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Exploring the boundaries of a light-driven molecular motor design

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_journal_name_full Org.Biomol.Chem.

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1. SUBMISSION DETAILS

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0. AUDIT DETAILS

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_audit_creation_method 'PLATON <TABLE ACC> option'
_audit_update_record
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_publ_contact_author_name # Name of author for correspondence
'Prof Ben Feringa'
_publ_contact_author_address # Address of author for correspondence
;

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9747 AG
NETHERLANDS
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;
Exploring the boundaries of a light-driven
molecular motor design: new sterically overcrowded alkenes with
preferred direction of rotation
;

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Altomare, A., Burla, M.C., Camalli, M., Cascarano, G.L.,
(1999) J. Appl. Cryst. 32, 115-119.

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Dordrecht, The Netherlands.

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;

Fig. 1. Chemical structural diagram (scheme 1) of the title compound

Fig. 2. Perspective PLUTO drawing of the molecule illustrating the
configuration and the adopted numbering scheme.

Fig. 3. Molecular packing viewed down unit cell axes.

Fig. 4. Perspective ORTEP drawing of the title compound.
Displacement ellipsoids for non-H are represented at the 50%
probability level.

The H-atoms are drawn with an arbitrary radius.

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loop_

_publ_author_name

'Ben Feringa'

'Harmen de Jong'

'Matthijs K.J. ter Wiel'

'Richard van Delden'

#####

data_c27h20s2

_database_code_depnum_ccdc_archive 'CCDC 231703'

#####

5. CHEMICAL DATA

_chemical_name_systematic

;

?

;

_chemical_name_common ?

_chemical_melting_point ?

_chemical_formula_moiety 'C27 H20 S2'

_chemical_formula_structural ?

Ex: 'C12 H16 N2 O6, H2 O' and '(Cd 2+)3, (C6 N6 Cr 3-)2, 2(H2 O)'

_chemical_formula_sum 'C27 H20 S2'

_chemical_formula_weight 408.59

_chemical_compound_source 'see text'

loop_

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_atom_type_description

_atom_type_scatter_dispersion_real

_atom_type_scatter_dispersion_imag

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H H 0.0000 0.0000 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
C C 0.0033 0.0016 'International Tables Vol C Tables 4.2.6.8 and 6.1.1.4'
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# 6. CRYSTAL DATA
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_symmetry_space_group_name_Hall '-P 2yn'
_symmetry_space_group_name_H-M 'P 21/n'
```

```
loop_
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```
_symmetry_equiv_pos_as_xyz
```

