

A spiro-bifluorene based 3D electron acceptor with dicyanovinylene substitution for solution processed non-fullerene organic solar cells

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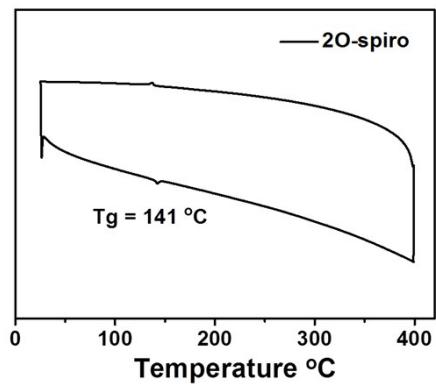
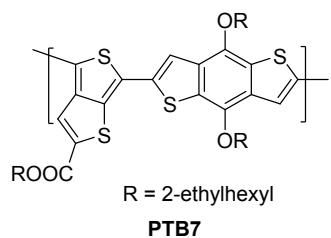


Figure S1. DSC thermograms of the **2O-spiro** on heating and cooling cycles, heating/cooling rate = 10 °C/min.



Scheme S1. The chemical structure of **PTB7**

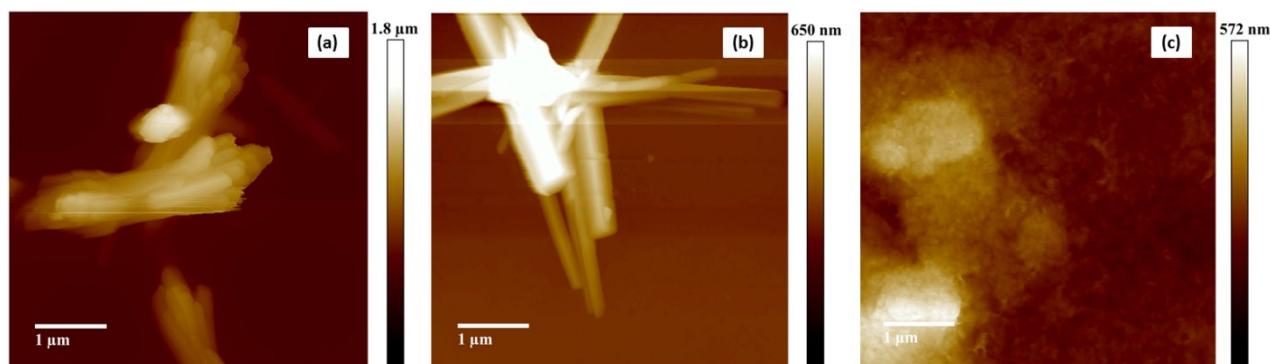


Figure S2. AFM height images of PTB7:4CN-spiro (1:1, w/w) blend processed by different solvents: (a) tetrachloroethane/DCB; (b) DCB; (c) tetrachloroethane/3%DIO.

