Supplementary Information

HSD1787, a tetrahydro-3*H*-pyrazolo[4,3-*f*]quinoline compound synthesized via Povarov reaction, potently inhibits proliferation of cancer cell lines at nanomolar concentrations

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Figure S41: ¹H NMR of compound 19

Figure S42: ¹³C NMR of compound 19

Single crystal X-ray diffraction data

Single crystals of compound 1 were coated with trace of fomblin oil and quickly transferred to the goniometer head of a Bruker Quest diffractometer with a fixed chi angle, a sealed tube fine focus X-ray tube, single crystal curved graphite incident beam monochromator, a Photon100 CMOS area detector and an Oxford Cryosystems low temperature device. Examination and data collection were performed with Mo K α radiation ($\lambda = 0.71073$ Å) at 150 K. Data were collected, reflections were indexed and processed, and the files scaled and corrected for absorption using APEX3 [1]. The space group was assigned and the structure was solved by direct methods using XPREP within the SHELXTL suite of programs [2] and refined by full matrix least squares against F² with all reflections using Shelxl2018 [3] using the graphical interface Shelxle [4]. H atoms attached to carbon and oxygen atoms were positioned geometrically and constrained to ride on their parent atoms, with carbon hydrogen bond distances of 0.95 Å for and aromatic C-H, 0.99 Å for aliphatic CH2 moieties, and 0.84 for hydroxyl OH atoms, respectively. Hydroxyl H atoms were allowed to rotate but not to tip to best fit the experimental electron density. Uiso(H) values were set to a multiple of U_{eq}(C/O) with 1.5 for OH, and 1.2 for C-H and CH₂ units, respectively. Complete crystallographic data, in CIF format, have been deposited with the Cambridge Crystallographic Data Centre. CCDC 2017003 contains the supplementary crystallographic data for this paper. These data can be obtained free of charge from The Cambridge Crystallographic Data Centre via www.ccdc.cam.ac.uk/data_request/cif.

- [1] Bruker (2016). Apex3 v2016.9-0, Saint V8.34A, SAINT V8.37A, Bruker AXS Inc.: Madison (WI), USA, 2013/2014.
- [2] a) SHELXTL suite of programs, Version 6.14, 2000-2003, Bruker Advanced X-ray Solutions, Bruker AXS Inc., Madison, Wisconsin: USA) b) Sheldrick GM. A short history of SHELX. *Acta Crystallogr A.* **2008**, *64(1)*, 112–122.
- [3] a) Sheldrick GM. University of Göttingen, Germany, **2018**. b) Sheldrick GM. Crystal structure refinement with SHELXL. *Acta Crystallogr Sect C Struct Chem.* **2015**, 71(1), 3–8.
- [4] Hübschle CB, Sheldrick GM, Dittrich B. ShelXle: a Qt graphical user interface for SHELXL. *J. Appl. Crystallogr.* **2011**, *44*(6), 1281–1284.

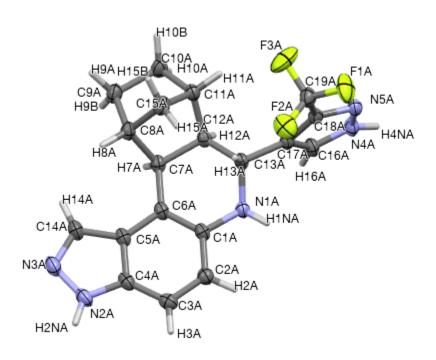


Table S1. Experimental details for single crystal X-ray diffraction of compound 1

	Compound 1	
Crystal data CCDC 2017003		
Chemical formula	$C_{19}H_{18}F_3N_5 \cdot 2(C_2H_6O)$	
$M_{ m r}$	465.52	
Crystal system, space group	Triclinic, P1	
Temperature (K)	150	
a, b, c (Å)	9.8976 (5), 15.4264 (8), 16.4770 (9)	
α, β, γ (°)	105.039 (2), 105.423 (2), 93.516 (2)	
$V(\mathring{\mathbf{A}}^3)$	2319.1 (2)	
Z	4	
F(000)	984	
$D_x (\mathrm{Mg \ m}^{-3})$	1.333	

Radiation type	Cu Kα	
No. of reflections for cell measurement	9714	
θ range (°) for cell measurement	2.9–79.6	
$\mu \text{ (mm}^{-1})$	0.87	
Crystal shape	Plate	
Colour	Colourless	
Crystal size (mm)	$0.31\times0.20\times0.04$	
Data collection		
	Bruker AXS D8 Quest CMOS	
Diffractometer	diffractometer with PhotonII charge-integrating pixel array detector	
	(CPAD)	
Radiation source	I-mu-S microsource X-ray tube	
Monochromator	Laterally graded multilayer (Goebel) mirror	
Detector resolution (pixels mm ⁻¹)	7.4074	
Scan method	ω and phi scans	
Absorption correction	Multi-scan SADABS 2016/2: Krause, L., Herbst-Irmer, R., Sheldrick G.M. & Stalke D., J. Appl. Cryst. 48 (2015) 3-10	
T_{\min}, T_{\max}	0.588, 0.754	
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	60418, 10031, 8494	
R _{int}	0.058	
θ values (°)	$\theta_{max}=80.9,\theta_{min}=2.9$	
$(\sin \theta/\lambda)_{\max} (\mathring{A}^{-1})$	0.640	

Range of h, k, l	$h = -12 \rightarrow 10, k = -19 \rightarrow 19, l = -21 \rightarrow 21$	
Refinement		
Refinement on	F^2	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.047, 0.134, 1.11	
No. of reflections	10031	
No. of parameters	740	
No. of restraints	390	
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement	
Weighting scheme	$w = \frac{1}{[\sigma^2(F_o^2)]} + (0.0646P)^2 + 0.5949P]$ where $P = (F_o^2 + 2F_c^2)/3$	
$(\Delta/\sigma)_{\max}$	0.001	
$\Delta \rho_{max}, \Delta \rho_{min} (e \ \mathring{A}^{-3})$	0.29, -0.26	
Extinction method	SHELXL2018/3 (Sheldrick 2018), $Fc^* = kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$	
Extinction coefficient	0.0021 (3)	

Computer programs: Apex3 v2017.3-0 (Bruker, 2017), *SAINT* V8.38A (Bruker, 2016), *SHELXS97* (Sheldrick, 2008), *SHELXL2018/*3 (Sheldrick, 2015, 2018), *SHELXLE* Rev946 (Hübschle *et al.*, 2011).

