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Fast and slow walking driven by chemical fuel[†]

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We demonstrate the fast forward and slow backward motion of a biped on a tetrahedral track using chemical fuel, cooperative binding and kinetic selectivity. Walking of the biped is based on its dibenzyl amine feet that bind to zinc porphyrin units and, upon protonation, to dibenzo 24-crown-8 sites affording pseudorotaxane linkages.

The myosin, dynein and kinesin family of motor proteins perform sophisticated, repetitive transport tasks essential to cellular functioning.¹ Importantly, these intracellular transport tasks are performed out-of-equilibrium² by the consumption of ATP³ as fuel. These complex, biological processes have instigated the design of both DNA-based⁴ as well as synthetic/semi-synthetic small-molecule walkers,^{5–8} often driven by chemical, light or biological inputs.

Molecular walkers have also been applied as transducers for catalytic reactions⁹ or cargo transport.^{4,10} However, most synthetic walkers rely on either strong covalent or kinetically inert coordination bonds, which firmly hold the biped on track. The desired sequential multi-step walking is then realized *via* chemical/light/semi-biological stimulation.^{11–13} In contrast, the concept of a multi-component walker held on track by dynamic interactions remains a largely unexplored area,^{14,15} and even more, if the system is chemically fueled.

In this report, we demonstrate how a multi-component assembly walks across a deck by exploiting the reversible switching between the $NH_{amine} \rightarrow zinc$ -porphyrin¹⁶ (= $NH_{amine} \rightarrow ZnPor$) and dibenzylammonium pseudorotaxane binding¹⁷ motifs. The process may be started with acid and reversed with base, or in a pulsed forward-backward motion using a fuel acid,¹⁸ for instance 3¹⁹ (Fig. 1). Upon protonation of [1•2] the dibenzylamine sites are

protonated and insert as secondary ammonium units into the dibenzo-24-crown-8 binding sites of deck 1 affording two [2]pseudorotaxane linkages in $[1 \cdot 2(H^{+})_{2}]$. With base the walking is reversed.

The design strategy for the walker-on-deck was based on recent reports from our group, where we demonstrated that $N_{\rm Pv} \rightarrow {\rm ZnPor}$ and pseudorotaxane linkages are orthogonal interactions.²⁰ The synthesis and characterization of deck 1 and other ligands used in this study (Fig. 1) are described in detail in the ESI.[†] Upon mixing deck 1 and biped 2 (1:1), the assembly [1•2] was quantitatively afforded. The walker-on-deck was fully characterized by ¹H, ¹H-¹H COSY and elemental analysis (ESI,† Fig. S21, S22 and S29). Further, the ¹H DOSY of [1•2] vielded a single, sharp trace with a $D = 4.40 \times 10^{-10} \text{ m}^2 \text{ s}^{-1}$ confirming the quantitative complexation between 1 and 2 (ESI,† Fig. S29). Foremost, [1•2] was identified by the changes in the ¹H-NMR signals of protons j'-H and g'-H of biped 2 as walker and those of proton r-H of deck 1 (Fig. 2c-e). The stark changes of the j'-H and g'-H signals were attributed to interaction with the porphyrin's shielding ring current upon axial $NH_{amine} \rightarrow ZnPor$ coordination, which consequently shifted the r-H proton signals of 1 upfield.

Similarly, **1** and **2** were mixed with TFA in a 1:1:2 ratio to assemble $[1 \cdot 2(H^+)_2]$. The double protonation of **2**, however, in parallel removed Zn^{2+} (*ca.* 50%) from the ZnPor sites of deck **1**. Instead, we achieved [2]pseudorotaxane formation with HPF₆ as acid by first generating $[2(H^+)_2](PF_6^-)_2$, which in a second step was combined with deck **1** (Fig. 2a and b).

Complex $[1 \cdot 2(H^+)_2](PF_6^-)_2$ was fully characterized by ¹H, ¹H-¹H COSY NMR and elemental analysis. Additionally, the ¹H DOSY of $[1 \cdot 2(H^+)_2](PF_6^-)_2$ yielded a single, sharp trace with $D = 3.30 \times 10^{-10} \text{ m}^2 \text{ s}^{-1}$ confirming quantitative complexation between 1 and $[2(H^+)_2](PF_6^-)_2$ (ESI,† Fig. S28). Most diagnostic were the changes in the ¹H-NMR signature of proton signals o-H, p-H of 1 and h'-H, i'-H adjacent to the ammonium units of $[1 \cdot 2(H^+)_2]$. The proton signals of o-H & p-H split into a complex multiplet due to the de-symmetrization induced by threading.

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Fig. 1 Ligands used in this study. (The cartoon version of the tetrahedral deck 1 is a 2D-projection).



Fig. 2 Partial comparison of ¹H-NMR (500 MHz, CD₂Cl₂, 298 K) of (a) free biped $[2(H^+)_2](PF_6^-)_2$, (b) [2]pseudorotaxane $[1 \bullet 2(H^+)_2](PF_6^-)_2$, (c) free deck 1, (d) $[1 \bullet 2]$, and (e) free biped 2.

