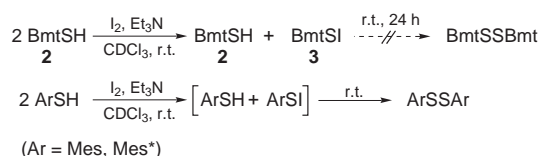


Scheme 2 ^a Estimated by ¹H NMR

amount of disulfide **4** (Scheme 2). The reaction of **3** with benzylamine (10 equiv.) readily afforded sulfenamide **5**. On the other hand, **3** was found to be unreactive toward thiol **2** bearing the same substituent. Oxidation of **2** with 0.5 equiv. of I₂ in the presence of triethylamine afforded a 1 : 1 mixture of **2** and **3**, no disulfide formation being detected even after 24 h at room temperature (Scheme 3). These results imply that two species which are otherwise incompatible can be present in the same system with retention of their reactivities toward other reagents when they bear a bowl-type substituent. Under the same conditions, 2,4,6-trimethylbenzenethiol (MesSH) was immediately oxidized to MesSSMes and even 2,4,6-tri-*tert*-butylbenzenethiol (Mes*SH) afforded Mes*SSMes* although the reaction was much slower. Apparently, the bowl-shaped structure of the Bmt group is more effective for prevention of dimerization than a Mes* group, where the functionality is more closely shielded.



Scheme 3 Mes* = C₆H₂But₃-2,4,6

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Notes and References

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§ Bmt denotes 4-*tert*-butyl-2,6-bis[(2,2'',6,6''-tetramethyl-*m*-terphenyl-2'-yl)methyl]phenyl.

¶ **3**: mp 252–257 °C; ¹H NMR (500 MHz, CDCl₃): δ 0.97 (s, 9H), 1.95 (brs, 24H), 3.70 (brs, 4H), 6.46 (s, 2H), 6.84 (brs, 8H), 6.94 (t, ³J = 7.3 Hz, 4H), 7.03 (d, ³J = 7.5 Hz, 4H), 7.33 (t, ³J = 7.5 Hz, 2H); ¹³C NMR (125 MHz, CDCl₃): δ 21.0 (q), 30.8 (q), 34.7 (s), 34.8 (t), 125.0 (d), 126.8 (d), 127.0 (d), 127.5 (d), 129.3 (d), 129.4 (s), 136.2 (s), 137.2 (s), 140.8 (s), 142.1 (s), 146.2 (s), 151.1 (s); UV–VIS (CHCl₃): λ_{max}(ε) = 328 nm (4000 dm³ mol^{−1} cm^{−1}).

|| Crystal data for **3**·0.5C₇H₈: C_{59.5}H₆₁SI, *M* = 935.10, monoclinic, space group *P*2₁/*n*, *a* = 15.668(8), *b* = 17.47(1), *c* = 18.71(1) Å, β = 105.91(4)°, *U* = 4926(4) Å³, *Z* = 4, *D*_c = 1.261 g cm^{−3}, μ = 7.30 cm^{−1}. The intensity data were collected at 293 K on a Rigaku AFC7R diffractometer with Mo-Kα radiation (λ = 0.71069 Å), and the structure was solved by direct methods and expanded using Fourier techniques. The non-hydrogen atoms were refined anisotropically. Hydrogen atoms were included but not refined. The final cycle of full-matrix least-squares refinement was based on 3789 observed reflections [*I* > 3.00σ(*I*)] and 539 variable parameters with *R*(*R*_w) = 0.058(0.045). CCDC 182/944.

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