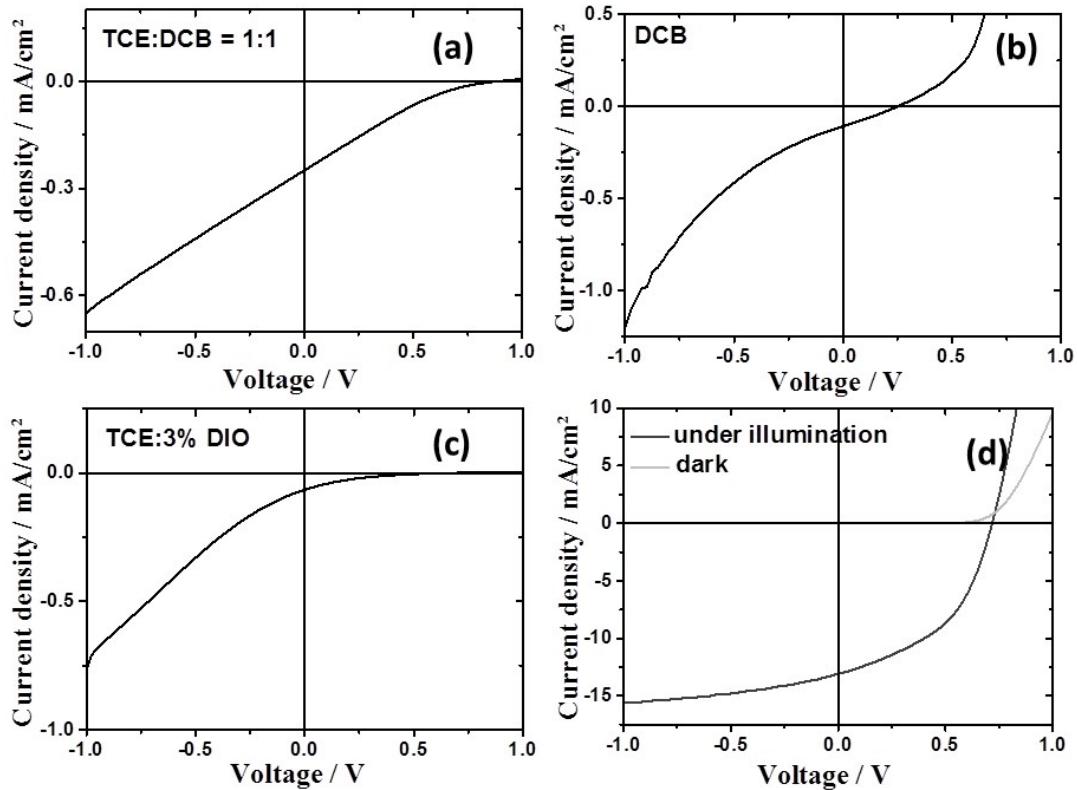
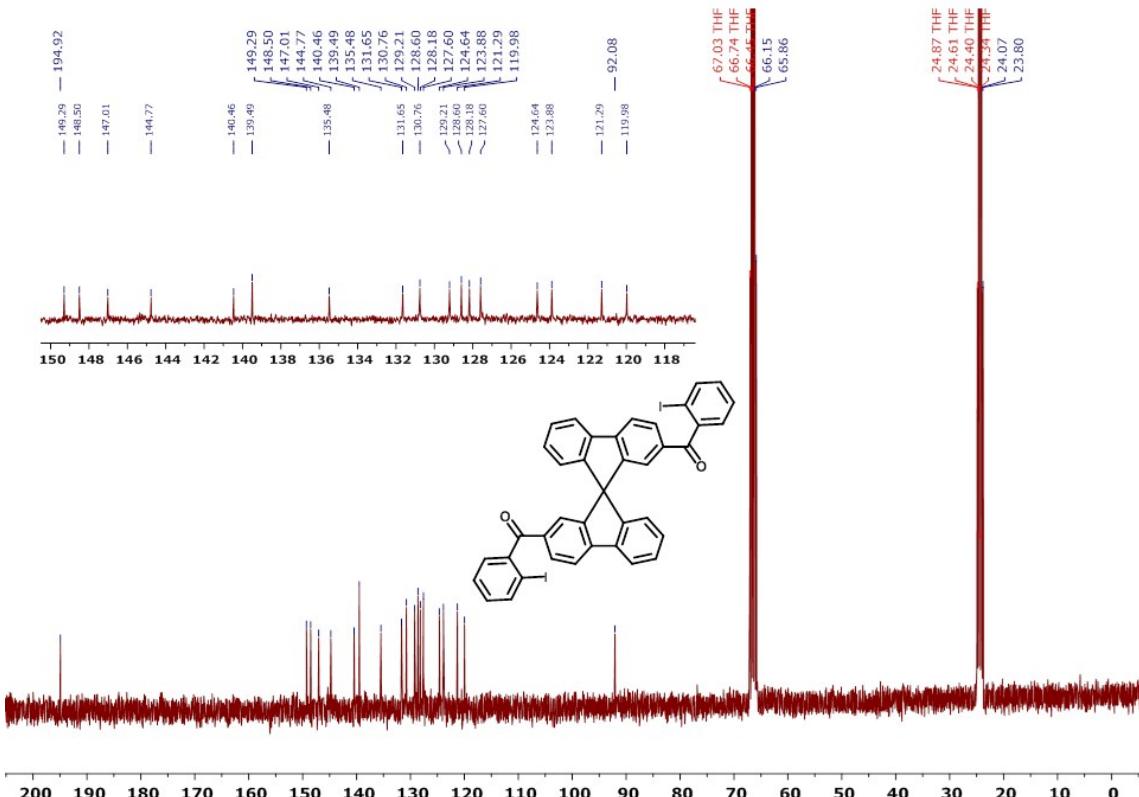
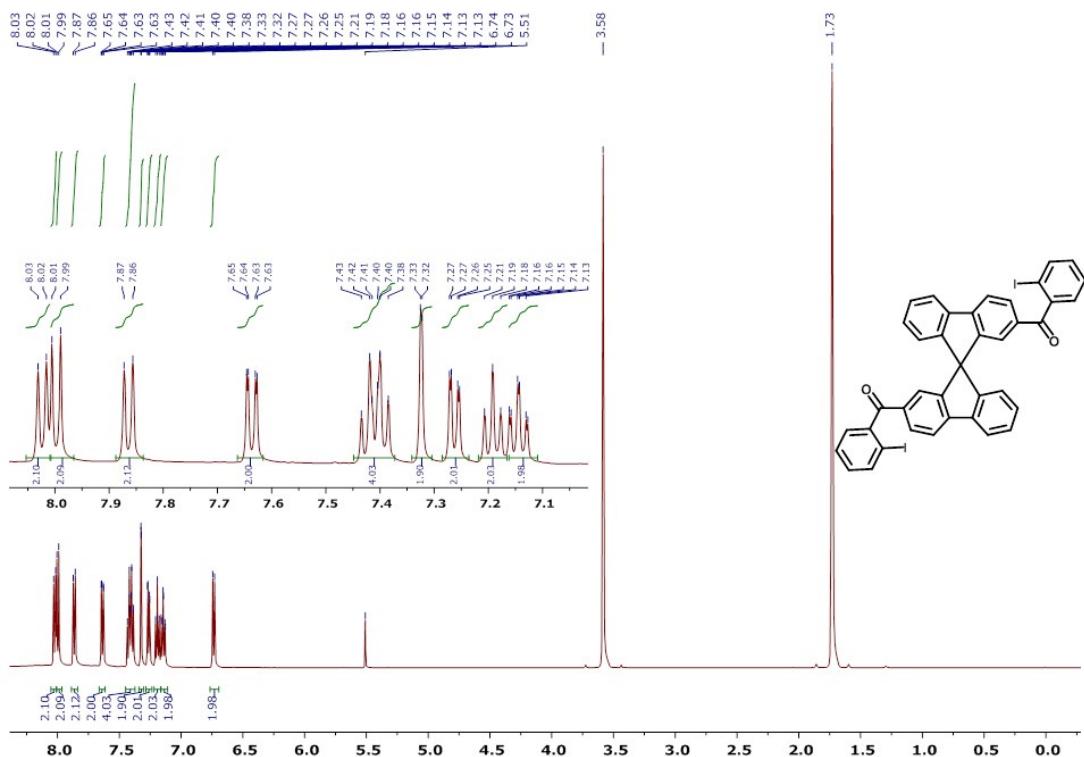


