# Inclusion and release of ant alarm pheromones from metal-organic frameworks

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**Electronic supplementary information** 

# 1. General experimental details

All reagents and solvents were purchased from commercial sources and used without further purification.

Powder X-ray diffraction (PXRD) patterns for all samples were recorded on a Bruker AXS D8 Advance diffractometer with copper  $K_{\alpha}$  radiation ( $\lambda = 1.5406$  Å) at 298 K. The beam slit was set to 1 mm, detector slit set to 0.2 mm and anti-scattering slit set to 1 mm. Samples were dried in ambient conditions and ground using a pestle and mortar. The powders were packed onto a flat plate and measured with a 2 $\theta$  range of 5 – 60°. The step size was 0.024° with the scan speed set to 0.3 s per step.

<sup>1</sup>H NMR spectra were recorded on the digested MOFs at 298 K on a Bruker Avance 300 MHz Ultrashield spectrometer or an Agilent 500 MHz ProPulse spectrometer. All <sup>1</sup>H NMR spectra were referenced to the residual *protio* peaks at  $\delta$  2.50 ppm for DMSO-*d*<sub>6</sub>. For the UiO-66 materials, a typical MOF digestion was carried out by adding 10 mg of a crystalline sample into 0.4 mL of DMSO-*d*<sub>6</sub> and 0.2 mL of a stock solution of NH<sub>4</sub>F in D<sub>2</sub>O (4.14 M). For the IRMOF materials, the MOF digestions were each carried out using approximately 10 mg of crystalline sample in 0.4 mL of DMSO-*d*<sub>6</sub> and 0.2 mL of a stock solution of 0.1 mL of 35 wt% DCl/D<sub>2</sub>O in 3 mL DMSO. In all cases, the mixtures were sonicated until all solids had completely dissolved.

IRMOF-1,<sup>15,18</sup> IRMOF-3,<sup>15,19</sup> IRMOF-NHPr,<sup>20</sup> IRMOF-NHBu,<sup>20</sup> IRMOF-NHOc<sup>20</sup> and Zn-MOF-74<sup>17</sup> were prepared as previously reported.

### 2. Synthesis of UiO-66-NHPr, [Zr<sub>6</sub>O<sub>4</sub>(OH)<sub>4</sub>(bdc-NHPr)<sub>6</sub>]

The synthesis of UiO-66-NHPr was performed by using an analogous procedure to that of UiO-66-NH<sub>2</sub>.<sup>23</sup> In a typical reaction, H<sub>2</sub>bdc-NHPr (234 mg, 1.05 mmol) along with ZrCl<sub>4</sub> (243 mg, 1.05 mmol) and DMF (12 mL) were loaded into a Teflon lined autoclave. The solution was stirred until the reactants had completely dissolved. The autoclave was placed in an oven and heated at 120 °C for 24 h. The resulting yellow powder was rinsed and centrifuged with MeOH (6000 rpm for 15 min) to remove unreacted H<sub>2</sub>bdc-NHPr and residual DMF in the pores. The washing procedure was repeated over 3 days with the solvent replaced every 24 h. Finally, the UiO-66-NH<sub>2</sub> powder was dried under vacuum at 120 °C for 12 h. Figure S1 shows the powder X-ray diffraction pattern for UiO-66-NHPr in comparison to that simulated from the crystal structure of UiO-66, confirming that UiO-66 and UiO-66-NHPr are isostructural.



Figure S1. PXRD pattern of UiO-66-NHPr (red) shown together with the simulated PXRD pattern calculated from the UiO-66 crystal structure.<sup>23</sup>

#### 3. Preparation of 3-octanone-loaded IRMOF-NHPr

#### Method A: Using four equivalents of 3-octanone

IRMOF-NHPr crystals (116 mg, 0.37 mmol) were added in a glass vial containing 5 mL DMF. 3-Octanone (0.23 mL, 1.48 mmol) was then added and the vial was sealed. The mixture was agitated briskly before leaving it to stand at room temperature for 3 days. The crystals were then isolated *via* filtration and rinsed once with fresh DMF (3 mL) to remove any residual 3-octanone on the crystal surfaces. The crystals were rinsed a further once or twice with fresh DMF (3 mL each wash). The NMR spectrum of the sample upon DCl digestion is shown in Figure S2.



Figure S2. The <sup>1</sup>H NMR spectrum for the digested 3-octanone-loaded IRMOF-NHPr.

The 3-octanone loading was determined by comparing the integrals at  $\delta$  7.14 ppm and  $\delta$  1.45 ppm which correspond to the aryl protons from D<sub>2</sub>BDC-NHPr, H<sub>i</sub> and alkyl protons, H<sub>d</sub> from 3-octanone, respectively. The ratio of D<sub>2</sub>bdc-NH<sub>2</sub> to 3-octanone was calculated as being approximately 1:0.57 which gives the chemical formula as [Zn<sub>4</sub>O(bdc-NHPr)<sub>3</sub>]·1.7oct (23.2 wt% 3-octanone loading).

## Method B: Using an excess of 3-octanone

IRMOF-NHPr crystals (116 mg, 0.37 mmol) were added in a glass vial containing neat 3octanone. The vial was sealed and the mixture was agitated briskly before leaving it to stand at room temperature for 3 days. The crystals were then isolated via filtration and rinsed once with fresh hexane (3 mL) to remove any residual 3-octanone on the crystal surfaces. The crystals were rinsed a further once or twice with fresh hexane (3 mL each wash).

The other MOFs in this study were loaded with 3-octanone or 4-methyl-3-heptanone using analogous procedures. In some cases, MOF samples were activated by chloroform exchange followed by heating under vacuum to remove DMF.

