

Supplementary information for

New Flavonoid – *N,N*-Dibenzyl(*N*-methyl)amine Hybrids: Multi-Target-Directed Agents for Alzheimer’s Disease Endowed with Neurogenic Properties

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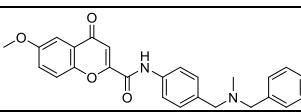
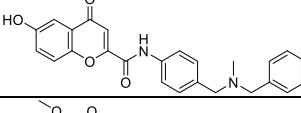
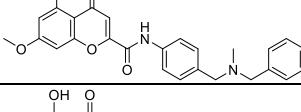
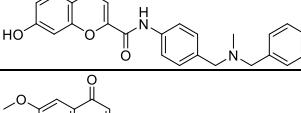
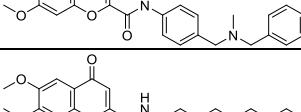
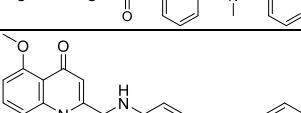
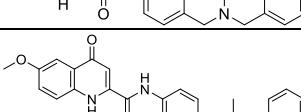
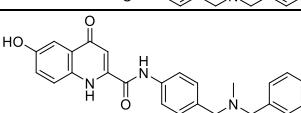
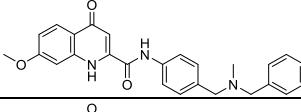
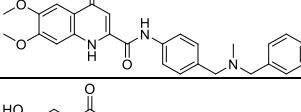
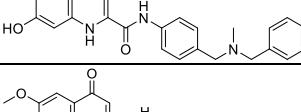
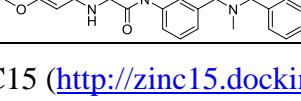
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Medicinal chemistry alerts of hybrids 1-13

Table S1. Medicinal chemistry alerts of hybrids **1-13** according to ZINC15 (<http://zinc15.docking.org/>) [1] and SwissADME (<http://www.swissadme.ch/>) [2] web sites.

	Structure	SMILES	Aggregator ^a	PAINS Alerts ^{a,b}	Brenk Alerts ^b
1		O=C(C1=CC(C2=C(O1)C=CC(OC)=C2)=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
2		O=C(C1=CC(C2=C(O1)C=CC(O)=C2)=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
3		O=C(C1=CC(C2=C(O1)C=C(OC)C=C2OC)=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
4		O=C(C1=CC(C2=C(O1)C=C(O)C=C2O)=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
5		O=C(C1=CC(C2=C(O1)C=C(OC)C(O)=C2)=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
6		O=C(C1=CC(C2=C(O1)C=C(OC)C(O)=C2)=O)NC3=CC=CC(CN(CC4=CC=CC=C4)=CC=C4)C=C3	No	0	0
7		O=C(C(NC1=C2C(OC)=CC=C1)=CC2=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
8		O=C(C(NC1=C2C=C(OC)C=C1)=CC2=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
9		O=C(C(NC1=C2C=C(O)C=C1)=CC2=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
10		O=C(C(NC1=C2C=CC(OC)=C1)=CC2=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
11		O=C(C(NC1=C2C=C(OC)C(OC)=C1)=CC2=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	0
12		O=C(C(NC1=C2C=C(O)C(O)=C1)=CC2=O)NC3=CC=C(CN(CC4=CC=CC=C4)C)C=C3	No	0	1 (catechol)
13		O=C(C(NC1=C2C=C(OC)C(OC)=C1)=CC2=O)NC3=CC=CC(CN(CC4=CC=CC=C4)C)C=C3	No	0	0

^a ZINC15 (<http://zinc15.docking.org/>). ^b SwissADME (<http://www.swissadme.ch/>).

***In silico* study of toxicity of hybrid 6**

Chart S1. Toxicity endpoints predicted at the level of “impossible” for hybrid **6** after *in silico* prediction with Derek Nexus v6.0.1 [3].