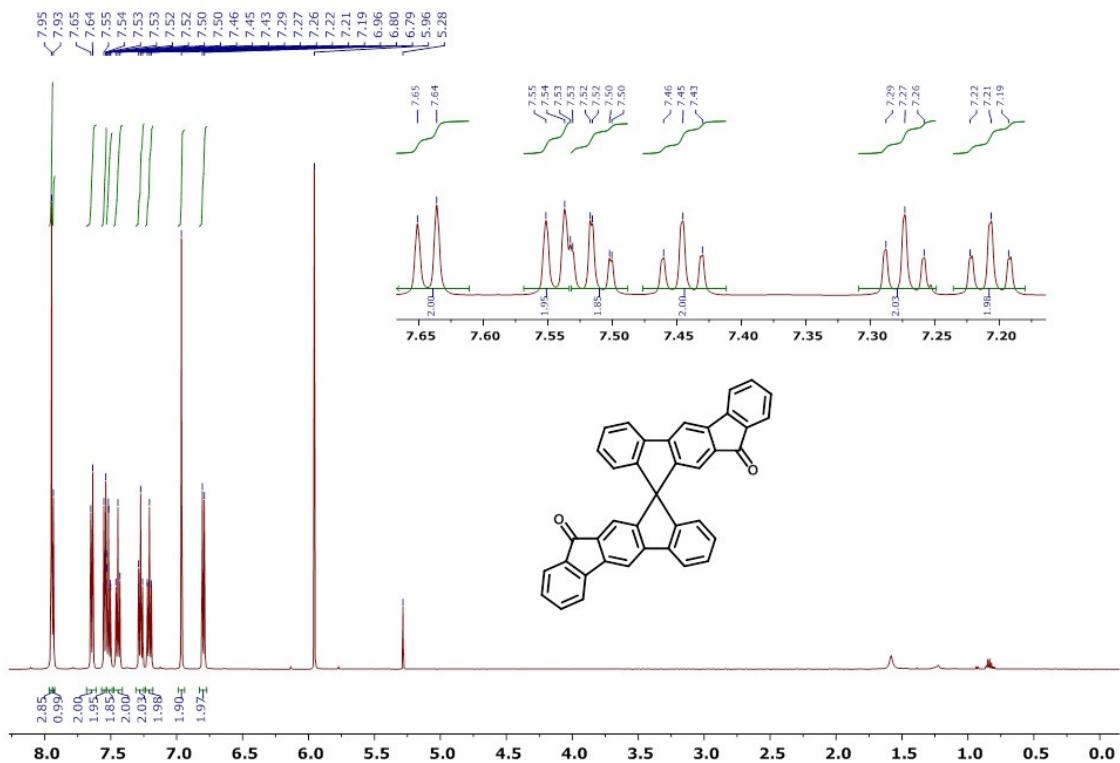
Figure S3. Current density-voltage (J-V) curves of the OPVs processed by different solvents: (a) tetrachloroethane/DCB; (b) DCB; (c) tetrachloroethane/3%DIO; (d) PTB7:PC₇₀BM solar cell processed from CB/3%DIO.

