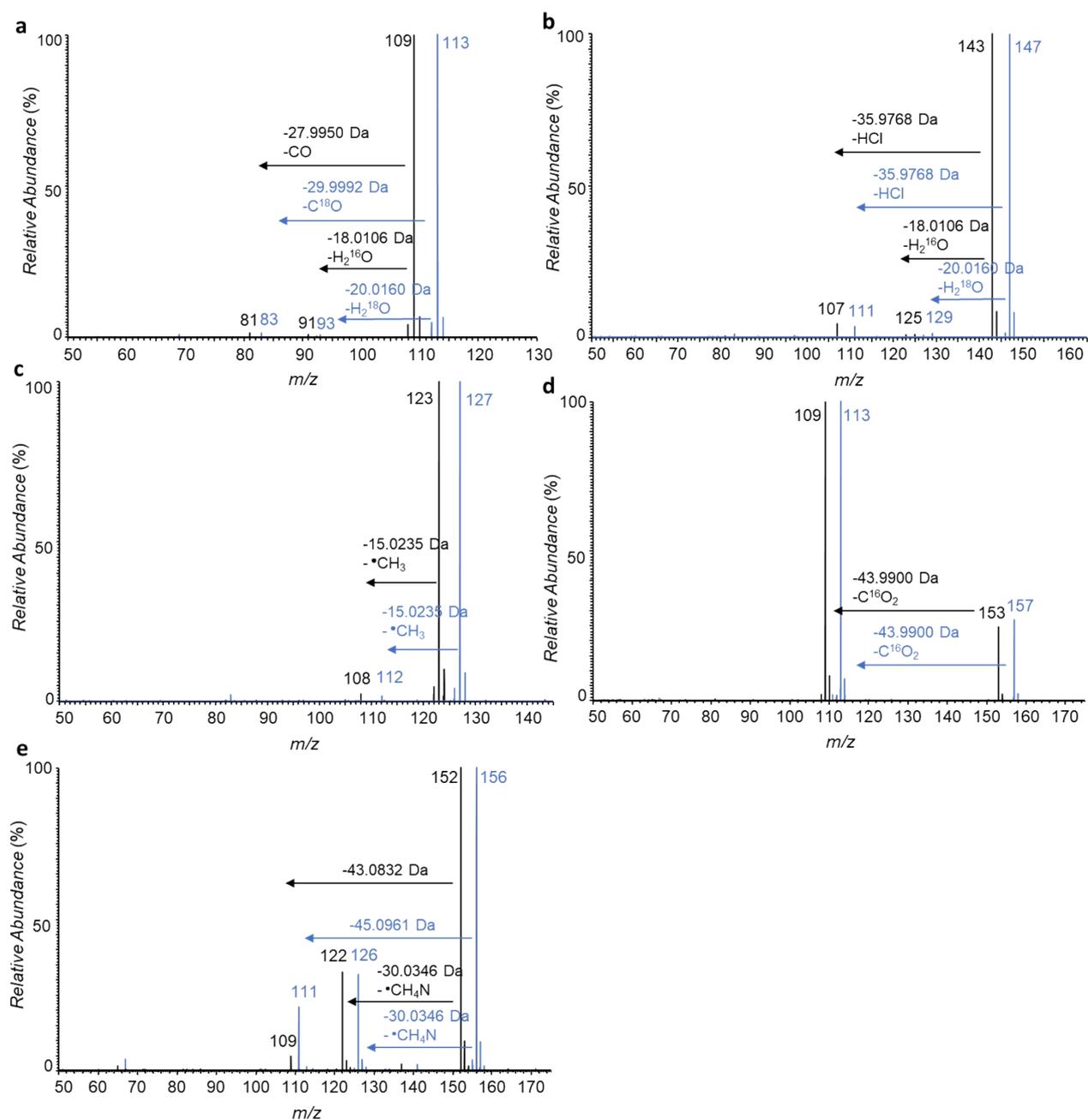


Fig. S1 HCD fragmentation spectra of unlabeled (black) and doubly ^{18}O -labeled catechol (a), 4-chlorocatechol (b), 4-methylcatechol (c), 3,4-dihydroxybenzoic acid (d), and dopamine (e).