5- α Reductase inhibition	Mitochondrial dysfunction
Adrenal gland toxicity	Mutagenicity <i>in vitro</i>
Alpha-2-mu-Globulin nephropathy	Mutagenicity <i>in vivo</i>
Anaphylaxis	Nephrotoxicity
Androgen receptor modulation	Neurotoxicity
Bladder disorders	Non-specific genotoxicity <i>in vitro</i>
Bladder urothelial hyperplasia	Non-specific genotoxicity <i>in vivo</i>
Blood in urine	Occupational asthma
Bone marrow toxicity	Ocular toxicity
Bradycardia	Oestrogen receptor modulation
Carcinogenicity	Oestrogenicity
Cardiotoxicity	Peroxisome proliferation
Cerebral oedema	Phospholipidosis
Chloracne	Photo-induced chromosome damage <i>in vitro</i>
Chromosome damage <i>in vitro</i>	Photo-induced non-specific genotoxicity <i>in vitro</i>
Chromosome damage <i>in vivo</i>	Photo-induced non-specific genotoxicity <i>in vivo</i>
Cumulative effect on white cell count and immunology	Photoallergenicity
Cyanide-type effects	Photocarcinogenicity
Developmental toxicity	Photomutagenicity <i>in vitro</i>
Glucocorticoid receptor agonism	Phototoxicity
Hepatotoxicity	Pulmonary toxicity
High acute toxicity	Respiratory sensitisation
Irreversible cholinesterase inhibition	Splenotoxicity
Irritation (of the gastrointestinal tract)	Teratogenicity
Kidney disorders	Testicular toxicity
Kidney function-related toxicity	Thyroid toxicity
Lachrymation	Uncoupler of oxidative phosphorylation
Methaemoglobinaemia	Urolithiasis