¹H-NMR and ¹³C-NMR Spectrum

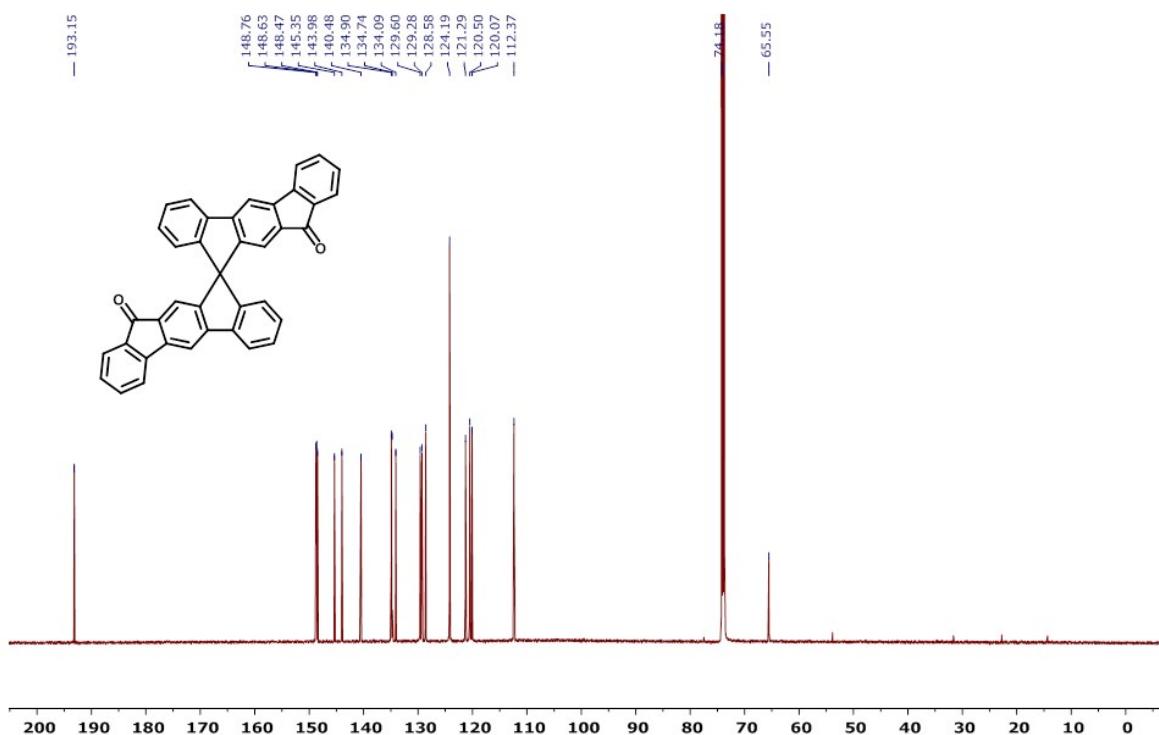
¹H NMR spectrum of Compound 1 in C₂D₂Cl₄