The structure is a racemic mixture, space group P-1. There are two crystallographically independent molecules in the cell. They are chemically identical (same conformation and configuration). There are also four ethanol molecules. A bit disordered, but well resolved. Amine H atoms are located (i.e., the H atom positions are verified by the data. Description of the minor disorder: The fluorine atoms of the two trifluoromethyl groups were refined as disordered by slight rotation. The major and minor disordered moieties were each restrained to have similar geometries. Uij components of ADPs for disordered atoms closer to each other than 2.0 Angstrom were restrained to be similar. Subject to these conditions the occupancy ratios refined to 0.807(18) to 0.193(18) (molecule A) and to 0.56(4) to 0.44(4)

(molecule B). Two of the four solvate ethanol molecules were refined as disordered by inversion at the methylene carbon atom. The two disordered moieties were restrained to have similar geometries as another not disordered ethanol molecule. Uij components of ADPs for disordered atoms closer to each other than 2.0 Angstrom were restrained to be similar. Subject to these conditions the occupancy ratio refined to 0.759(7) to 0.241(7) (disorder of O4 / O5) and to 0.619(6) to 0.381(6) (disorder of O3 / O6).

Figure S1: ¹H spectra of compound 1 in DMSO-d₆

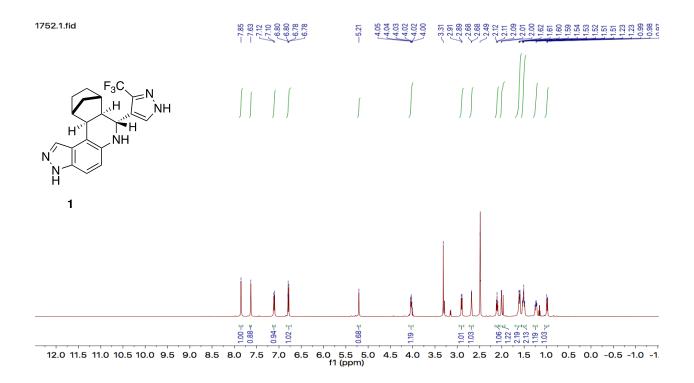


Figure S2: ¹³C spectra of compound 1 in DMSO-d₆

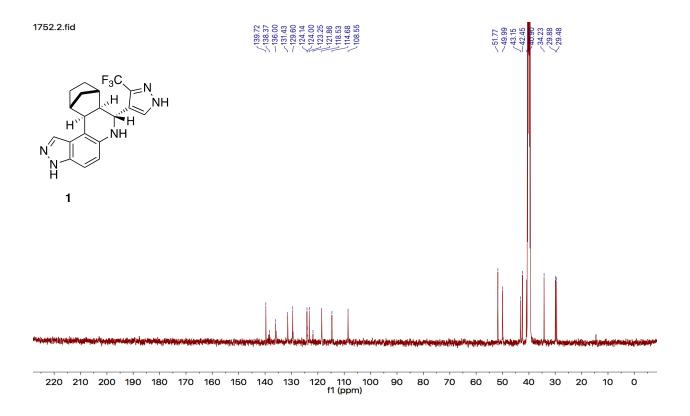


Figure S3: ¹H spectra of compound 2 in DMSO-d₆

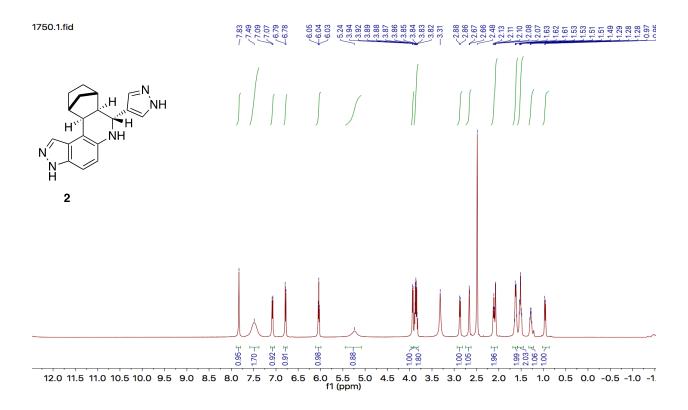


Figure S4: ¹³C spectra of compound 2 in DMSO-d₆

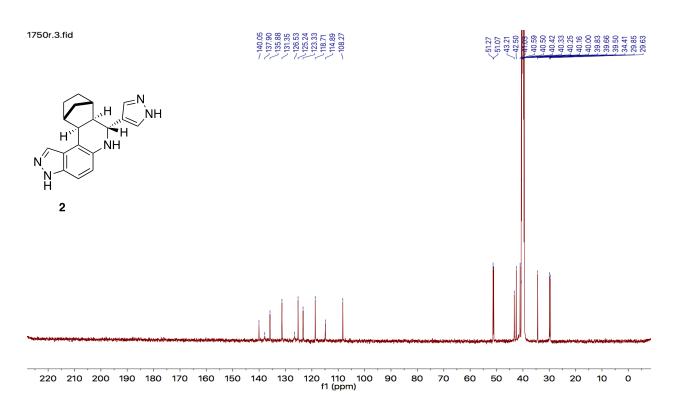


Figure S5: ¹H spectra of compound 3 in methanol-d₄

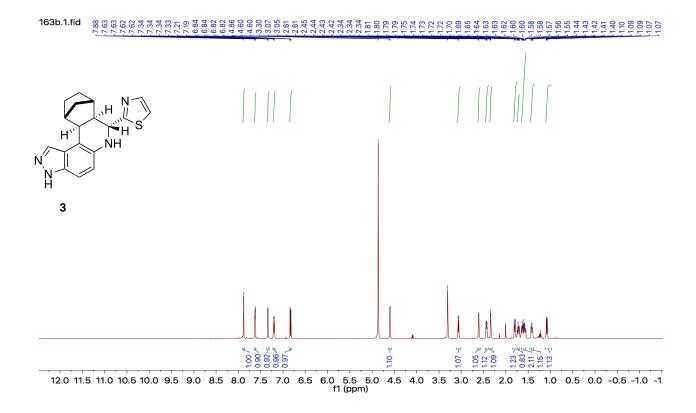


Figure S6: ¹³C spectra of compound 3 in methanol-d₄

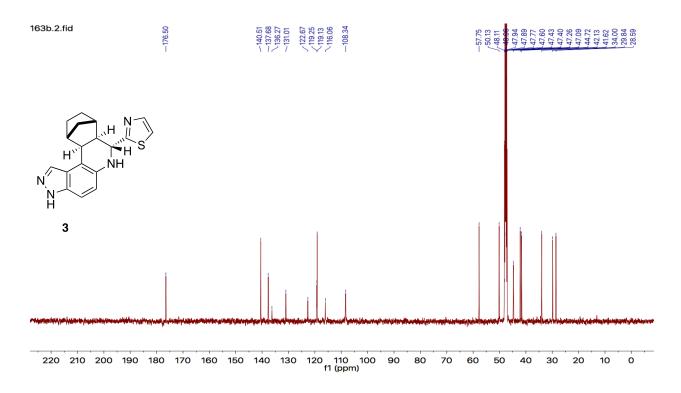


Figure S7: 1 H spectra of compound 4 in methanol- d_{4}

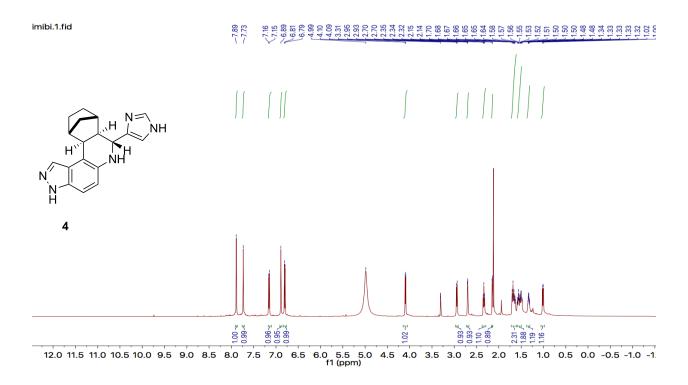


Figure S8: ¹³C spectra of compound 4 in methanol-d₄

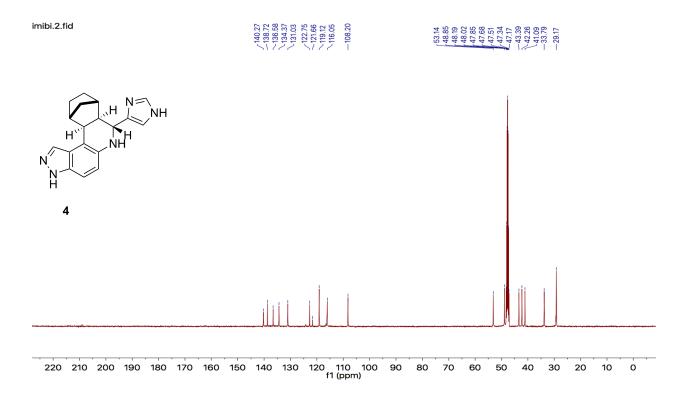


Figure S9: ¹H spectra of compound 5 in methanol-d₄

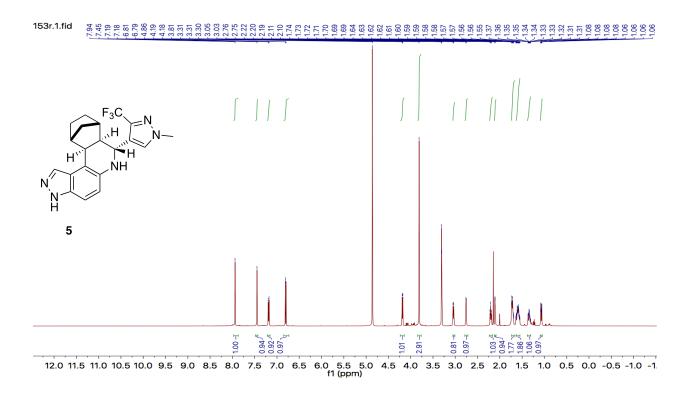


