1,3-Dipolar cycloaddition of isatin-derived azomethine ylides with 2*H*-azirines: stereoselective synthesis of 1,3diazaspiro[bicyclo[3.1.0]hexane]oxindoles

Anikó Angyal^{a,b}, András Demjén^a, Veronika Harmat^c, János Wölfling^b, László G. Puskás^a, Iván Kanizsai^{a,*}

^aAVIDIN Ltd., Alsó kikötő sor 11/D, Szeged, H-6726, Hungary.

^bDepartment of Organic Chemistry, University of Szeged, Dóm tér 8, H-6720, Szeged, Hungary.

^cEötvös Loránd University, Institute of Chemistry, Laboratory of Structural Chemistry and Biology, and MTA-ELTE Protein Modeling Research Group, Pázmány P. sétány 1/A, H-1117, Budapest, Hungary.

Tel.: +36-62/202107; fax: +36-62/202108; e-mail: i.kanizsai@avidinbiotech.com

Supporting Information

Table of Contents

1.	Reaction optimization	S2
2.	¹ H and ¹³ C NMR spectra for all products	S3
3.	1D and 2D NMR spectra of compounds (±)-4a and (±)-4b	S32
4.	1D and 2D NMR spectra of compound (±)-12a	S37
5.	1D and 2D NMR spectra of compound (±)-17a	S40
6.	1D and 2D NMR spectra of compound (±)-31a	S43
7.	Crystallographic studies of compounds (±)-4a and (±)-31a	S46
	7.1. Experimental	S46
	7.2. Results	S47
8.	References	

1. Reaction optimization

Table S1. Screening of reaction conditions

0			EtOOC,			
	N +	Соон	+ <u>N</u> CC	DOEt so	Ivent Ar	
	1	2m	(±)-3a		(±)-31	H a
Entry ^a	Solvent	Solvent (mL)	Temp. (°C)	Time (h)	(±)-3a (equiv.)	Yield (%) ^b
1	MeOH	1	rt	24	1	10
2	EtOH	1	rt	24	1	25
3	IPA	1	rt	24	1	44
4	TFE	1	rt	24	1	6
5	MeCN	1	rt	24	1	10
6	MeNO ₂	1	rt	24	1	5
7	DMF	1	rt	24	1	28
8	EtOAc	1	rt	24	1	2
9	THF	1	rt	24	1	6
10	Toluene	1	rt	24	1	-
11	CHCl ₃	1	rt	24	1	16
12	DCM	1	rt	24	1	7
13	IPA	1	rt	24	2	51
14	IPA	1	rt	24	3	60
15	IPA	0.25	rt	24	3	55
16	IPA	0.5	rt	24	3	63
17	IPA	1.5	rt	24	3	62
18	IPA	2	rt	24	3	68
19	IPA	2.5	rt	24	3	63
20	IPA	3	rt	24	3	61
21	IPA	2	0 °C	48	3	58
22	IPA	2	60 °C	4	3	52
23	IPA	2	80 °C	2.5	3	56

^{*a*}Reaction conditions: isatin (0.125 mmol), L-proline (0.15 mmol), 2*H*-azirine, anhydrous solvent, argon atmosphere. ^{*b*}Determined by HPLC analysis.