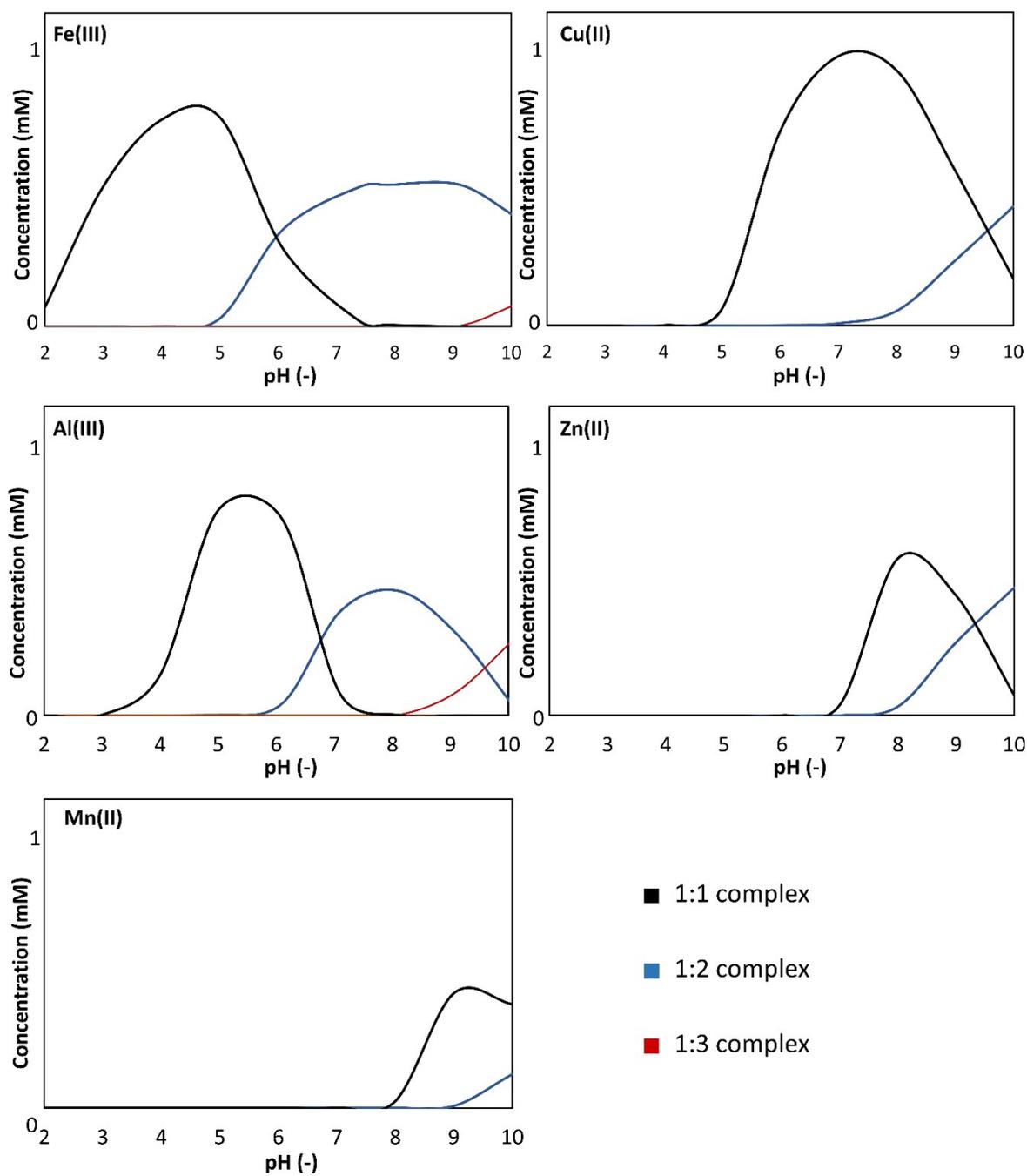


Fig. S2 Predicted concentration of 1:1, 1:2, and 1:3 complexes of catechol with Fe(III), Cu(II), Al(III), Zn(II) and Mn(II) at different pH. Visual MINTEQ (version 3.1) was used for the simulation. The metal-catechol and metal hydrolysis species and stability constants ($\log K$) were obtained from the standard database in Visual MINTEQ 3.1 and not fixed to a constant ionic strength.

