LETTERS 2003 Vol. 5, No. 21 3951-3954

ORGANIC

Conformational and Electronic Engineering of Twisted Diphenylacetylenes

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Received August 13, 2003

ABSTRACT



Three tethered diphenylacetylene derivatives were prepared by alkyne metathesis. In these cycles, the twist angle between the two benzene rings is variable and determined by the nature of the linker. The engineering of the twist angle leads to a change of the UV–vis spectra of the cycles. The larger the twist angle in the macrocycles, the more blue shifted their λ_{max} (UV–vis), the lower their fluorescence quantum yield, and the lower field shifted their ¹³C NMR signals of the alkyne carbons are.

This contribution presents the synthesis of diphenylacetylene derivatives **12**, **13**, and **15** by ring-closing alkyne metathesis. The optical properties of **12**, **13**, and **15** are dependent upon the size of the twist angle α . Increasing the angle α leads to larger HOMO–LUMO gaps and lower emissive efficiencies.

Diphenylacetylenes (DPA) are fascinating in their own right and as building blocks for larger structures, oligo- and poly(phenyleneethynylene)s. DPAs are rigid, and their only degree of freedom is the rotation of the arene units around the central alkyne. This rotational freedom has an impact upon the optical and electronic properties of DPAs 1-6 as well upon that of the poly(*p*-phenyleneethynylene)s (PPEs).¹⁻⁶

The conformational change has been invoked to explain the large solvatochromic and thermochromic effects that the PPEs show and the change in fluorescence observed between the pair **1a**/2 and **2**/6 upon binding to metal cations.¹ While the PPEs do show large effects in absorption, the DPAs seem to show only a change in emission intensity, but not in their absorption spectra. We found that curious and were interested to "dial in" the twist angle α between the two arene rings in DPAs to examine their electronic properties. There was only sparse information regarding how to change the twist angle in DPAs.⁷

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A series of bridged DPAs was checked by us via fast AM1 calculations. Xylylene linkers between the diphenylacetylenes led to structures 12-14 with varying α when going from 12 to 14. Ab initio calculations of these topologies (RHF 6-31G**) refined the picture and showed that 13 is the least twisted cycle ($\alpha = 40^{\circ}$) and that both 12 and 14 are considerably more distorted (Figure 1). While these calculations



Figure 1. Computationally obtained (RHF 6-31G**, Spartan 2002 on a Windows platform) geometries for **12** (left), **13** (middle), and **14** (right). The α values are 58.82°, 39.12°, and 88.02°.

tions give gas phase structures, the 6-31G** basis set is generally reliable for ground-state geometries. Noticeable was the significant shift in the HOMO and LUMO energies upon the twisting action in the three isomeric cycles (Figure 1). While the HOMO-LUMO values do not directly correspond to a physical property they qualitatively match changes in the absorption spectra. An increasing angle α led to an increased HOMO-LUMO gap according to the calculations. Because these cycles are quite flexible, we performed an energy profile calculation with respect to the rotation around the CC triple bond. In 13 and 14, only one minimum is observed (see Figure 1), but the conformational behavior of 12 is more complex. Two energy minima, one at 0° and one at approximately 60°, are found (Figure 2a). According to different levels of theory either one is somewhat more stable. AM1 and RHF 6-31G** predict the twisted form of 12 to be slightly more stable while B3LYP 6-31G** predicts the planar form of 12 to be more stable. The two conformers are separated by a barrier that was calculated to be between 4 and 7 kcal mol^{-1} . As a comparison, the rotational barrier of 1b was calculated (Figure 2b). All levels of theory show that the planar syn form in which both methoxy groups are



Figure 2. (a) Rotational profile of **12** calculated on three levels of theory. (b) Rotational profile of **1b** calculated on three levels of theory. (c) Rotational profile of diphenylacetylene calculated on three levels of theory. In a-c, the geometry optimization was performed on the AM1 level and the ab initio calculations were single point calculations utilizing optimized AM1 geometries. Abbreviations: HF, restricted Hartree–Fock; BLP, Becke–Lee– Yang–Parr functional (B3LYP). *x*-axis: torsion angle (deg).

located at the same side of the molecule is 0.7-0.8 kcal mol⁻¹ less stable than the planar anti form, where both methoxy groups avoid each other. The difference between the calculations is in the height of the barrier, which is nonexistent in the AM1 calculation and approximately 1 kcal mol⁻¹ in the DFT calculation.

All of the calculations were single-point energy calculations performed on optimized AM1 geometries. To asses the precision of these calculations, the rotational barrier of diphenylacetylene was computationally evaluated. This system had been calculated⁵ earlier in its planar as well in its 90° twisted conformation. The barrier of rotation was calculated to 0.86 kcal mol⁻¹.⁵ When utilizing AM1 optimized geometries and the B3LYP 6-31G** method this barrier is well reproduced with a value of 0.97 kcal mol⁻¹. The relative error utilizing AM1 geometries in combination with single-point ab initio calculations is <0.1 kcal mol⁻¹. The use of AM1-calculated geometries is therefore reasonable. The control calculation backs up the calculational results obtained for **1b** and **12–14**.