¹H-NMR, ¹³C-NMR and HRMS data of selected flavonoid – DBMA hybrids

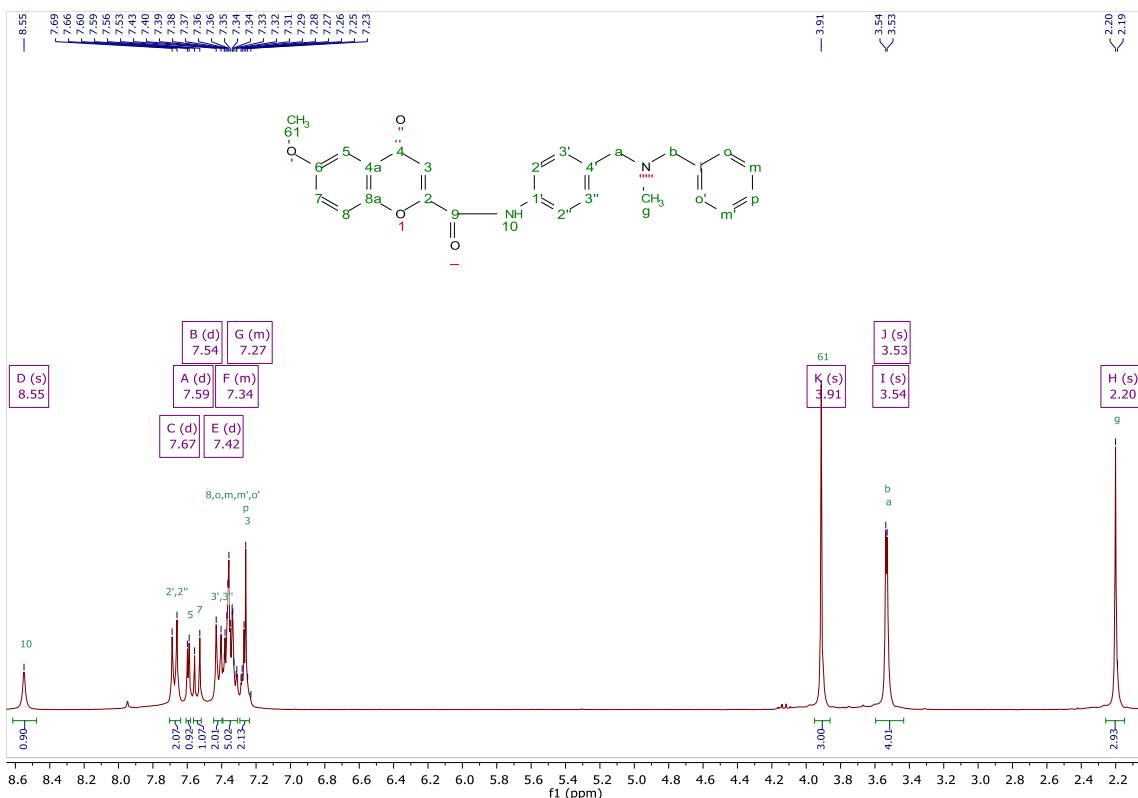


Figure S1. ¹H NMR of **1** in CDCl₃

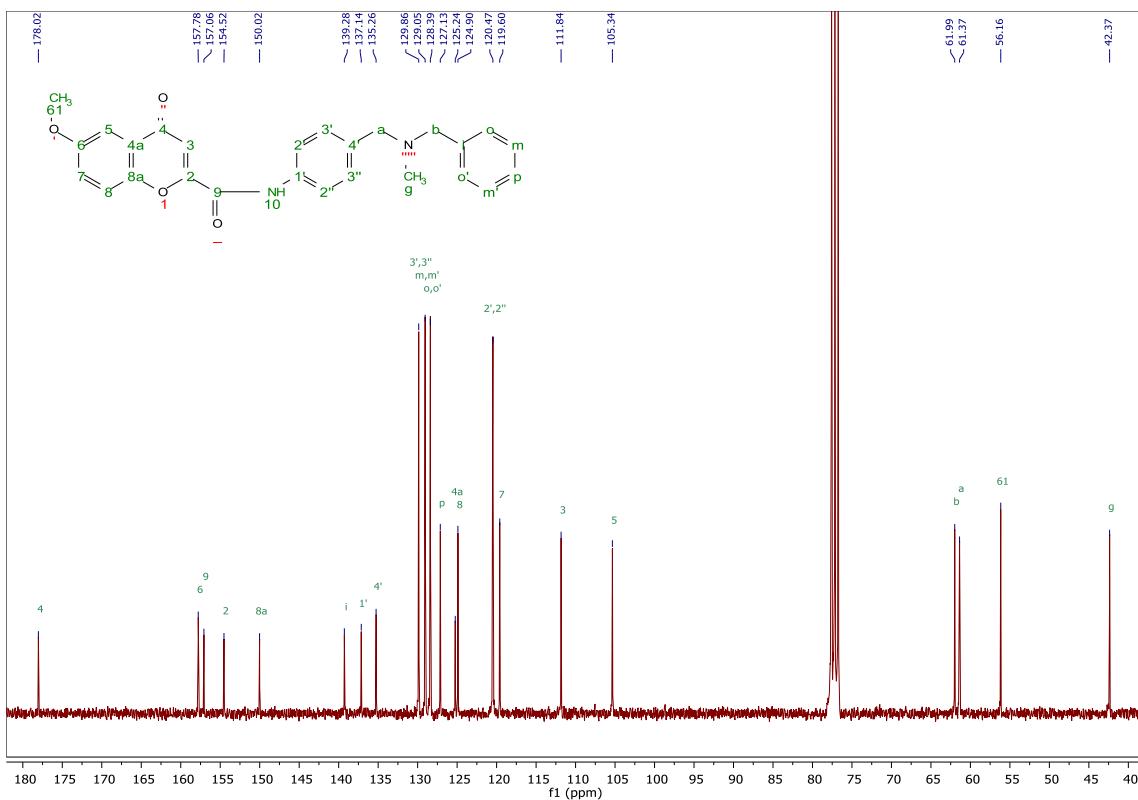


Figure S2. ¹³C NMR of **1** in CDCl₃

Data File	5521_mev_303_01.d	Sample Name	mev_303
Sample Type	Sample	Position	Vial 22
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Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Success
DA Method	Default.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C26 H24 N2 O4	1.166	428.1754	29865	C26 H24 N2 O4	428.1736	4.17

Compound Label	RT	Algorithm	Mass
Cpd 1: C26 H24 N2 O4	1.166	Find By Formula	428.1754