The proton signals h'-H & i'-H of $[1 \bullet 2(H^+)_2]$ (4.27–4.23 ppm in $[2(H^+)_2](PF_6^-)_2)$ shifted downfield to 4.74 ppm (broad) in $[1 \bullet 2(H^+)_2]$ after [2]pseudorotaxane formation (Fig. 2a and b).

Before studying the dissipative system, we tested the stepwise deprotonation-driven walking using DBU as base (Scheme 1).



Scheme 1 Stepwise walking of biped 2 on deck 1 by step-by-step deprotonation of the bis-pseudorotaxane $[1{-}2(H^+)_2](PF_6^-)_2$ by titration with DBU. The intermediately formed $[1{-}2(H^+)](PF_6^-)$ then undergoes gradual deprotonation to the di- $NH_{Amine} \rightarrow$ ZnPor bound complex $[1{-}2]$.

Pseudorotaxane $[1 \bullet 2(H^+)_2](PF_6^-)_2$ was titrated with DBU (increments of 0.1 equiv.) using the proton r-H signal as an indicator (Fig. 3). Upon addition of DBU, the proton r-H signal split into three sets: two sets from $[1 \cdot 2(H^+)]$ with two distinct ZnPor units (unbound: 10.40 ppm; bound: 10.36 ppm) and one set from $[1 \circ 2(H^{\dagger})_2]$ with two degenerate ZnPor sites (10.41 ppm). From 0.1-2.0 equiv. of DBU, the integration of the r-H signal at 10.36 ppm (w.r.t. to an internal standard) furnished the amount of $[1 \bullet 2(H^+)]$ in the mixture of $[1 \bullet 2(H^+)_2]$, $[1 \bullet 2(H^+)]$, and $[1 \bullet 2]$ as shown in Fig. 3a. Further, a plot of the percentage of $[1 \bullet 2(H^+)]$ vs. the equiv. of added DBU revealed some interesting findings: First, there is a steady growth (from 0 to 86%) of $[1 \bullet 2(H^+)]$ during the addition of 0.1-1.2 equiv. of DBU followed by its decay to form [1•2]. Secondly, the plot clearly shows that at least 86% (peak of the inverted parabolic-fit) of the back walking of $[1 \bullet 2(H^+)_2] \rightarrow [1 \bullet 2]$ occurs stepwise by mono-deprotonation. After adding 1.0 equiv. of DBU (ESI,[†] Fig. S31), the mixture is composed of $[1 \bullet 2(H^+)]$ (80%) and $\left[1 \bullet 2(H^{+})_{2}\right]$ (20%). After 1.4–2.0 equiv. of DBU had been added, the diagnostic j', g'-H proton signals of [1•2] around 6.7 ppm appeared, adding further support to the stepwise deprotonation pathway.

One would expect the monoprotonated $[1 \cdot 2(H^+)]$ to undergo sliding of the amine foot between both zinc porphyrin sites and/or the ammonium foot between the two degenerate dibenzo24-crown-8 units. Surprisingly, a ¹H-¹H ROESY-NMR using the o,p-H and r-H as diagnostic signals revealed exchange peaks neither between the two crown ethers nor between both



Fig. 3 (a) Partial ¹H-NMR stack showing the changes in the proton r-H signal upon titration of $[1 \cdot 2(H^+)_2](PF_6^-)_2$ with DBU. (b) Percentage of $[1 \cdot 2(H^+)](PF_6^-)$ in the mixture against equiv. of DBU added w.r.t to the initial assembly $[1 \cdot 2(H^+)_2](PF_6^-)_2$.



Scheme 2 Chemically fueled switching between $[1 \bullet 4]$ and the bis-[2]pseudorotaxane $[1 \bullet 4(H^+)_2]$ using the fuel acid 3.

zinc porphyrin units (ESI,[†] Figure S27). Accordingly, the amine foot did not exchange between the two zinc porphyrins and by extension there was no chemical exchange between the two distinct r-H signals in $[1 \cdot 2(H^+)]$. Thus, the biped takes a steady walking motion from $[1 \cdot 2(H^+)_2]$ through $[1 \cdot 2(H^+)]$ to $[1 \cdot 2]$.