The synthesis of 12, 13, and 15 starts (Scheme 1) with the benzylation of 7 followed by propynylation utilizing a



Pd-catalyzed coupling of the Sonogashira type.⁸ The bisethers 8-11 are isolated in 28-56% yield. Alkyne metathesis with either the Grela⁹ system or our own preactivated variant¹⁰ of the Mortreux¹¹ catalyst furnished the cycles **12**, **13**, and **15** in 18-19% isolated yield after repeated chromatography on silica gel. Major byproducts are scarcely soluble oligomers and polymers that could not be purified/ separated. Repeated attempts to close **10** to **14** under different conditions failed with a variety of catalytic in situ systems.

To gain an understanding of the structures of the cycles, we attempted to grow single-crystalline specimens. Cycle 13 is a colorless glass that resisted single-crystal growth. The ortho cycles 12 and 15 formed single crystals from hexafluorobenzene (12) and hexane/dichloromethane (15), respectively. The use of hexafluorobenzene as a crystallization solvent seems critical because it favors aromatic faceto-face interactions.¹²⁻¹⁴ The structures of **12** and **15** are shown in Figure 3a,b. There are three independent molecules of 12 in the unit cell. Each of them has a different angle α $(22.4-26.6^{\circ})$. The energy potential for rotation is quite soft for 12, but that is not too unexpected. In the case of 15, only one twist angle (29.9°) is observed. The twist angle of 12 and 15 in the solid state is significantly smaller than calculated, but this is a packing effect in combination with the soft rotational profile in 12 and 15. While the X-ray data are an important proof of the topology of the cycles, and show a fascinating solid state ordering, they probably do not correctly represent the conformation of the cycles in solution (Figure 3).



Figure 3. (a) ORTEP representation of 12; $\alpha = 26.6^{\circ}$. (b) Ball and stick representation of 15; $\alpha = 29.9^{\circ}$.

The UV-vis spectra of 12, 13, 15, and 1b are shown in Figure 4. While 1b has its λ_{max} at 332 nm, the *ortho* cycles 12 and 15 show their λ_{max} at 313-315 nm, almost 20 nm blue-shifted. All of these spectra display a fine structure. The *meta* cycle 13 features a broad and undistinguished UV-vis spectrum that is intermediate between that of 1b and the

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Figure 4. (a) UV-vis spectra (chloroform) of 1b and the cycles 12 and 13. (b) Emission spectra (chloroform, height normalized) of 1b and the cycles 12 and 13. Emission of 15 is superimposable to that of 12 and therefore not shown.

other cycles. The fine structure in the spectra of **12** and **13** cannot be assigned to any specific vibration(s), and the numeric value of the fine structure in **1b** (2059 cm⁻¹) might be assigned to a CC triple-bond stretch. The fluorescence spectra of **1b** and **12**, **13**, and **15** are similar but their quantum yields are significantly different (Table 1). The fluorescence quantum yield (chloroform) of **13** is 69%, and **12** and **15** feature quantum yields in the range of 10-22%. The parent

Table 1. Spectroscopic Data of Cycles 12, 13, and 15

	1b	12	13	15
UV-vis (CHCl ₃ , nm)	330	313	309	315
ϵ at λ_{\max}		20 000	17 300	21 700
fluorescence (nm)	348	352	350	350
quantum yield (CHCl ₃)	0.45	0.12	0.69	0.10
CH ₂ Cl ₂	0.75	0.33	0.77	0.21
hexanes	0.54	0.16	0.66	0.08
ethyl ether	0.55	0.15	0.69	0.07
¹³ C NMR alkyne shift,	0	+1.20	+0.71	+1.64
ightarrow ppm (1b) ^{<i>a</i>}				

^{*a*} For the influence of mechanical strain on the ¹³CNMR spectral shifts of alkynes, see: Lee, D. C.; Sahoo, S. K.; Cholli, A. L.; Sandman, D. J. *Macromolecules* **2002**, *35*, 4347–4355. Cholli, A. L.; Sandman, D. J.; Maas, W. *Macromolecules* **1999**, *32*, 4444–4446.

compound **1b** has a fluorescence quantum yield of 45%. The quantum yields differ considerably upon the solvent utilized, but the trend is unchanged by the choice of solvents. Coinciding with the quantum yields and the blue shift in the absorption are the ¹³C NMR chemical shifts of the alkyne carbons that deshield in the same way, reaching a maximum of 1.64 ppm in **15**. Again, the meta cycle **13** is the one that resembles **1b** most.

In conclusion, we have shown that the optical gap and the fluorescence quantum yield in DPAs **1b**, **12**, **13**, and **15** vary in dependence upon the twist angle of the two constituting phenyl rings. A self-consistent picture emerges from experimental and calculational results, in which an increased torsion angle leads to a decreased fluorescence quantum yield *and an enlarged optical gap*.

Acknowledgment. We thank the National Science Foundation (CHE 0138659) and the USC NanoCenter for generous support. U.B. is a Camille Dreyfus Teacher–Scholar (2000–2004).

Supporting Information Available: Experimental procedures and characterization for all compounds and CIFs for **12** and **15**. This material is available free of charge via the Internet at http://pubs.acs.org.

OL035532+