#### 4. Stability studies

Figure S3 shows PXRD patterns for IRMOF-NHPr following activation when left to stand in air. As can be seen from the reduction in peak intensities, crystallinity is lost over the course of 1 h. Figures S4 and S5 show PXRD patterns for IRMOF-NHPr following loading with excess 3-octanone and 4-methyl-3-heptanone, respectively. Loading with either pheromone has been shown to stabilise the MOF with respect to decomposition in air.



**Figure S3.** PXRD patterns for IRMOF-NHPr: simulated PXRD pattern from the single crystal X-ray diffraction data for IRMOF-3 (black),<sup>19</sup> PXRD pattern for IRMOF-NHPr upon activation (red), PXRD pattern for activated IRMOF-NHPr after 15 min in air (blue), PXRD pattern for activated IRMOF-NHPr after 30 min in air (green), PXRD pattern for activated IRMOF-NHPr after 1 h in air (pink).





(red), PXRD pattern for 3-octanone-loaded IRMOF-NHPr after 15 min in air (blue), PXRD pattern for 3octanone-loaded IRMOF-NHPr after 30 min in air (green), PXRD pattern for 3-octanone-loaded IRMOF-NHPr after 1 h in air (pink), PXRD pattern for 3-octanone-loaded IRMOF-NHPr after 2 days in air (brown), PXRD pattern for 3-octanone-loaded IRMOF-NHPr after 5 days in air (orange).



Figure S5. PXRD patterns for IRMOF-NHPr after loading with 4-methyl-3-heptanone (mhp): simulated PXRD pattern from the single crystal X-ray diffraction data for IRMOF-3 (black),<sup>19</sup> PXRD pattern for IRMOF-NHPr upon activation (red), PXRD pattern for mhp-loaded IRMOF-NHPr after 15 min in air (blue), PXRD pattern for mhp-loaded IRMOF-NHPr after 30 min in air (green), PXRD pattern for mhp-loaded IRMOF-NHPr after 1 h in air (pink), PXRD pattern for mhp-loaded IRMOF-NHPr after 2 days in air (brown), PXRD pattern for mhp-loaded IRMOF-NHPr after 5 days in air (orange).

#### 5. Release studies

Pheromone-loaded MOFs were placed in unsealed vials at room temperature and exposed to air. At regular intervals, small samples were taken, digested as described in Section 1 and analysed by <sup>1</sup>H NMR spectroscopy, with integrals used to calculate the pheromone remaining in the samples. Each analysis was carried out three times.

Headspace analysis was also undertaken using GC-MS over an eight-day period. The results are summarised in Figure S6, and clearly show evidence for 3-octanone above the solid samples.



Figure S6. Headspace analysis on a sample of 3-octanone-loaded IRMOF-NHPr, showing the mean area of the 3-octanone peak.

#### 6. Simulation Methods

All simulations were carried out at a loading of 0.156 molecules of 3-octanone molecules / formula unit corresponding to 1.25 molecules / unit cell. To improve statistics, the simulations where carried out in a simulation box comprising 2 × 2 × 2 unit cells. Molecular dynamics simulations were carried out in the NVT ensemble with flexible models for both the frameworks and the 3-octanone molecules using LAMMPS (Large-scale Atomic/Molecular Massively Parallel Simulator).<sup>S1,S2</sup> Cif files of the frameworks were converted to LAMMPS data files using a cif to LAMMPS interface.<sup>S3</sup> The DREIDING force field was employed to describe the framework<sup>S4</sup> and the GAFF force field for the 3-octanone molecules.<sup>S5,S6</sup> The exclusion rules of GAFF and DREIDING are different and in the present work the exclusion

rules of DREIDING were used. Partial charges for the MOFs and the 3-octanone molecules were calculated using the RESP (Restrained Electrostatic Potential) method<sup>S7</sup> implemented in CP2K.<sup>S8,S9</sup> A cut-off radius of 18 Å was used for non-bonded interactions (electrostatics and Lennard-Jones) and tail corrections were applied. Long range electrostatic interactions were computed using the particle-particle-particle-mesh (P<sup>3</sup>M) algorithm.<sup>S10</sup> The temperature was maintained at 300 K using the Nose Hoover thermostat.<sup>S11,S12</sup> A time step of 1 fs was employed. After 6 ns of equilibration, data were collected over a production run of 20 ns. Three replicas were run for each simulation.

#### 7. Behavioural Bioassays

The experiments were conducted in a *Eucalyptus* plantation, 20 km from Juiz de Fora, MG, Brazil using thirty small (1-2 foraging entrance holes) *Atta sexdens* colonies. The ant species necessitated use of a mhp-loaded MOF sample as opposed to an oct-loaded sample. A 500 mg sample of  $[Zn_4O(bdc-NHPr)_3]$  mhp was washed with hexane, air-dried to remove traces of DMF and then stored as 50 mg subsamples at  $-20^{\circ}$ C until use. As a positive control, 50 µl of 4-methyl-3-heptanone was applied to a 1 cm diameter rubber disk, left for 5 min to absorb at room temperature, and then stored at  $-20^{\circ}$ C until use. Treatments consisted of 10 g of citrus pulp bait contained sulfluramid insecticide (Mirex-S, Atta-Kill) placed immediately beside the main foraging trail of the colony, 10 cm from the entrance hole. The bait was either applied by itself as a control, or had either a rubber disk with 4-methyl-3-heptanone pheromone (positive control), or a 50 mg sample of  $[Zn_4O(bdc-NHPr)_3]$  mhp placed on top of the bait pile. The numbers of leaf-cutting ant workers approaching to within 2 cm of the bait piles were then counted for 15 min. The numbers of workers counted at the bait piles were compared between the three treatments using a Kruskal-Wallis test in SPSS 25.

# 8. References

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