With these findings at hand, we decided to study the operation of dissipative walking using fuel 3. Unexpectedly, ligand 2 was not suitable for this purpose as $[2(H^+)_2]$ proved to be insoluble. To remedy this problem, ligand 4 was designed as an analog of 2 with bis-octoxy chains for higher solubility. Consequently, now ligands 1, 4 were mixed to quantitatively furnish [1•4] which was fully characterized by ¹H, ¹H-¹H COSY, ¹H DOSY ($D = 3.50 \times 10^{-10} \text{ m}^2 \text{ s}^{-1}$) and elemental analysis. (ESI,† Fig. S25, S26 and S30) Subsequently, 2.5 equiv. of fuel acid 3 was added for conversion of $[1 \bullet 4]$ to $[1 \bullet 4(H^{+})_{2}]$ (Scheme 2). ¹H-NMR indicated the disappearance of the drastically upfield shifted broad signals of protons j'-H, g'-H in [1•4] along with a shift of the r-H proton from 10.37 to 10.39 ppm as in the free deck 1 (Fig. 4). Proton signals j',g'-H shifted from 6.65 and 6.58 ppm in [1•4] to 7.42 and 7.28 ppm in $[1•4(H^+)_2]$, respectively. Moreover, proton signals o,p-H of deck 1 showed an intricate splitting into two sets similar to those in $[1 \cdot 2(H^+)_2]$, which supported formation of bis-[2]pseudorotaxane $[1 \bullet 4(H^+)_2]$.

The stepwise walking from $[1 \bullet 4(H^+)_2]$ back to $[1 \bullet 4]$ *via* stepby-step deprotonation was also monitored by ¹H-NMR. The dithreaded bis-[2]pseudorotaxane $[1 \bullet 4(H^+)_2]$ prevailed in the mixture from *t* = 8 to 80 min as evidenced by the single r-H signal at 10.39 ppm along with the split o,p-H signals (Fig. 4: Phase 1). From *t* = 80 to 125 min we clearly see the emergence of two distinct sets of r-H signals (10.39 and 10.38 ppm) also accompanied by



Fig. 4 Comparison of partial ¹H-NMR spectra as a function of time (500 MHz, CD₂Cl₂, 298 K) during the first fueled motion cycle.

shifts of the β_2 , β_3 -H signals in the range of 9.4–9.6 ppm (Phase 2 in Fig. 4). The r-H signals at 10.39 and 10.38 ppm correspond to the di-threaded $[1 \cdot 4(H^+)_2]$ and mono-threaded $[1 \cdot 4(H^+)]$, respectively. At time t = 115-130 min, the full conversion of $[1 \cdot 4(H^+)_2]$ to the monoprotonated $[1 \cdot 4(H^+)]$ was finished, as indicated by nearly complete disappearance of $[1 \cdot 4(H^+)_2]$ (10.39 ppm) and simultaneous increase in the signal corresponding to $[1 \cdot 4(H^+)]$ (10.38 ppm). Alike, the β -protons of $[1 \cdot 4(H^+)_2]$ disappeared while the signals from $[1 \cdot 4(H^+)]$ simultaneously increased showing the same stepwise walking protocol.

From t = 130 to 230 min, we then observe the complete conversion of the mixture into [1•4] (Phase 3 in Fig. 4) due to the second deprotonation. The gradual emergence of the h', i'-H signals (2.55-2.65 ppm) (Fig. S33, ESI⁺) and g', j'-H signals (6.64-6.71 ppm) (Phase 3 in Fig. 4) are attributed to the reformation of [1•4]. Although, fuel 3 has been utilized to drive several dissipative networks²¹⁻²³ and a standalone molecular rotor,²⁴ this report represents the first use of 3 to drive a walkeron-deck. The reproducibility of the walking process was tested twice by running two consecutive fueled walking cycles with minimal disruption by the waste products from the chemical fuel consumption apart from a slightly longer 2nd cycle operation time (362 min vs. 230 min) (Fig. 5a and b). From the data, we recognize that walking from $[1 \bullet 4]$ to $[1 \bullet 4(H^{+})_{2}]$ is fast, while the return *via* the intermediate $[1 \bullet 4(H^+)]$ is slow, as if a brake pad²⁵ would apply. For functional utility in the future this opens the opportunity to take up a load in the long-lived $[1 \bullet 4(H^+)_2]$ and carry it to [1•4].

In summary, we have demonstrated unprecedented reversible and stepwise walking of a small biped on a tetrahedral track using a fuel acid. Further, the three-step walking motion is based on a novel, pulsed switching protocol between orthogonally self-sorted $NH_{\text{Amine}} \rightarrow \text{ZnPor}$ and ammonium \subset crownether pseudorotaxane interactions. The combination of multiple self-sorting^{26,27} and fueling emphasizes the potential of leveraging dissipative processes^{28–30} to achieve sophisticated functions in multi-component, dynamic supramolecular assemblies.

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Fig. 5 (a) Plot of chemical shift of r-H of **1** *versus* time during the first cycle from $[1\bullet4]$ to $[1\bullet4(H^+)_2]$ and back. The three phases in the transition (Phases 1–3) are populated by different proportions of the assemblies. (b) Plot of chemical shift of proton r-H of **1** *versus* time over two fueled cycles monitored by ¹H-NMR. The biped **4** was twice driven from $[1\bullet4]$ to $[1\bullet4(H^+)_2]$ and automatically back to $[1\bullet4]$ with minimal degradation in function.

Conflicts of interest

There are no conflicts to declare.

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