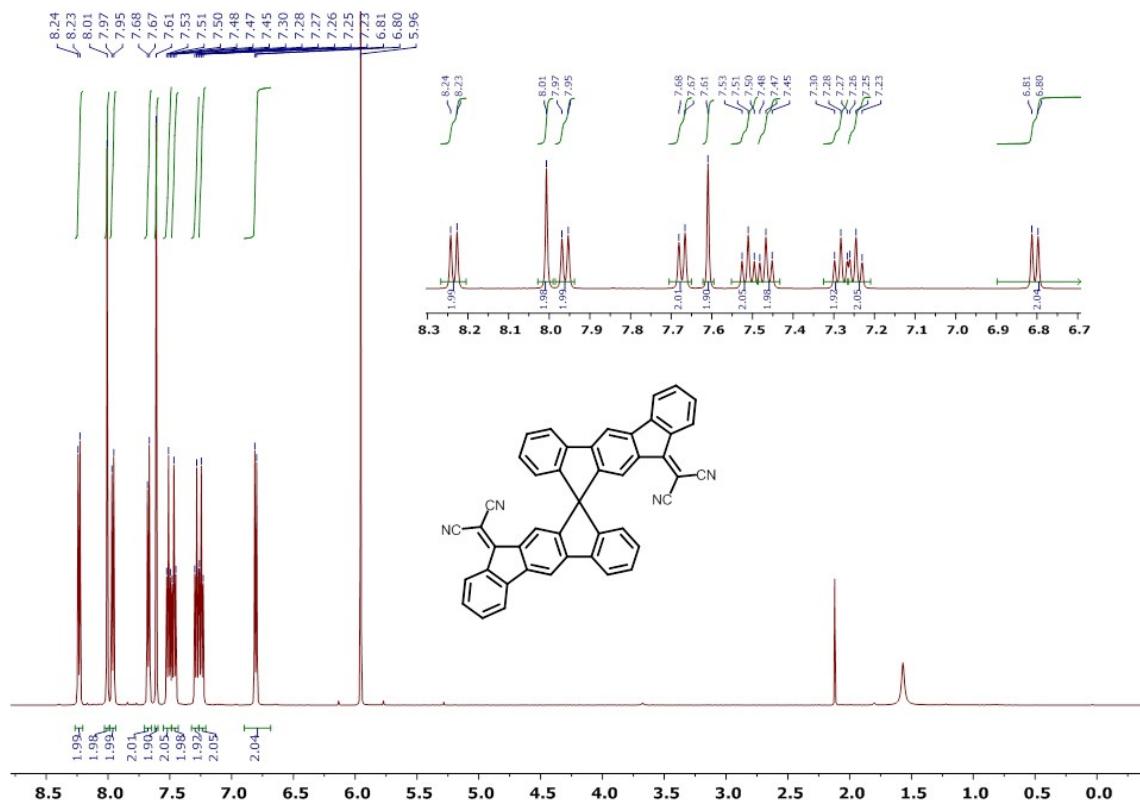
¹H NMR spectrum of Compound 2O-spiro in C₂D₂Cl₄