Figure S10: ¹³C spectra of compound 5 in methanol-d₄

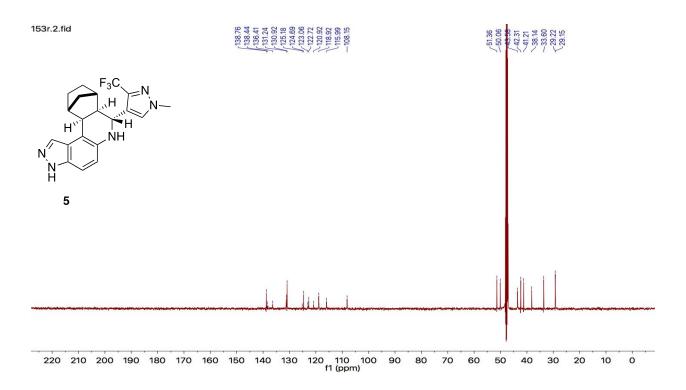


Figure S11: ¹H spectra of compound 6 in DMSO-d₆

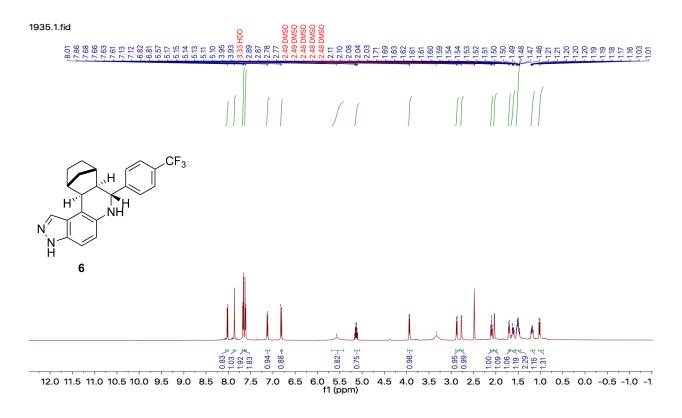


Figure S12: 13 C spectra of compound 6 in DMSO- d_6

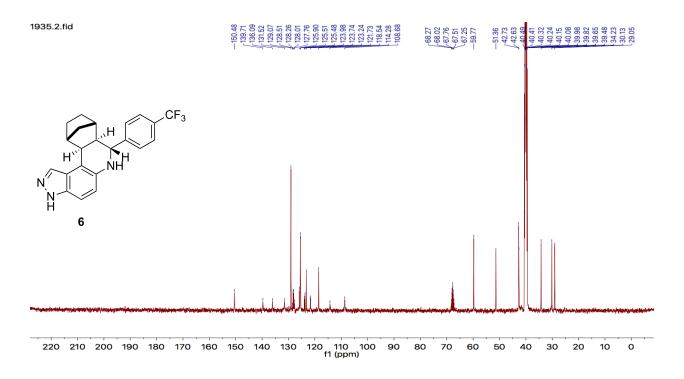


Figure S13: ¹H spectra of compound 7 in DMSO-d₆

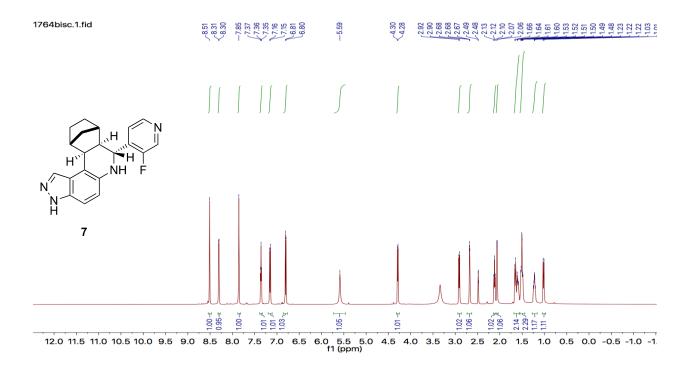


Figure S14: ¹³C spectra of compound 7 in DMSO-d₆

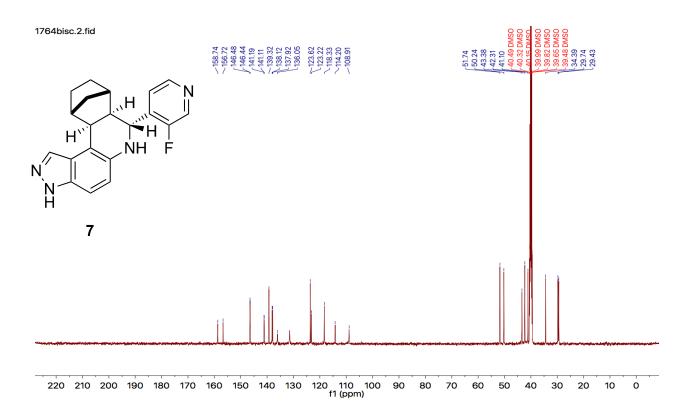


Figure S15: ¹H spectra of compound 8 in methanol-d₄

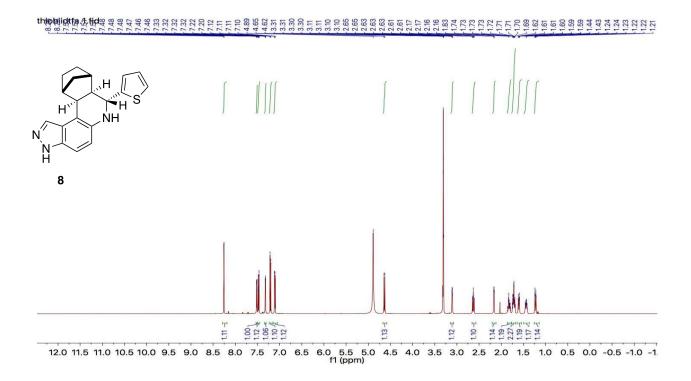


Figure S16: ¹³C spectra of compound 8 in methanol-d₄

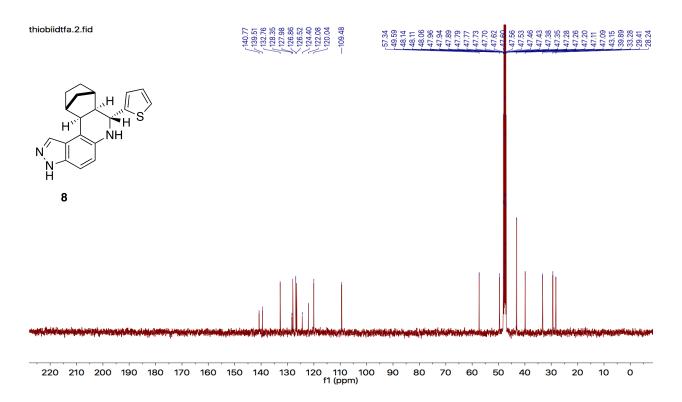


Figure S17: ¹H spectra of compound 9 in methanol-d₄

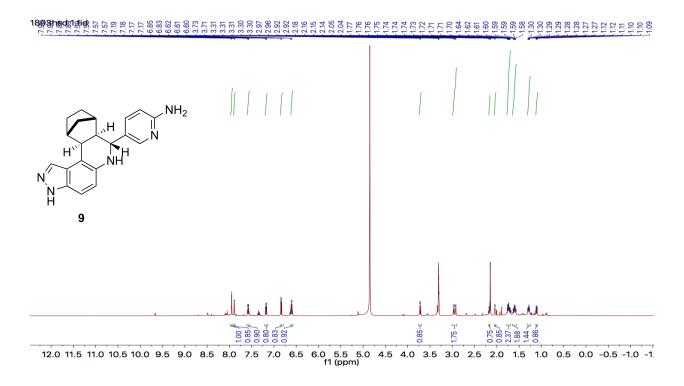


Figure S18: ¹³C spectra of compound 9 in methanol-d₄

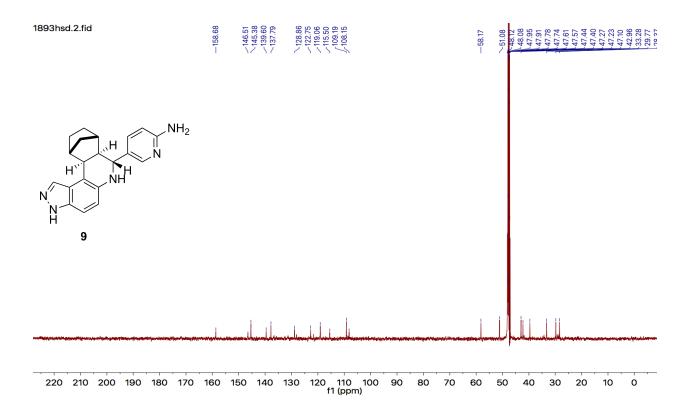
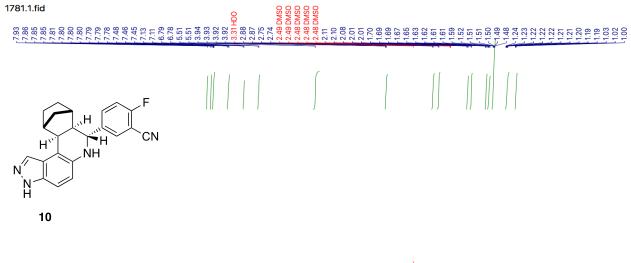


Figure S19: ¹H spectra of compound 10 in DMSO-d₆



12.0 11.5 11.0 10.5 10.0 9.5 9.0 8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 0.0 -0.5 -1.0 -1. f1 (ppm)