2. ¹H and ¹³C NMR spectra for all products



Figure S1. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-5a



Figure S2. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-5a



Figure S3. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-6a



Figure S4. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-6a



Figure S5. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-7a



Figure S6. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-7a



Figure S7. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-8a



Figure S8. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-8a



Figure S9. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-9a



Figure S10. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-9a



Figure S11. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-10a



Figure S12. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-10a



Figure S13. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-11a



Figure S14. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-11a



Figure S15. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-13a



Figure S16. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-13a



Figure S17. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-14a



Figure S18. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-14a



Figure S19. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-15a



Figure S20. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-15a



Figure S21. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-16a



Figure S22. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-16a



Figure S23. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-18a



Figure S24. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-18a



Figure S25. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-19a



Figure S26. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-19a



Figure S27. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-20a



Figure S28. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-20a



Figure S29. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-21a



Figure S30. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-21a



Figure S31. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-22a



Figure S32. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-22a



Figure S33. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-23a



Figure S34. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-23a



Figure S35. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-24a



Figure S36. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-24a



Figure S37. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-25a



Figure S38. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-25a



Figure S39. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-26a



Figure S40. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-26a



Figure S41. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-27a



Figure S42. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-27a



Figure S43. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-28a



Figure S44. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-28a



Figure S45. ¹H NMR spectrum (500 MHz, DMSO-*d*₆) of the compound (±)-29a



Figure S46. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-29a



Figure S48. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-30a



Figure S49. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-32a



Figure S50. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-32a



Figure S51. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-33a



Figure S52. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-33a



Figure S53. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-34a



Figure S54. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-34a



Figure S55. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-35a



Figure S56. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-35a



Figure S57. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-36a



Figure S58. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-36a



3. 1D and 2D NMR specra of compounds (±)-4a and (±)-4b

Figure S59. ¹H NMR spectrum (500 MHz, CDCl₃) of the compound (±)-4a



Figure S60. ¹³C NMR spectrum (126 MHz, CDCl₃) of the compound (±)-4a



Figure S61. HSQC spectrum of the compound (±)-4a in CDCl₃



Figure S62. HMBC spectrum of the compound (±)-4a in CDCl₃



Figure S63. COSY spectrum of the compound (±)-4a in CDCl₃



Figure S64. COSY spectrum of the compound (±)-4a in CDCl₃



Figure S65. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-4b



Figure S66. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-4b



Figure S67. HSQC spectrum of the compound (±)-4b in DMSO- d_6



Figure S68. HMBC spectrum of the compound (±)-4b in DMSO- d_6



Figure S69. NOESY spectrum of the compound (\pm)-4b in DMSO- d_6

4. 1D and 2D NMR specra of compound (±)-12a



Figure S70. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-12a



Figure S71. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-12a



Figure S72. HSQC spectrum of the compound (±)-12a in DMSO- d_6



Figure S73. HMBC spectrum of the compound (±)-12a in DMSO- d_6



Figure S74. COSY spectrum of the compound (±)-12a in DMSO- d_6



Figure S75. NOESY spectrum of the compound (\pm)-12a in DMSO- d_6

5. 1D and 2D NMR specra of compound (±)-17a



Figure S76. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-17a



Figure S77. ¹³C NMR spectrum (126 MHz, DMSO-*d*₆) of the compound (±)-17a



Figure S78. HSQC spectrum of the compound (±)-17a in DMSO- d_6



Figure S79. HMBC spectrum of the compound (±)-17a in DMSO- d_6



Figure S80. NOESY spectrum of the compound (±)-17a in DMSO- d_6

6. 1D and 2D NMR specra of compound (±)-31a



Figure S81. ¹H NMR spectrum (500 MHz, DMSO- d_6) of the compound (±)-31a



Figure S82. ¹³C NMR spectrum (126 MHz, DMSO- d_6) of the compound (±)-31a



Figure S83. HSQC spectrum of the compound (±)-31a in DMSO- d_6



Figure S84. HMBC spectrum of the compound (±)-31a in DMSO- d_6



Figure S85. COSY spectrum of the compound (±)-31a in DMSO- d_6



Figure S86. NOESY spectrum of the compound (±)-31a in DMSO- d_6

Crystallographic studies of compounds (±)-4a and (±)-31a Experimental

Racemic compounds (\pm) -4a and (\pm) -31a were crystallized from dichloromethane and tetrahydrofuran, respectively. X-ray diffraction data were collected on a Rigaku XtaLab Synergy-R diffractometer using Cu -K α radiation (λ = 1.54184 Å). Data reduction was carried out using the software provided with the diffractometer (CrysAlisPro 1.171.40.14e (Rigaku OD, 2018)). The structure was solved by direct methods using Olex2 v1.2 and refined by fullmatrix least-squares techniques (SHELXL-2014/7) on $F^{2,1,2}$ Hydrogen atoms were refined in the riding positions for both structures. The crystallographic parameters, data collection, and structure refinement details are summarized in Table S2. Compound (±)-4a crystallized in space group $P2_12_12_1$, with one molecule per asymmetric unit. The crystal showed partial merohedral twinning, with fraction of 0.2068 for the inverted structure. Flack-x parameter refined to a value of 0.2(2). Compound (±)-31a crystallized in space group P21/n. The asymmetric unit contains one molecule of (\pm) -31a and a tetrahydrofuran molecule with occupancy of 0.5, with a crystallographic inversion center within the tetrahydrofuran ring. Geometric and rigid-bond restraints were applied to the disordered tetrahydrofuran molecule. The structures were analyzed using Mercury program.³ Validation was carried out using CheckCIF/PLATON.⁴ The figures were created using Mercury and PyMOL.⁵ Structures of (±)-4a and (±)-31a were deposited with the Cambridge Crystallographic Data Centre and can be obtained free of charge with CCDC deposition numbers CCDC 1890661 and CCDC 1890659, respectively.