```
x,y,z
```

```
1/2-x,1/2+y,1/2-z
```

```
-x,-y,-z
```

```
1/2+x,1/2-y,1/2+z
```

```
_cell_length_a      8.827(2)
_cell_length_b      21.647(4)
_cell_length_c      11.075(2)
_cell_angle_alpha    90
_cell_angle_beta     100.493(3)
_cell_angle_gamma    90
_cell_volume         2080.8(7)
_cell_formula_units_Z 4
```

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_cell_measurement_temperature 293(1)
_cell_measurement_reflns_used  3334
_cell_measurement_theta_min    2.53
_cell_measurement_theta_max    21.79
_cell_special_details
```

```
;
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The final unit cell was obtained from the xyz centroids of
3334 reflections after integration using the SAINT software
package (Bruker, 2000).
```

```
;
```

```
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_exptl_crystal_colour           colorless
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_exptl_crystal_size_mid         0.15
_exptl_crystal_size_min         0.08
_exptl_crystal_size_rad         ?
_exptl_crystal_density_meas     ?
_exptl_crystal_density_diffn    1.304
_exptl_crystal_density_method   'not measured'
_exptl_crystal_F_000            856
_exptl_absorpt_coefficient_mu    0.267
_exptl_absorpt_correction_type   Multi-Scan
_exptl_absorpt_process_details  '(SADABS, Sheldrick, Bruker, 2000)'
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```
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# 7. EXPERIMENTAL DATA
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_exptl_special_details
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```
;
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```
?
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;
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_diffrn_radiation_monochromator  'parallel mounted graphite'
_diffrn_radiation_detector
;
CCD area-detector
;
_diffrn_measurement_device_type
;
Bruker Smart Apex
;
_diffrn_measurement_method       'phi and omega scans'
_diffrn_special_details          ?
_diffrn_detector_area_resol_mean '4096x4096 / 62x62 (binned 512)'
_diffrn_standards_number         ?
_diffrn_standards_interval_count ?
_diffrn_standards_interval_time ?

loop_
_diffrn_standard_refl_index_h
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_diffrn_standard_refl_index_l
? ? ?

# number of measured reflections (redundant set)
_diffrn_reflns_number            14823
_diffrn_reflns_av_R_equivalents  0.0379
_diffrn_reflns_av_sigmaI/netI    0.0446
_diffrn_reflns_limit_h_min       -10
_diffrn_reflns_limit_h_max       10
_diffrn_reflns_limit_k_min       -25
_diffrn_reflns_limit_k_max       25
_diffrn_reflns_limit_l_min       -13
_diffrn_reflns_limit_l_max       12
_diffrn_reflns_theta_min         2.53
_diffrn_reflns_theta_max         25.02
_diffrn_measured_fraction_theta_max 0.999
_diffrn_reflns_theta_full        25.02
_diffrn_measured_fraction_theta_full 0.999

_diffrn_reflns_reduction_process
;
Intensity data were corrected for Lorentz and polarization
effects, decay and absorption and reduced to  $F_o \sim F^2$ 
using SAINT (Bruker, 2000) and SADABS (Sheldrick, 2001)
;
# number of unique reflections
_reflns_number_total             3673
_reflns_number_gt                2408
_reflns_threshold_expression     >2sigma(I)

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_computing_cell_refinement       'SAINT, Bruker Version 6.02A, 2000'
_computing_data_reduction        'XPREP, Bruker Version 5.1/NT, 2000'
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SIR-97 (Altomare et al., 1997)
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_computing_structure_refinement  'SHELXL-97 (Sheldrick, 1997)'
_computing_molecular_graphics
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PLUTO (Meetsma, 2001)
PLATON (Spek, 1994)

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PLATON (Spek, 1990)
SHELXL (Sheldrick, 1997)
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#=====

# 8. REFINEMENT DATA

_refine_special_details
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Refinement of F^2 against ALL reflections. The weighted R-factor wR and
goodness of fit S are based on F^2, conventional R-factors R are based
on F, with F set to zero for negative F^2. The threshold expression of
F^2 > 2sigma(F^2) is used only for calculating R-factors(gt) etc. and is
not relevant to the choice of reflections for refinement. R-factors based