Figure S20: ¹³C spectra of compound 10 in DMSO-d₆

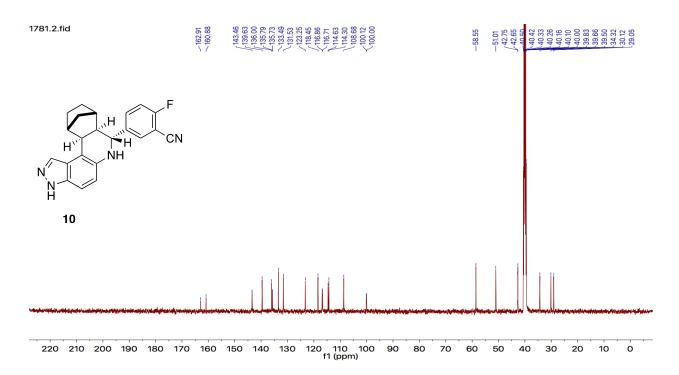


Figure S21: ¹H spectra of compound 11 in DMSO-d₆

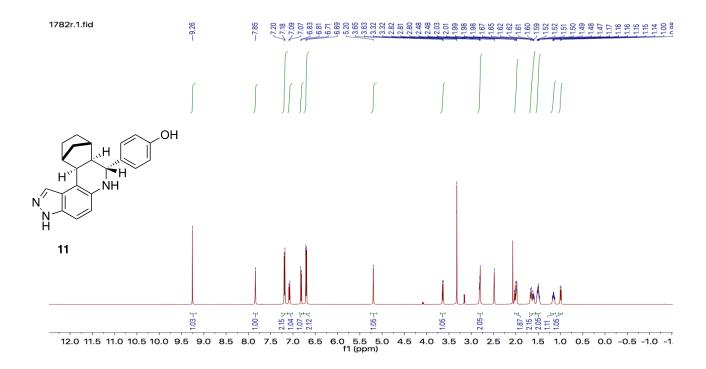


Figure S22: ¹³C spectra of compound 11 in DMSO-d₆

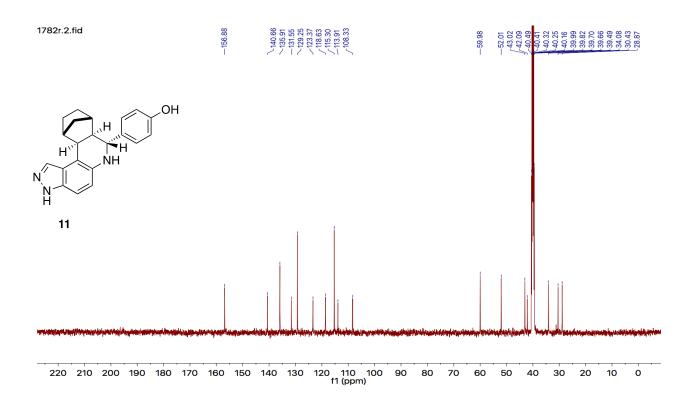


Figure S23: ¹H spectra of compound 12 in DMSO-d₆

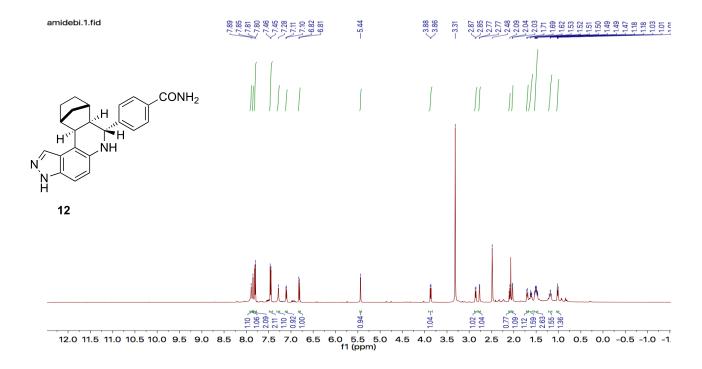


Figure S24: ¹³C spectra of compound 12 in DMSO-d₆

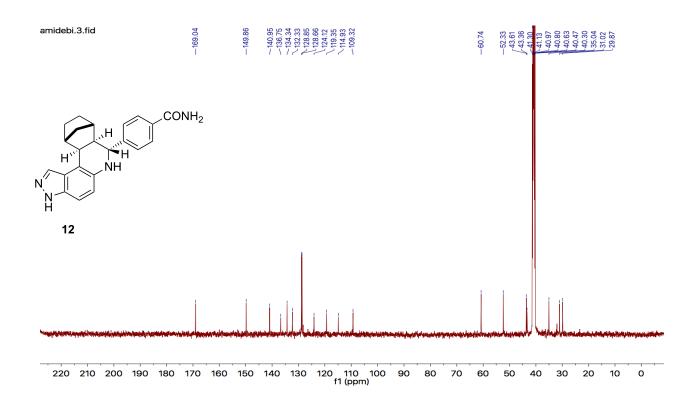


Figure S25: ¹H spectra of compound 13 in DMSO-d₆