Table S2. X-ray crystallographic data and structural refinements details for (\pm) -4a and (\pm) -
31a

Compound	(±)-4a	(±)-31a
CCDC reference number	CCDC 1890661	CCDC 1890659
Asymmetric unit contents		
Empirical formula	$C_{21}H_{23}N_3O_3$	$C_{20}H_{25}N_3O_{3.5}$
Formula weight	363.41	363.43
Temperature (K)	299.0 (2)	298.0 (2)
Wavelength (Å)	1.54184	1.54184
Crystal system	orthorhombic	monoclinic
Space group	P2 ₁ 2 ₁ 2 ₁	P2 ₁ /n

Unit cell dimensions <i>a</i> , <i>b</i> , <i>c</i>	9.7567(1),	10.65608(8),16.5098(1),10.94821(9)
(Å)	13.6713(1),	90.0, 99.8206(8), 90.0
α, β,	14.3251(1)	
γ (°)	90.0, 90.0, 90.0	
V(Å)	1910.77(3)	1897.89(3)
Ζ	4	4
D_{calc} (g /cm ³)	1.263	1.272
$\theta_{\min}, \theta_{\max}$ (°)	6.476, 75.450	4.897, 75.480
Reflections collected/unique	21399 / 3820	41156, 3824
R _{int}	0.0192	0.0241
Completeness to θ_{max} (%)	98.2	97.1
Data/restraints/parameters	3820 / 0 / 253	3824 / 5 / 264
Goodness-of-fit (GOF) on	1.062	1.039
F^2		
$R1 / wR2 [I > 2\sigma(I)]$	0.0311 / 0.0820	0.0440 / 0.1185
R1 / wR2 (all data)	0.0317 / 0.0833	0.0462 / 0.1205
Largest difference peak/hole	0.154 / -0.163	0.245 -0.202
(e/ Å ³)		

7.2. Results

Configuration of chirality centers were confirmed by the crystal structures of (\pm) -4a and (\pm) -31a. The structures are shown in Figure S87, and selected geometric parameters are compiled in Table S3. Interestingly, (\pm) -4a crystallized in space group P2₁2₁2₁ containing only one of the enantiomers in the unit cell. However it showed partial racemic twinning, resulting a ratio of 0.8 : 0.2 for the two enantiomers in the crystal investigated. (\pm) -31a crystallized in a centrosymmetric space group, thus contains the 1 : 1 raceme mixture. The five membered ring (N2-C10-N3-C8-C9) is in an envelope conformation with the N3 being the puckering atom because of the steric constraints caused by the fused cyclopropane moiety. In (\pm) -31a the C10-N3-C14-C15-C16 ring shows an envelope conformation similarly, with the atom opposite to the sterically constrained part being the puckering atom (C15). Minor differences between respective bond lengths and bond angles of the two structures can be explained by the attachment of the second five-membered ring in (\pm) -31a. Crystal packing of (±)-4a (Figure S88) shows a 3D hydrogen bonded network of molecules. In order to understand the formation of partial racemic twinnig, we compared the direction dependence of molecular contacts in the structure. A column-like association of molecules is connected by N1-H1...N2 hydrogen bonds parallel to unit cell axis *b*. Perpendicular to this direction, longer N3-H3...O2 hydrogen bonds connect the 'column' to neighboring molecules and the phenyl groups fit between the phenyl and oxindole moieties of neighboring molecules in a cog-like way, forming edge-to-face aromatic contacts with the oxindole moieties of those. In the third dimension a smooth edge of the column is formed by its oxindole and acetyl groups. By comparing the shapes of columns formed by the inverted molecules, we speculate that these inverted columns rotated around unit cell axis c by 180° could fit this interaction pattern (Figure S89).

Crystal packing of (±)-31a (Figure S90) shows hydrogen bonded layers of molecules parallel to **b** and **a**+**c** vectors, with disordered tetrahydrofuran molecules sitting between the layers. The aromatic ring is involved mainly in C-H.. π type interactions established with aliphatic groups.



Figure S87. Crystal structures of compounds (±)-4a and (±)-31a, respectively. Atomic displacement ellipsoids are contoured at the 0.3 probability level.