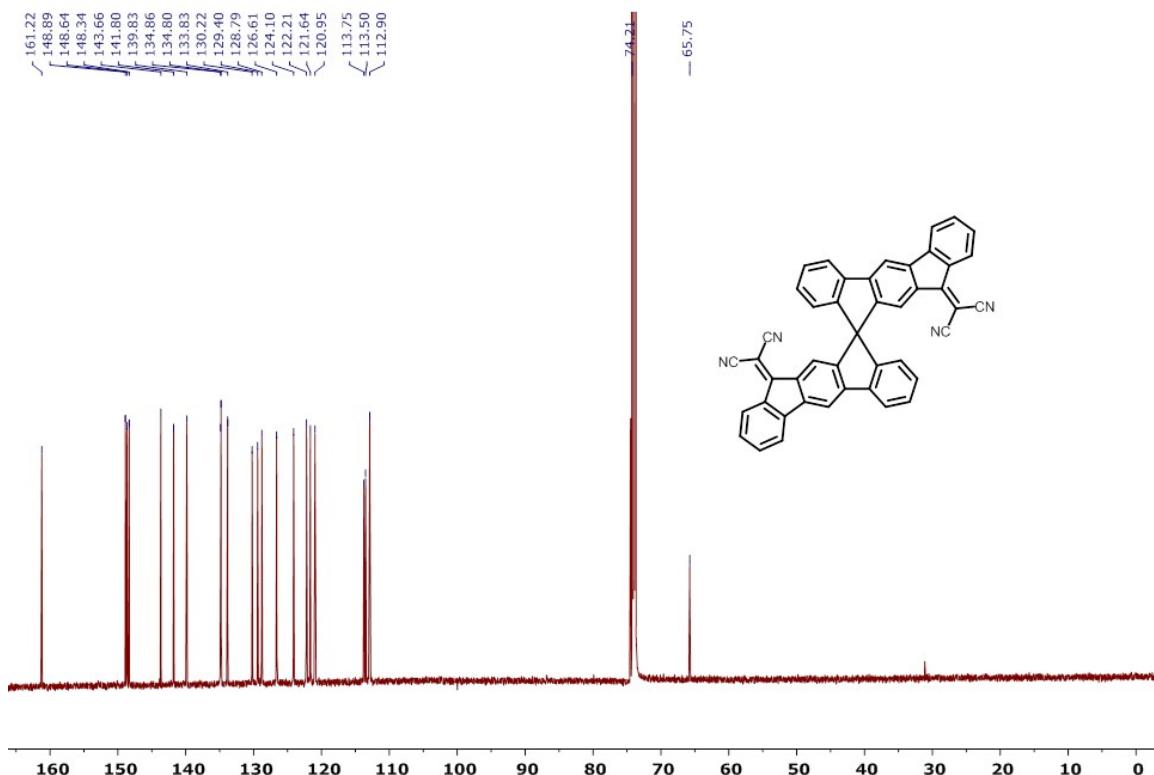
¹³C NMR spectrum of Compound 2O-spiro in C₂D₂Cl₄



¹H NMR spectrum of Compound 4CN-spiro in C₂D₂Cl₄



^{13}C NMR spectrum of Compound 4CN-spiro in $\text{C}_2\text{D}_2\text{Cl}_4$



HRMS spectra

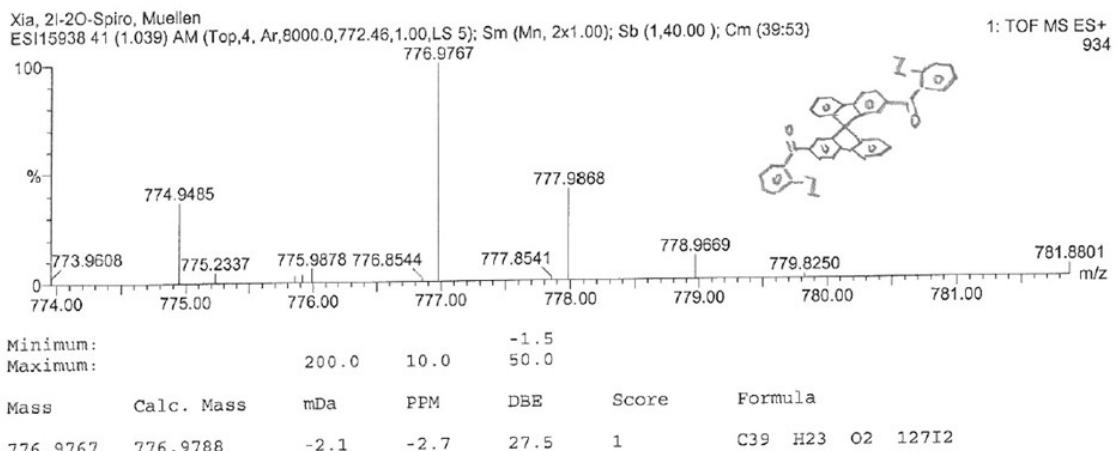
Elemental Composition Report

Page 1

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0
Isotope cluster parameters: Separation = 1.0 Abundance = 1.0%

Monoisotopic Mass, Odd and Even Electron Ions
19 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass)