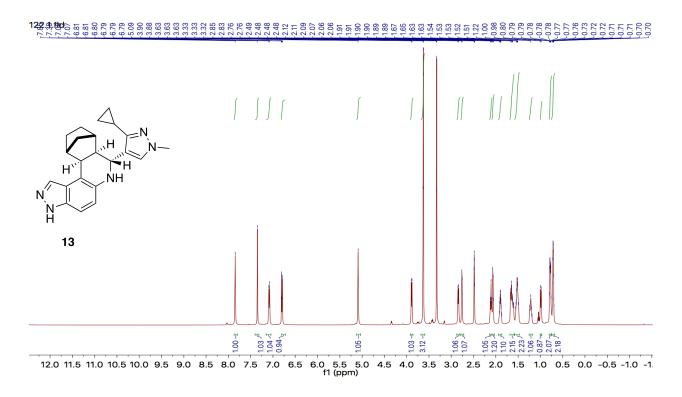


Figure S26: ¹³C spectra of compound 13 in DMSO-d₆

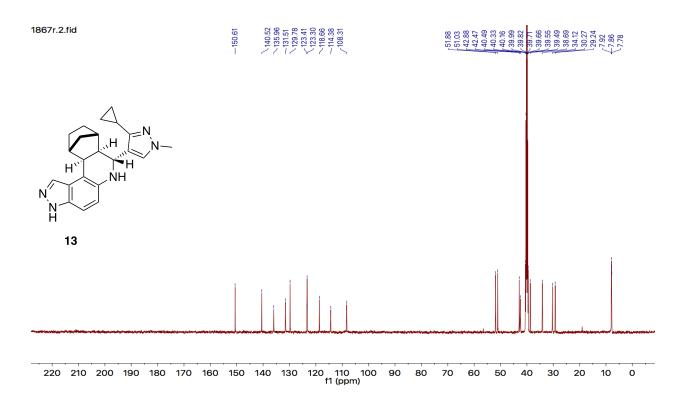


Figure S27: ¹H spectra of compound 14 in DMSO-d₆

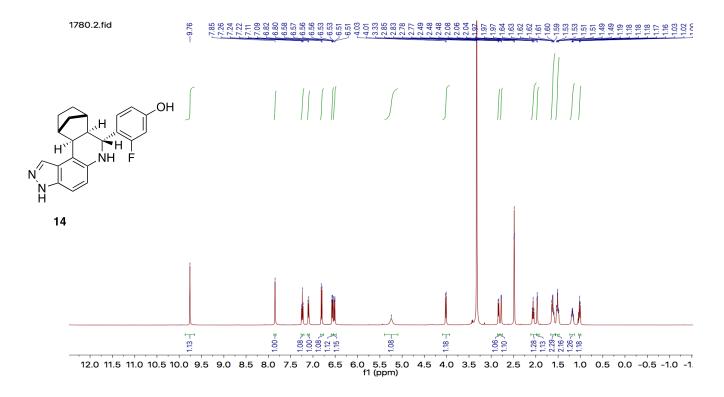


Figure S28: ¹³C spectra of compound 14 in DMSO-d₆

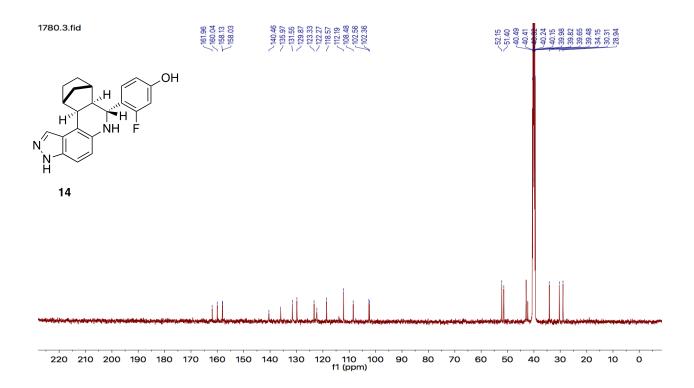


Figure S29: ¹H spectra of compound 15 in DMSO-d₆

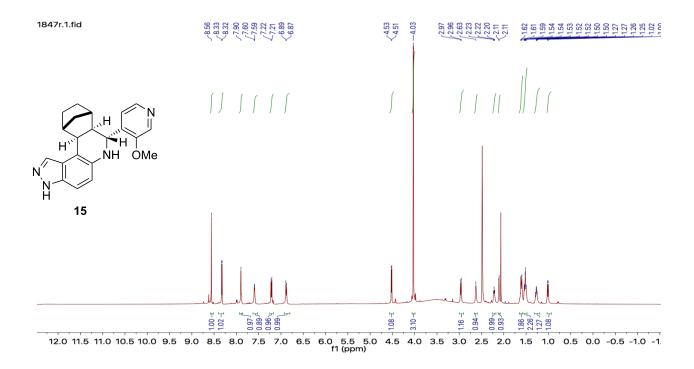


Figure S30: ¹³C spectra of compound 15 in DMSO-d₆

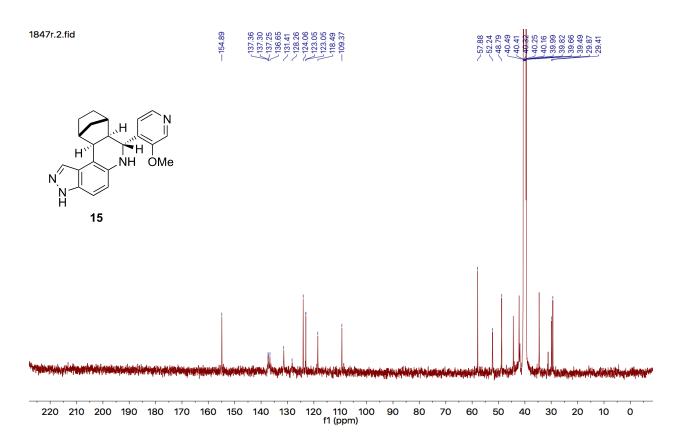


Figure S31: ¹H spectra of compound 16 in DMSO-d₆