Compound	(±)-4a	(±)-31a		
Geometric parameter				
Bond lengths (Å), bond angles and torsion angles(°)				
N2-C10	1.492(2)	1.487(2)		
C10-N3	1.467(2)	1.475(2)		
N3-C8	1.471(2)	1.458 (2)		
C8-C9	1.541(2)	1.536 (2)		
C9-N2	1.487(2)	1.471(2)		
C9-C11	1.482(2)	1.499(2)		
C11-N2	1.483(2)	1.471(2)		
N2-C10-N3	108.4(1)	108.3(1)		
C10-N3-C8	105.6(1)	107.8(1)		
N3-C8-C9	105.3(1)	104.9(1)		
C8-C9-N2	106.9(1)	108.8(1)		
C9-N2-C10	105.7 (1)	105.8(1)		
C9-C11-N2	60.20(9)	59.39(8)		
C11-N2-C9	59.9(1)	61.25(8)		
N3-C8-C7	107.7(1)	106.4(1)		
N3-C8-C5	113.2(1)	117.5(1)		
C9-C8-C7	111.0(1)	111.1(1)		
C9-C8-C5	117.3(1)	115.4(1)		
N2-C11-C13-O3	75.1	179.7		
C13-O3-C20-C21 (C13-O3-C17-	177.3	175.68		
C18)				
RMS deviation of the ring atoms from the plane (Å)				
N2-C10-N3-C8-C9	0.1199	0.0897		
C10-N3-C14-C15-C16	-	0.1608		
C1-C2-C3-N1-C4-C5-C6-C7-	0.0107	0.0520		
C8-O1				

Table S3. Selected geometric parameters of (\pm) -4a and (\pm) -31a. (Variances in the last digitsare shown in parentheses.) Atom names are listed for (\pm) -4a and (\pm) -31a

Angle between ring planes				
N2-C10-N3-C8-C9 / N2-C9-C11	80.05(8)	79.57(7)		
N2-C10-N3-C8-C9 / C1-C2-C3-	85.52(4)	81.83(3)		
N1-C4-C5-C6-C7-C8-O1				
Hydrogen bonds (for D-HA hydrogen bond: DA distance/ HA distance / D-H-				
A angle)				
N3-H3O2ª	3.245 / 2.384 / 161.26			
C11-H11O2ª	3.472 / 2.502 / 170.41			
N1-H1N2 ^b	2.981 / 2.151 / 161.99			
N1-H1O2 ^c		2.974 / 2.151 / 159.91		
C14-H14AO1 ^d		3.301 / 2.456 / 145.35		

^a O2 atom of the symmetry equivalent molecule trans formed by [x+1/2, -y+1/2, -z]^b N2 atom of the symmetry equivalent molecule trans formed by [-x, y-1/2, -z+1/2]

^c O2 atom of the symmetry equivalent molecule trans formed by [-x+3/2, y-1/2, -z+3/2]

^d O1 atom of the symmetry equivalent molecule trans formed by [x-1/2, -y+1/2, z-1/2]



Figure S88. Crystal packing of (±)-4a (left: view along unit cell axis *a*; right: view along unit cell axis *b*). Intermolecular hydrogen bonds N3-H3...O2 and N1-H1...N2 are shown in cyan. While N3-H3...O2 hydrogen bonds form a 2D network parallel to the *ac* plane (right); N1-H1...N2 hydrogen bonds connect molecules along unit cell axis *b* (left). (Hydrogen atoms are not shown.)



Figure S89. Accommodation of a series of (\pm) -4a molecules (referred to as "column") (orange carbon atoms) in the crystal structure (**a**, left), with neighboring molecules shown (green carbon atoms). The phenyl groups fit in a cog-like way, the oxindole and acetyl groups form a smooth edge. Three types of hydrogen bonds are shown in yellow, orange and red. **a**, (view along unit cell axis *b*) right: a column cut out from the inverted structure fits in the cog-like and smooth edges and can form hydrogen bonds shown in yellow and red (yellow carbon atoms). The inverted column was rotated around axis *c* by 180°. **b**, and **c**, Comparison of the shapes of rod-like molecule structures: cog-like and smooth edges, respectively. (Hydrogen atoms are not shown.)



Figure S90. Crystal packing of (\pm) -31a (left: view along unit cell axis *a*; right: view along unit cell axis *b*). Intermolecular hydrogen bonds N1-H1...O2 and C14-H14A...O1 are shown in cyan. Hydrogen bonds form a 2D network parallel to unit cell axis *b* and **a**+**c**. The disordered solvent molecules are accommodated between layers od hydrogen bonded molecules. (Hydrogen atoms are not shown.)

8. References

- Dolomanov, O. V.; Bourhis, L. J.; Gildea, R. J.; Howard, J. A. K.; Puschmann, H. J. Appl. Cryst. 2009, 42, 339–341.
- 2. Sheldrick, G. M. Acta Cryst. 2015, C71, 3-8.
- Macrae, C. F.; Edgington, P. R.; McCabe, P.; Pidcock, E.; Shields, G. P.; Taylor, R.; Towler, M.; van de Streek, J. J. Appl. Cryst. 2006, 39, 453–457.
- 4. Spek, A. L. Acta Cryst. 2009, D65, 148–155.
- DeLano, W. L. (2002) The PyMOL Molecular Graphics System, DeLano Scientific, San Carlos, CA