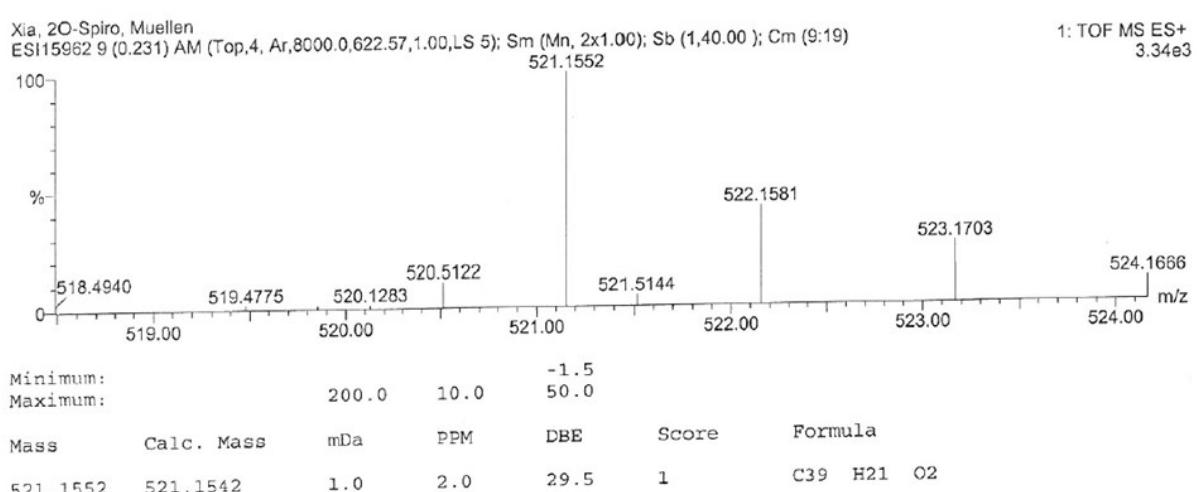
Elemental Composition Report

Page 1

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0
Isotope cluster parameters: Separation = 1.0 Abundance = 1.0%

Monoisotopic Mass, Odd and Even Electron Ions
14 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass)



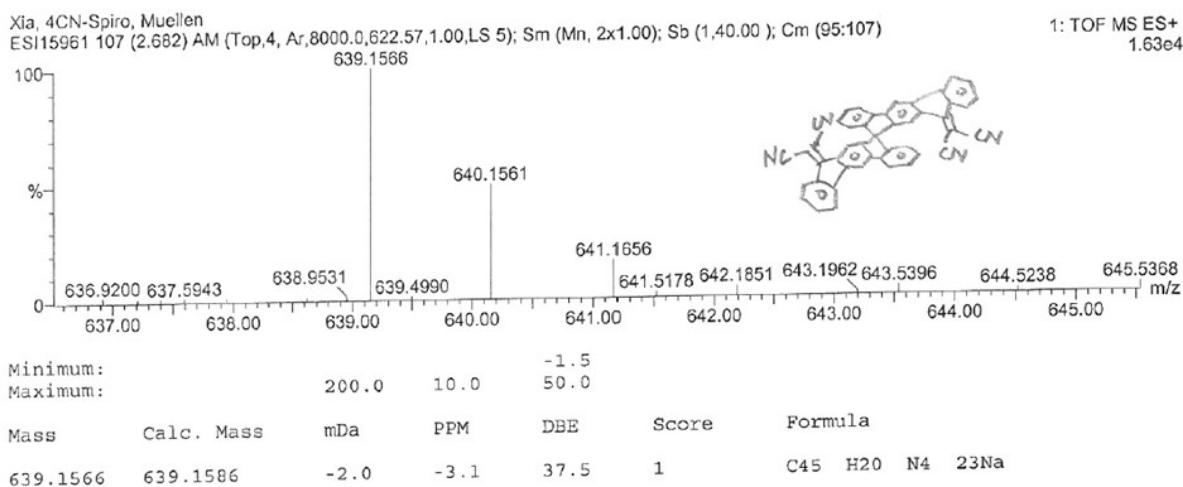
Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Isotope cluster parameters: Separation = 1.0 Abundance = 1.0%

Monoisotopic Mass, Odd and Even Electron Ions

9 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass)

**Crystal data of 2O-spiro and 4CN-spiro**

The crystal structures were solved by the software Mercury 3.1 and the crystal images were created with the software PowerPoint 2010.