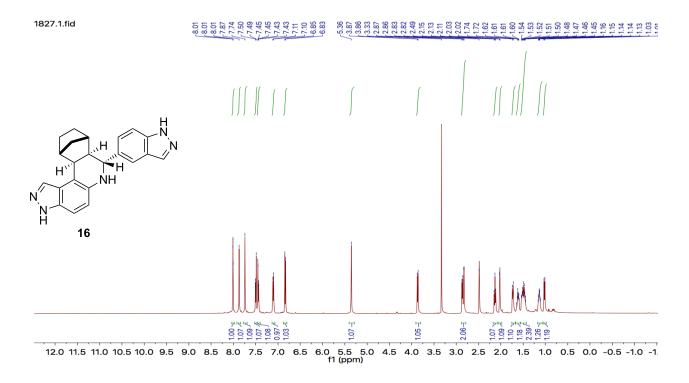


Figure S32: ¹³C spectra of compound 16 in DMSO-d₆

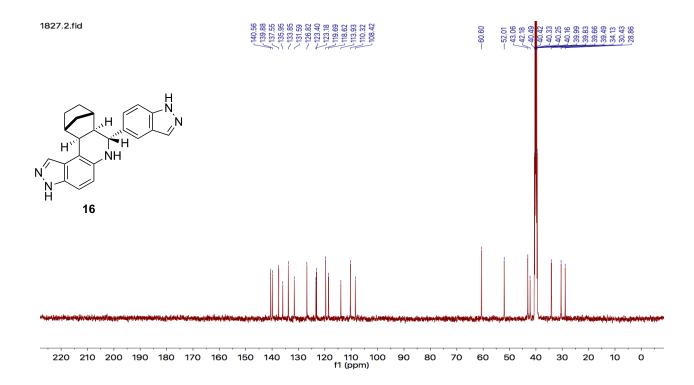


Figure S33: ¹H spectra of compound 17 in DMSO-d₆

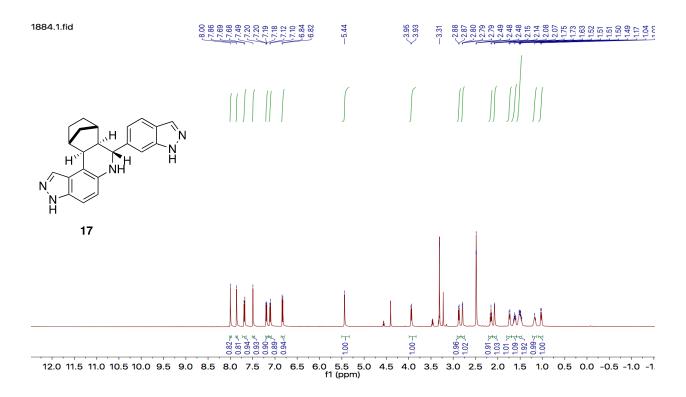


Figure S34: ¹³C spectra of compound 17 in DMSO-d₆

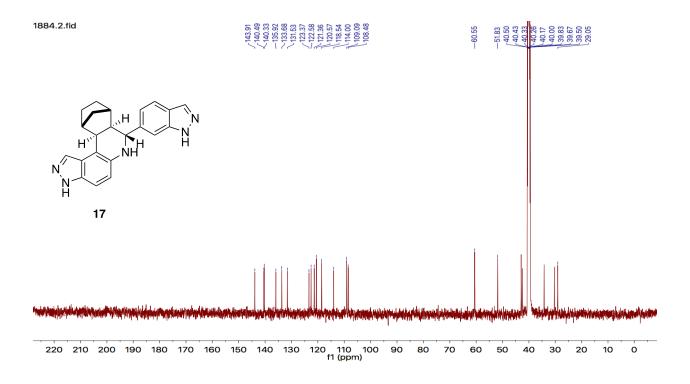


Figure S35: ¹H spectra of compound HSD1787 in DMSO-d₆

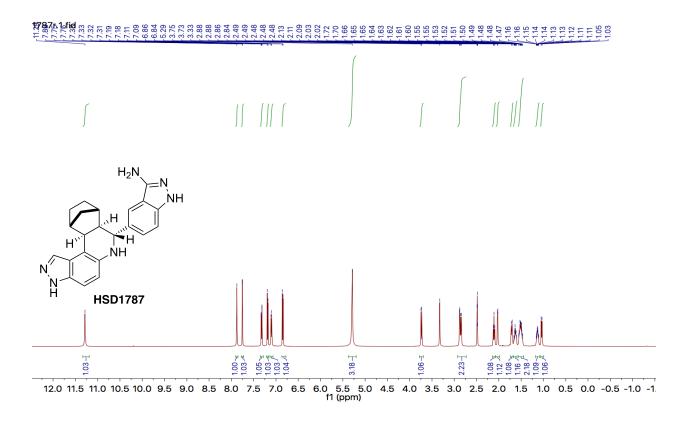


Figure S36: ¹³C spectra of compound HSD1787 in DMSO-d₆

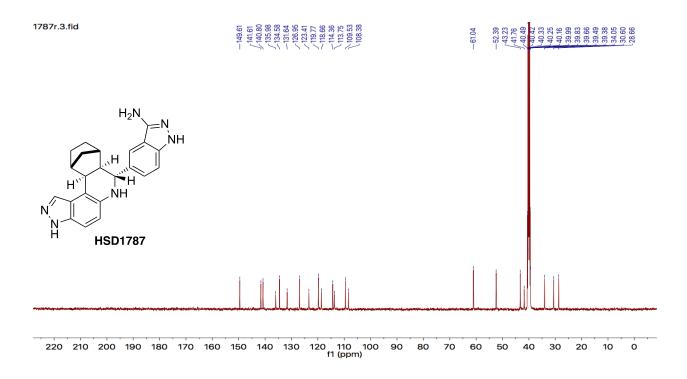