MS Zoomed Spectrum

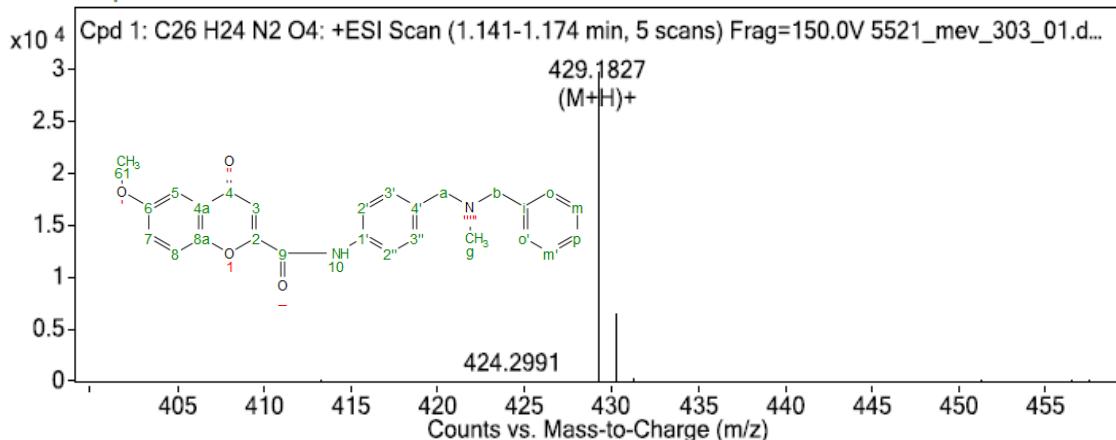


Figure S3. HRMS for hybrid 1

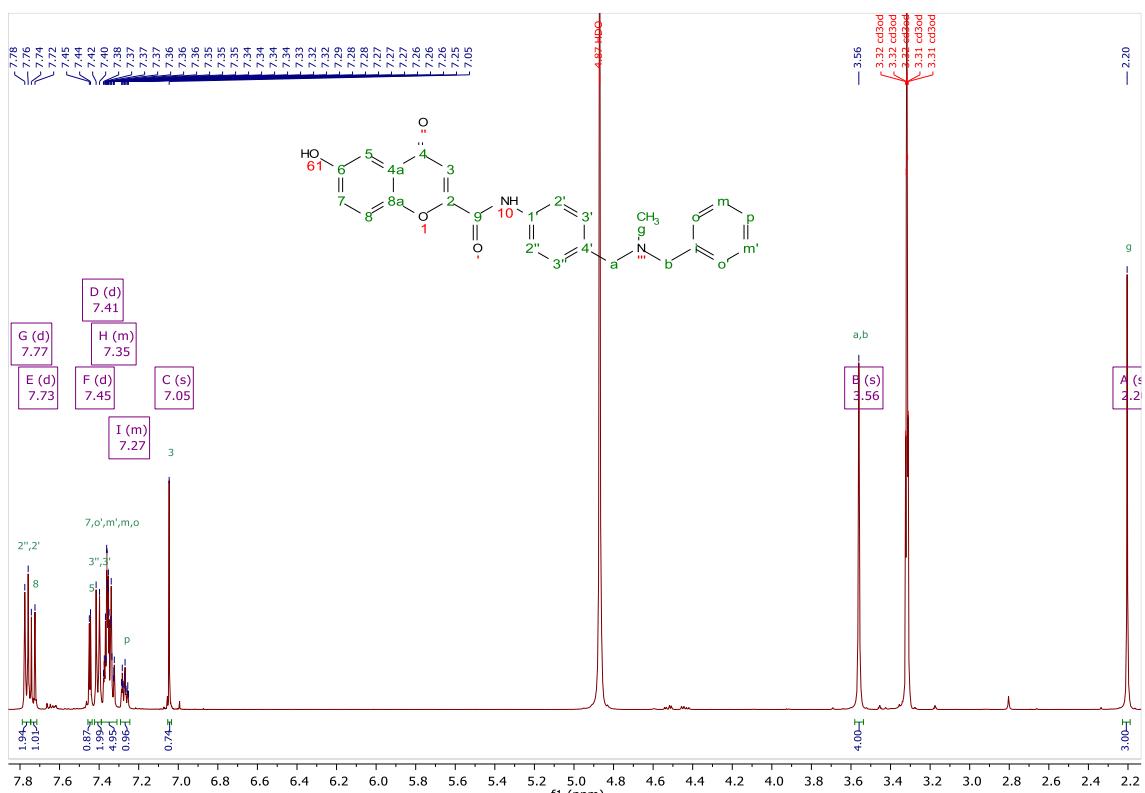


Figure S4. ¹H NMR of 2 in MeOD

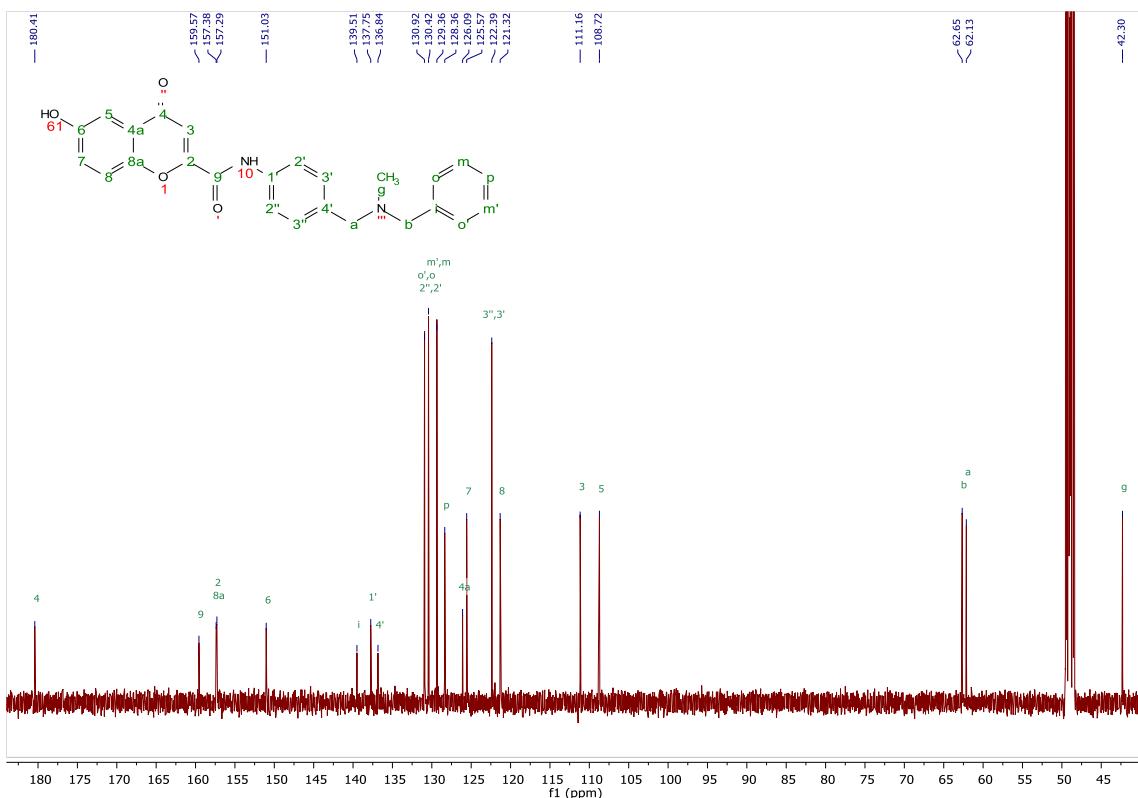


Figure S5. ^{13}C NMR of **2** in MeOD

Data File	6347_mev_419_01.d	Sample Name	mev_419
Sample Type	Sample	Position	Vial 7
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Success
DA Method	01_busqueda.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C25 H22 O4 N2	1.242	414.1583	492443	C25 H22 O4 N2	414.158	0.91

Compound Label	RT	Algorithm	Mass
Cpd 1: C25 H22 O4 N2	1.242	Find By Formula	414.1583