Table 1. Crystal data and structure refinement for compound 2O-spiro.

formula	C ₃₉ H ₂₀ O ₂		
molecular weight	520.55 g mol ⁻¹		
absorption	$\mu = 0.08 \text{ mm}^{-1}$		
crystal size	0.26 x 0.27 x 0.3 mm ³ yellow block		
space group	P 2 ₁ /c (monoclinic)		
lattice parameters	a = 17.6736(8) \AA		
(calculate from	b = 15.8144(5) \AA	$\beta = 109.009(3)^\circ$	
21390 reflections with	c = 19.7873(9) \AA		
2.5° < θ < 28.4°	V = 5228.9(4) \AA^3	z = 8	F(000) = 2160
temperature	-80°C		
density	$d_{\text{xray}} = 1.322 \text{ g cm}^{-3}$		

data collection

diffractometer	STOE IPDS 2T
radiation	Mo-K α Graphitmonochromator
Scan – type	ω scans
Scan – width	1°
scan range	$2^\circ \leq \theta < 28^\circ$ $-18 \leq h \leq 23 \quad -20 \leq k \leq 18 \quad -26 \leq l \leq 26$
number of reflections:	
measured	34000
unique	12573 ($R_{\text{int}} = 0.0551$)
observed	7447 ($ F /\sigma(F) > 4.0$)

daten correction, structure solution and refinement

corrections	Lorentz and polarisation correction.
Structure solution	Program: SIR-2004 (Direct methods)
refinement	Program: SHELXL-2014 (full matrix). 739 refined parameters, weighting scheme: $w=1/[\sigma^2(F_o^2) + (0.0514*P)^2+0.24*P]$ with $(\text{Max}(F_o^2,0)+2*F_c^2)/3$. H-atoms at calculated positions and refined with isotropic displacement parameters, non H-atoms refined anisotropically.
R-values	$wR2 = 0.1168$ ($R1 = 0.048$ for observed reflections, 0.0975 for all reflections)
goodness of fit	S = 1.003
maximum deviation	
of parameters	0.001 * e.s.d
maximum peak height in diff. Fourier synthesis	0.27, -0.17 e \AA^{-3}

Table 2. Crystal data and structure refinement for compound 4CN-spiro.

formula	$2(\text{C}_{45}\text{H}_{20}\text{N}_4), \text{CH}_2\text{Cl}_2$
molecular weight	1318.23 g mol $^{-1}$
absorption	$\mu = 1.273 \text{ mm}^{-1}$ correction with 6 faces

transmission	$T_{\min} = 0.785, T_{\max} = 0.987$
crystal size	0.01 x 0.01 x 0.2 mm ³ brown needle
space group	P -1 (triclinic)
lattice parameters	a = 14.0610(12) Å $\alpha = 99.814(7)^\circ$
(calculate from 15604 reflections with	b = 15.0528(14) Å $\beta = 90.613(7)^\circ$
2.7° < θ < 51.2°)	c = 33.272(3) Å $\gamma = 92.172(7)^\circ$
	V = 6933.3(11) Å ³ z = 4 F(000) = 2712
temperature	-60°C
density	$d_{\text{xray}} = 1.263 \text{ gcm}^{-3}$

data collection

diffractometer	STOE IPDS 2T
radiation	Cu-K _α mirror system IμS
Scan – type	ω scans
Scan – width	1°
scan range	$3^\circ \leq \theta < 68^\circ$
number of reflections:	$-15 \leq h \leq 15 \quad -18 \leq k \leq 17 \quad -39 \leq l \leq 39$
measured	94677
unique	23561 ($R_{\text{int}} = 0.3682$)
observed	3389 ($ F /\sigma(F) > 4.0$)

data correction, structure solution and refinement

corrections	Lorentz and polarisation correction.
Structure solution	Program: SHELXD-2013
refinement	Program: SHELXL-2013 (full matrix). 1819 refined parameters, weighting scheme: $w=1/[\sigma^2(F_o^2) + (0.0486*P)^2]$ with $(\text{Max}(F_o^2, 0) + 2*F_c^2)/3$. H-atoms at calculated positions and refined with isotropic displacement parameters, non H-atoms refined anisotropically.
R-values	$wR2 = 0.2318$ ($R1 = 0.0725$ for observed reflections, 0.3689 for all reflections)
goodness of fit	S = 0.68

maximum deviation of parameters	0.001 * e.s.d
maximum peak height in diff. Fourier synthesis	0.32, -0.32 eÅ ⁻³
remark	structure contains four different molecules and a number of solvent molecules. One potential place for solvent molecules could not be located. Squeeze was use for moldelling the electron density.