Figure S37: ¹H spectra of compound 18 in DMSO-d₆

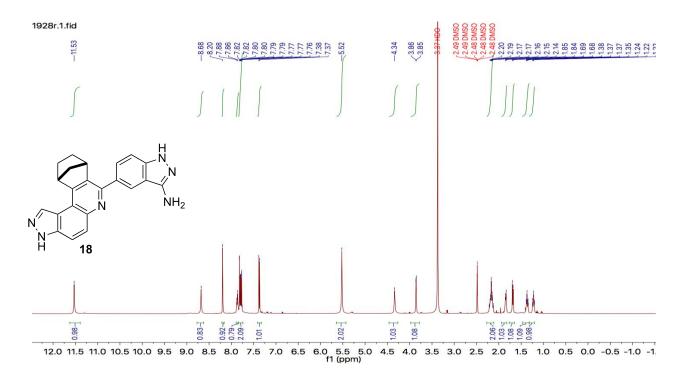


Figure S38: 13 C spectra of compound 18 in DMSO- d_6

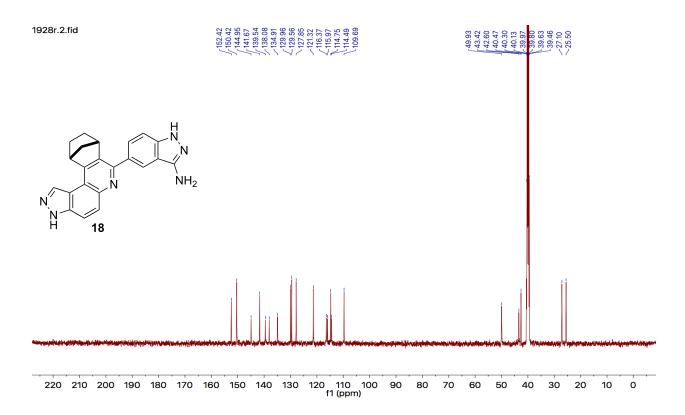


Figure S39: ¹H spectra of compound S-1 in DMSO-d₆

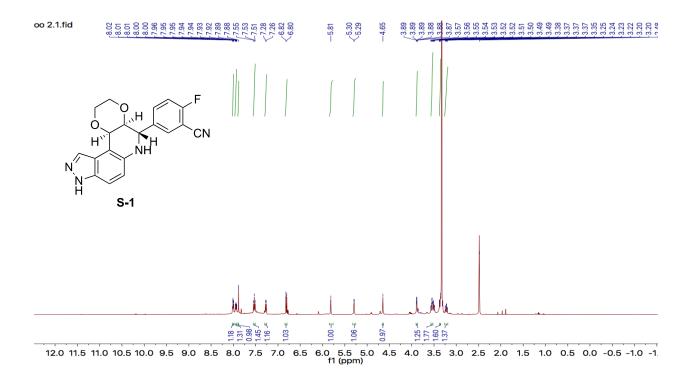


Figure S40: ¹³C spectra of compound S-1 in DMSO-d₆

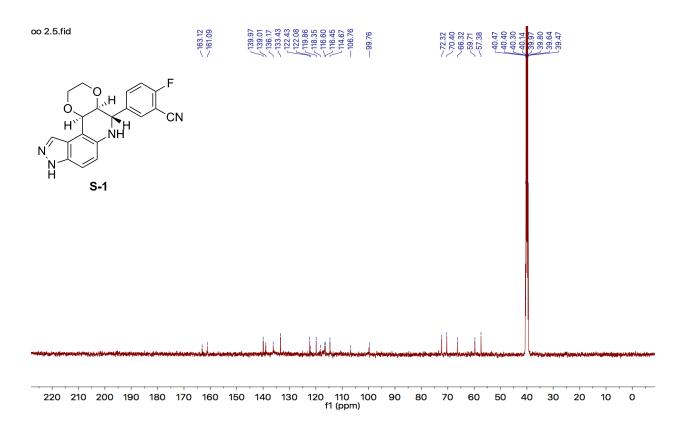


Figure S41: 1 H spectra of compound 19 in DMSO- d_{6}

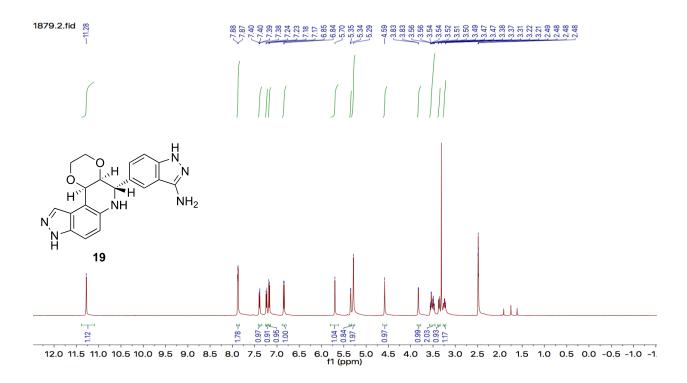


Figure S42: ¹³C spectra of compound 19 in DMSO-d₆