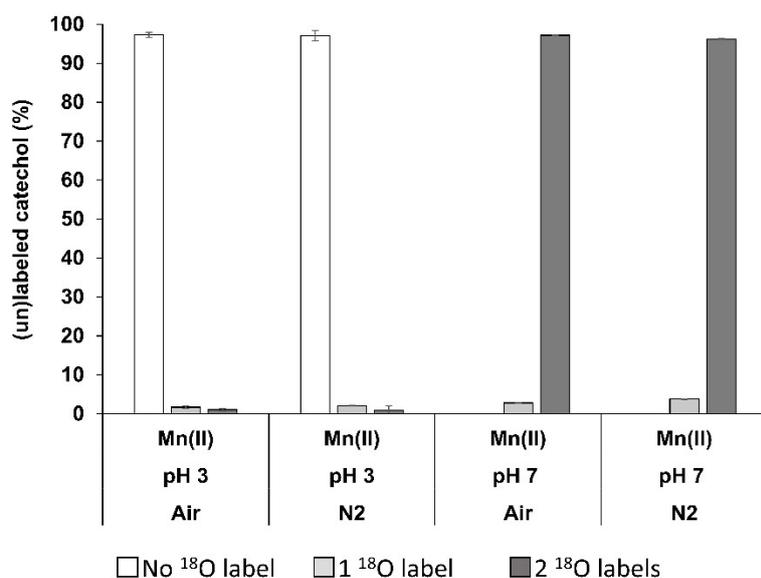


Fig. S3 Percentages of unlabeled, singly labeled, and doubly labeled catechol after equimolar incubations (1 mM) with MnCl₂ at 37 °C for 2 h, at pH 3 or 7 in presence of air or after purging with N₂. Purging with N₂ was performed for 2 min through the H₂¹⁸O solvent and headspace of the Eppendorf tube. Data are presented as average and standard deviation of two separate incubations. The large difference in ¹⁶O/¹⁸O exchange between pH 3 and 7 matches with the low tendency of Mn(II) to form catecholato complexes at low pH.

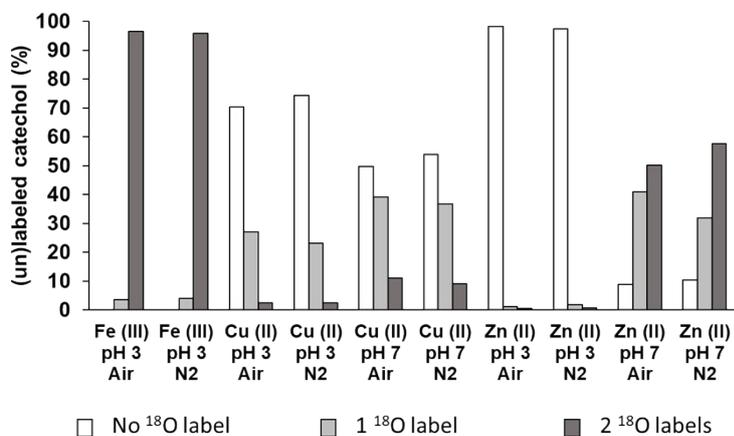


Fig. S4 Relative abundance of unlabeled, singly labeled and doubly labeled catechol in equimolar (1 mM) incubations of catechol with FeCl₃, CuCl₂, or ZnCl₂ at pH 3 or 7 in presence of air or after purging with N₂. Purging with N₂ was performed for 2 min through the H₂¹⁸O solvent and headspace of the Eppendorf tube. Incubation times were 30 min for the samples containing FeCl₃ and 24 h for the samples containing CuCl₂ or ZnCl₂.