on F^2 are statistically about twice as large as those based on F, and R-
factors based on ALL data will be even larger.
;

_refine_ls_structure_factor_coef Fsqd
_refine_ls_matrix_type full
_refine_ls_weighting_scheme calc
_refine_ls_weighting_details
'calc w=1/[\s^2*(Fo^2)+(0.0598P)^2+0.4521P] where P=(Fo^2+2Fc^2)/3'
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_atom_sites_solution_secondary direct
_atom_sites_solution_hydrogens difmap
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_refine_ls_extinction_method none
_refine_ls_extinction_coef ?
_refine_ls_abs_structure_details ?
_refine_ls_abs_structure_Flack ?
_refine_ls_number_reflns 3673
_refine_ls_number_parameters 331
_refine_ls_number_restraints 0
_refine_ls_number_constraints ?
_refine_ls_R_factor_all 0.0834
_refine_ls_R_factor_gt 0.0498
_refine_ls_wR_factor_ref 0.1342
_refine_ls_wR_factor_gt 0.1179
_refine_ls_goodness_of_fit_ref 1.022
_refine_ls_restrained_S_all 1.022
_refine_ls_shift/su_max 0.000
_refine_ls_shift/su_mean 0.000

_refine_diff_density_max 0.342
_refine_diff_density_min -0.302
_refine_diff_density_rms 0.037

_vrn_publ_code_void_volume 0.0
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# 9. ATOMIC COORDINATES AND DISPLACEMENT PARAMETERS

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C2 C Uani 0.0279(4) 0.3816(2) -0.0955(3) 1.000 0.0887(14) . . .
C3 C Uani 0.0987(5) 0.4355(2) -0.0668(3) 1.000 0.0926(16) . . .
C4 C Uani 0.1717(4) 0.44998(15) 0.0551(3) 1.000 0.0763(12) . . .
C5 C Uani 0.2578(5) 0.50398(19) 0.0842(5) 1.000 0.0998(19) . . .
C6 C Uani 0.3316(5) 0.51568(18) 0.2001(5) 1.000 0.0977(18) . . .
C7 C Uani 0.3224(4) 0.47323(17) 0.2943(4) 1.000 0.0827(16) . . .
C8 C Uani 0.2386(3) 0.42051(15) 0.2700(3) 1.000 0.0651(11) . . .
C9 C Uani 0.1605(3) 0.40705(12) 0.1499(2) 1.000 0.0592(9) . . .
C10 C Uani 0.0759(3) 0.35128(13) 0.1191(2) 1.000 0.0564(9) . . .
C11 C Uani 0.0613(3) 0.30195(11) 0.2093(2) 1.000 0.0509(8) . . .
C12 C Uani 0.1211(3) 0.24143(14) 0.1709(3) 1.000 0.0605(10) . . .
C13 C Uani 0.0067(4) 0.21327(16) 0.0646(3) 1.000 0.0872(12) . . .
C14 C Uani 0.0730(5) 0.1597(2) 0.0078(4) 1.000 0.1394(19) . . .
C15 C Uani -0.0046(3) 0.30976(11) 0.3090(2) 1.000 0.0515(8) . . .
C16 C Uani -0.0097(3) 0.26206(13) 0.4038(3) 1.000 0.0606(10) . . .
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C18 C Uani -0.0560(4) 0.1593(2) 0.4730(6) 1.000 0.0979(18) . . .
C19 C Uani -0.0052(5) 0.1766(3) 0.5921(5) 1.000 0.110(2) . . .
C20 C Uani 0.0388(4) 0.2358(3) 0.6206(4) 1.000 0.0912(18) . . .
C21 C Uani 0.0337(3) 0.27903(14) 0.5278(3) 1.000 0.0692(10) . . .
C22 C Uani -0.0502(3) 0.39281(13) 0.4542(2) 1.000 0.0616(10) . . .
C23 C Uani -0.1252(4) 0.44605(15) 0.4818(3) 1.000 0.0760(12) . . .
C24 C Uani -0.2277(4) 0.47493(16) 0.3924(4) 1.000 0.0811(14) . . .
C25 C Uani -0.2589(4) 0.45126(15) 0.2751(3) 1.000 0.0728(11) . . .
C26 C Uani -0.1859(3) 0.39850(13) 0.2468(3) 1.000 0.0610(10) . . .
C27 C Uani -0.0783(3) 0.36848(12) 0.3348(2) 1.000 0.0539(8) . . .
H2 H Uiso -0.023(4) 0.3676(15) -0.187(3) 1.000 0.112(11) . . .
H3 H Uiso 0.105(4) 0.4640(16) -0.123(3) 1.000 0.112(13) . . .
H5 H Uiso 0.258(4) 0.5342(17) 0.019(4) 1.000 0.123(13) . . .
H6 H Uiso 0.382(4) 0.5526(16) 0.215(3) 1.000 0.099(12) . . .
H7 H Uiso 0.373(3) 0.4777(13) 0.372(3) 1.000 0.074(10) . . .
H8 H Uiso 0.233(3) 0.3920(10) 0.329(2) 1.000 0.045(7) . . .
H12 H Uiso 0.219(3) 0.2492(11) 0.150(2) 1.000 0.067(8) . . .
H12' H Uiso 0.138(3) 0.2104(12) 0.238(2) 1.000 0.072(8) . . .
H13 H Uiso -0.092(5) 0.1945(17) 0.099(4) 1.000 0.143(15) . . .
H14 H Uiso 0.16414 0.17253 -0.02117 1.000 0.2089 . . .
H14' H Uiso -0.00126 0.14441 -0.05987 1.000 0.2089 . . .
H14" H Uiso 0.09879 0.12758 0.06782 1.000 0.2089 . . .
H17 H Uiso -0.096(3) 0.1899(10) 0.300(2) 1.000 0.047(7) . . .
H18 H Uiso -0.088(3) 0.1212(15) 0.447(3) 1.000 0.078(10) . . .
H19 H Uiso -0.012(4) 0.1489(17) 0.657(4) 1.000 0.124(13) . . .
H20 H Uiso 0.066(4) 0.2501(15) 0.712(3) 1.000 0.110(12) . . .
H23 H Uiso -0.101(3) 0.4621(12) 0.565(3) 1.000 0.083(9) . . .
H24 H Uiso -0.278(3) 0.5104(12) 0.409(2) 1.000 0.066(8) . . .
H25 H Uiso -0.337(4) 0.4716(15) 0.218(3) 1.000 0.102(11) . . .
H26 H Uiso -0.208(3) 0.3812(11) 0.167(2) 1.000 0.064(8) . . .

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_atom_site_aniso_U_33
_atom_site_aniso_U_23
_atom_site_aniso_U_13
_atom_site_aniso_U_12
S1 0.0896(6) 0.1333(8) 0.0607(5) -0.0188(5) -0.0147(4) 0.0014(5)