MS Zoomed Spectrum

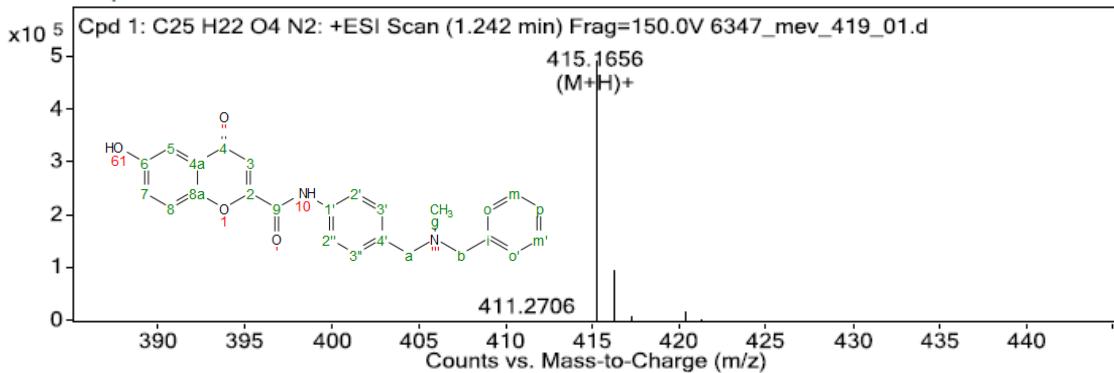


Figure S6. HRMS for compound **2**

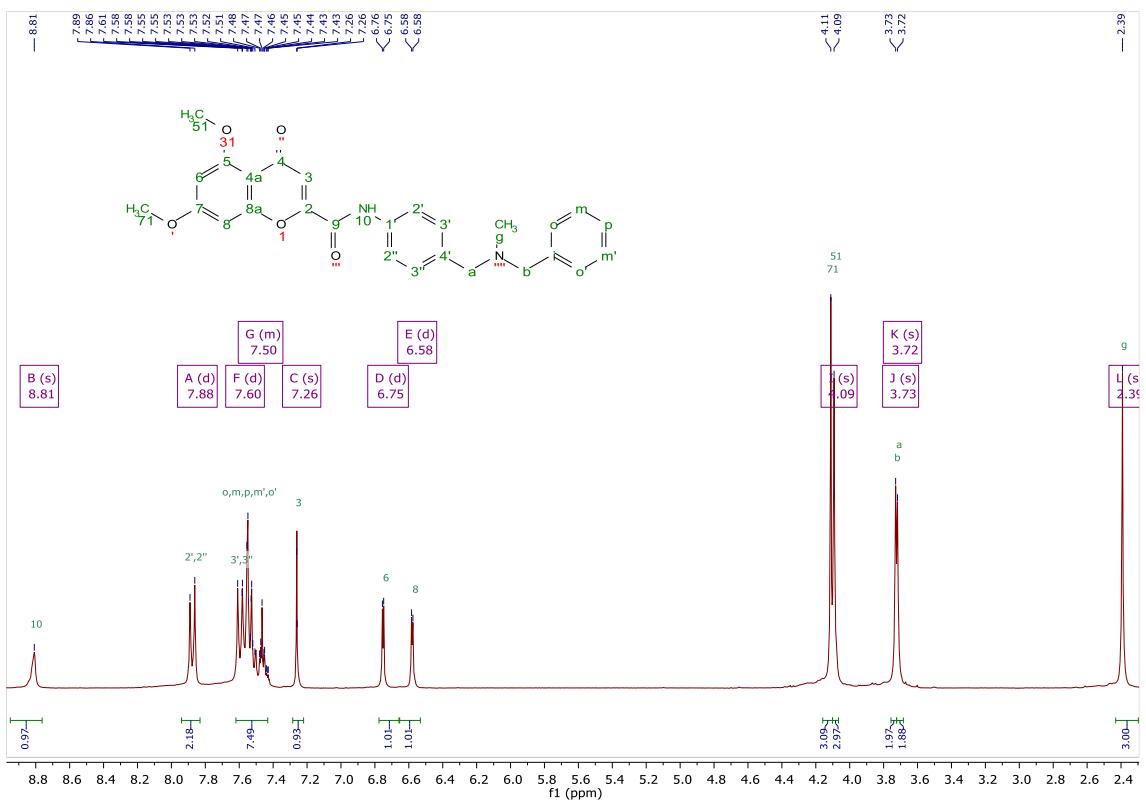


Figure S7. ^1H NMR of **3** in CDCl_3

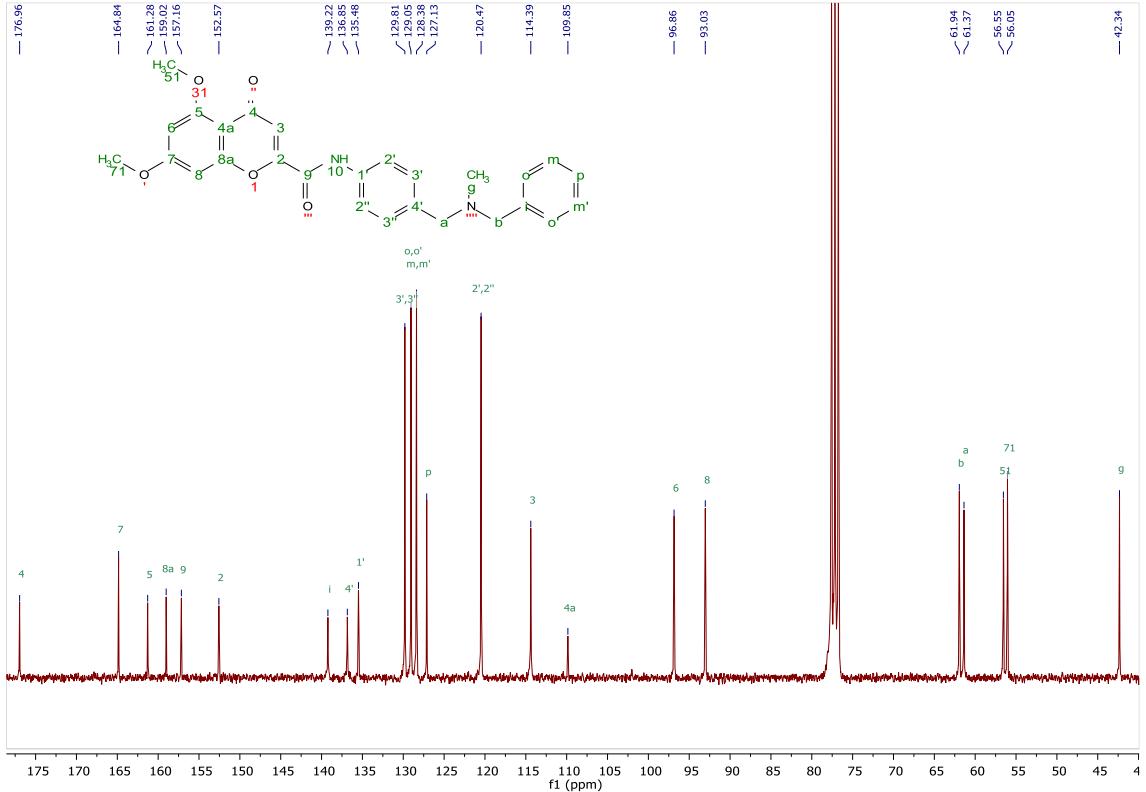


Figure S8. ^{13}C NMR of **3** in CDCl_3

Data File	5522_mev_304_01.d	Sample Name	mev_304
Sample Type	Sample	Position	Vial 23
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	All Ions Missed
DA Method	Default.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C27 H26 N2 O5	1.15	458.1863	77121	C27 H26 N2 O5	458.1842	4.64

Compound Label	RT	Algorithm	Mass
Cpd 1: C27 H26 N2 O5	1.15	Find By Formula	458.1863