RP-UHPLC-PDA-ESI-MS chromatograms and spectra

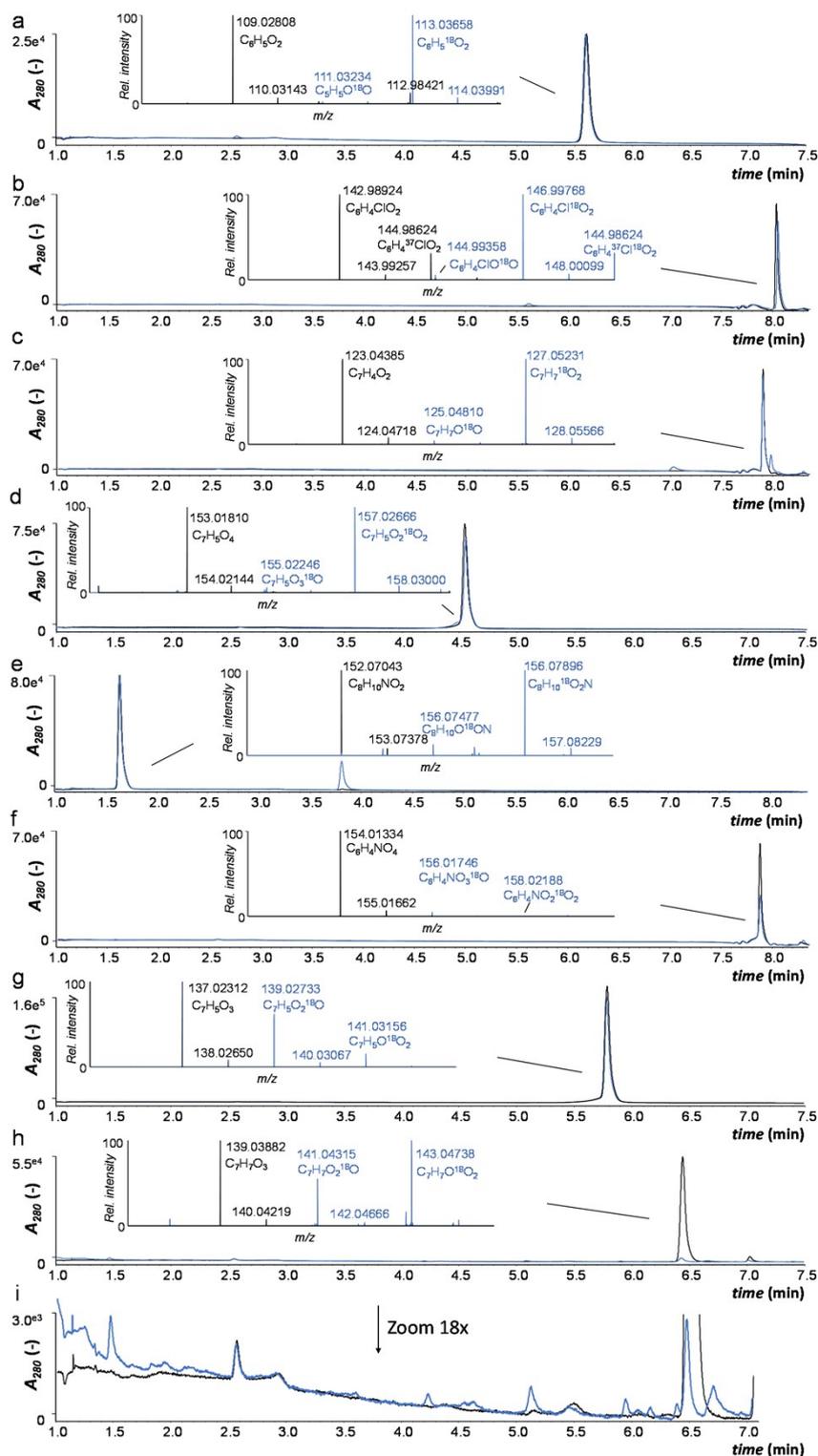


Fig. S5 Overlaid RP-UHPLC-UV₂₈₀ chromatograms and HRMS spectra of catechol (a), 4-chlorocatechol (b), 4-methylcatechol (c), 3,4-dihydroxybenzoic acid (d), dopamine (e), 4-nitrocatechol (f), 3,4-dihydroxybenzaldehyde (g), and 4-methoxycatechol (h) after incubation at 37 °C in absence (black) and presence (blue) of FeCl₃. Incubation times were 1 h for a-c and e, 4 h for d and 24 h for f-i. Chromatogram i is an 18-fold zoomed version of h, displaying several (unidentified) degradation products of 4-methoxycatechol. Equimolar concentrations (0.1 mM) were used in all incubations. The inserts depict the overlaid high resolution Full MS spectra under the corresponding peaks. Masses without molecular formula label correspond to the natural ¹³C isotope peak of the preceding peak.