```

S2 0.0773(5) 0.1227(7) 0.0479(4) -0.0074(4) 0.0071(3) 0.0019(5)
C1 0.0671(17) 0.097(2) 0.0434(15) 0.0077(15) 0.0114(12) 0.0224(15)
C2 0.088(2) 0.123(3) 0.057(2) 0.018(2) 0.0186(17) 0.030(2)
C3 0.100(3) 0.115(3) 0.069(2) 0.043(2) 0.032(2) 0.039(2)
C4 0.076(2) 0.072(2) 0.089(2) 0.0224(18) 0.0369(18) 0.0209(17)
C5 0.108(3) 0.082(3) 0.121(4) 0.031(3) 0.052(3) 0.020(2)
C6 0.096(3) 0.064(2) 0.146(4) 0.004(3) 0.056(3) -0.002(2)
C7 0.072(2) 0.085(3) 0.097(3) -0.015(2) 0.031(2) -0.0036(18)
C8 0.0592(17) 0.074(2) 0.0675(19) 0.0002(17) 0.0258(14) -0.0036(15)
C9 0.0561(15) 0.0692(18) 0.0579(16) 0.0086(14) 0.0256(13) 0.0141(14)
C10 0.0492(14) 0.0766(18) 0.0454(14) 0.0067(13) 0.0139(11) 0.0119(13)
C11 0.0473(13) 0.0636(16) 0.0403(13) -0.0029(12) 0.0039(10) 0.0009(12)
C12 0.0521(16) 0.0735(19) 0.0555(16) -0.0078(15) 0.0091(13) -0.0003(14)
C13 0.084(2) 0.095(2) 0.077(2) -0.0250(18) -0.0004(18) 0.0015(19)
C14 0.128(3) 0.139(3) 0.138(4) -0.075(3) -0.011(3) 0.033(3)
C15 0.0465(13) 0.0617(16) 0.0453(14) 0.0021(12) 0.0060(11) -0.0005(12)
C16 0.0492(14) 0.0726(19) 0.0627(17) 0.0130(14) 0.0174(12) 0.0051(13)
C17 0.0632(19) 0.079(2) 0.088(2) 0.012(2) 0.0244(18) -0.0028(16)
C18 0.072(2) 0.077(3) 0.154(4) 0.036(3) 0.045(2) 0.005(2)
C19 0.076(2) 0.148(5) 0.116(4) 0.075(4) 0.043(2) 0.029(3)
C20 0.069(2) 0.134(4) 0.076(3) 0.038(3) 0.0272(18) 0.023(2)
C21 0.0554(16) 0.102(2) 0.0544(16) 0.0211(16) 0.0212(13) 0.0130(15)
C22 0.0560(16) 0.0746(19) 0.0575(17) -0.0074(14) 0.0189(13) -0.0080(14)
C23 0.074(2) 0.083(2) 0.077(2) -0.0240(19) 0.0300(17) -0.0136(18)
C24 0.079(2) 0.063(2) 0.110(3) -0.008(2) 0.040(2) 0.0018(17)
C25 0.0631(19) 0.073(2) 0.087(2) 0.0079(18) 0.0261(17) 0.0046(16)
C26 0.0551(16) 0.0690(19) 0.0609(18) 0.0016(15) 0.0161(14) 0.0020(14)
C27 0.0512(14) 0.0640(16) 0.0496(14) 0.0012(12) 0.0175(11) -0.0040(12)

#=====

10. MOLECULAR GEOMETRY

_geom_special_details

;

Bond distances, angles etc. have been calculated using the rounded fractional coordinates. All esds are estimated from the variances of the (full) variance-covariance matrix. The cell esds are taken into account in the estimation of distances, angles and torsion angles

;

loop_

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_geom_bond_atom_site_label_2

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_geom_bond_site_symmetry_1

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S1 C1 1.754(3) . . . yes

S1 C13 1.790(4) . . . yes

S2 C21 1.753(3) . . . yes

S2 C22 1.756(3) . . . yes

C1 C2 1.398(5) . . . no

C1 C10 1.388(3) . . . no

C2 C3 1.334(6) . . . no

C3 C4 1.421(5) . . . no

C4 C5 1.400(5) . . . no

C4 C9 1.419(4) . . . no

C5 C6 1.353(8) . . . no

C6 C7 1.404(6) . . . no

C7 C8 1.360(5) . . . no

C8 C9 1.413(4) . . . no

C9 C10 1.428(4) . . . no

C10 C11 1.484(3) . . . no

C11 C12 1.503(4) . . no
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