MS Zoomed Spectrum

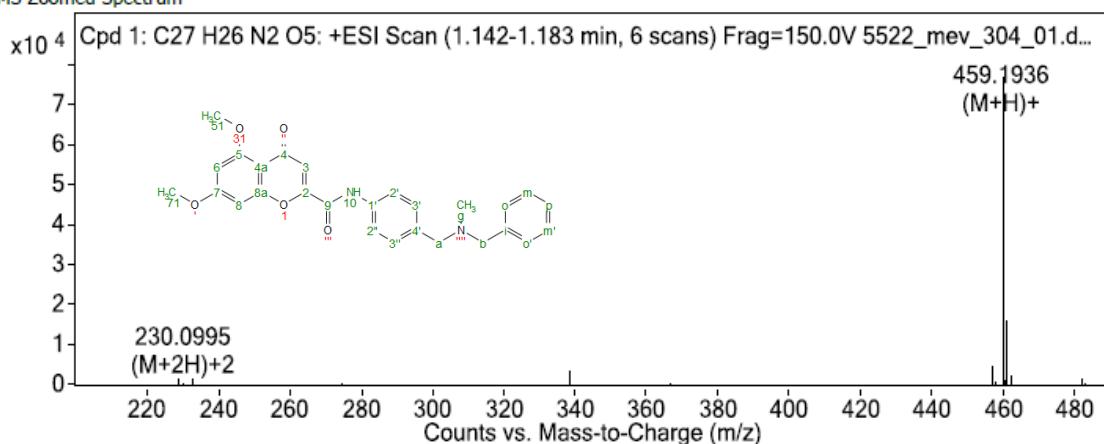


Fig S9. HRMS for compound 3

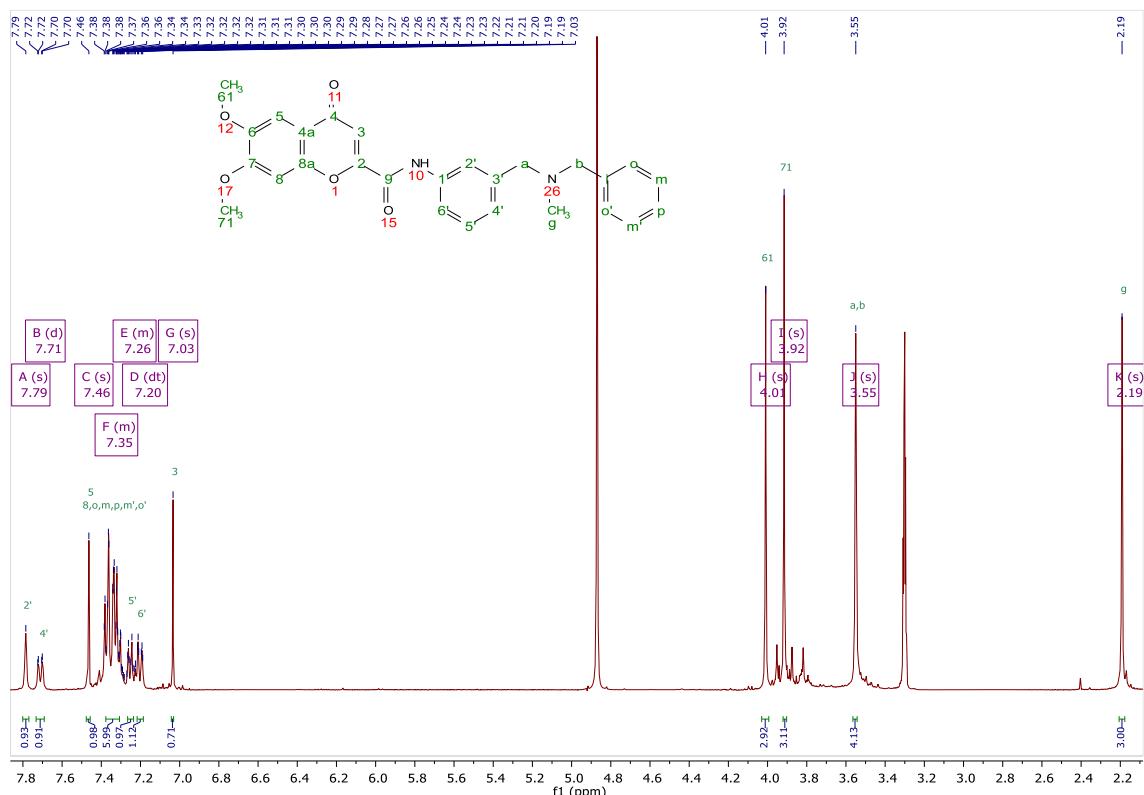


Figure S10. ¹H NMR of 6 in MeOD

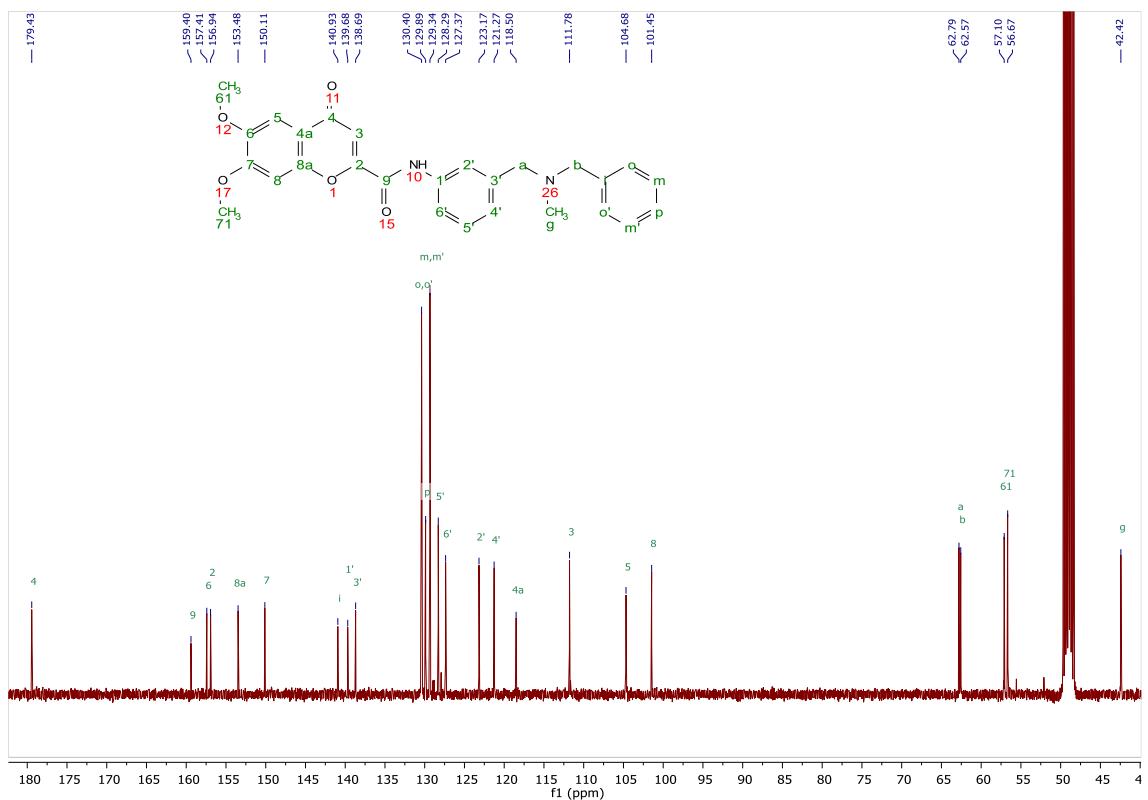


Figure S11. ^{13}C NMR of **6** in MeOD

Data File	7789_mev_658_01.d	Sample Name	mev_658
Sample Type	Sample	Position	Vial 3
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Some Ions Missed
DA Method	defecto.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C27 H26 N2 O5	0.689	458.18405	182379	C27 H26 N2 O5	458.18417	-0.27

Compound Label	RT	Algorithm	Mass
Cpd 1: C27 H26 N2 O5	0.689	Find By Formula	458.18405

MS Zoomed Spectrum

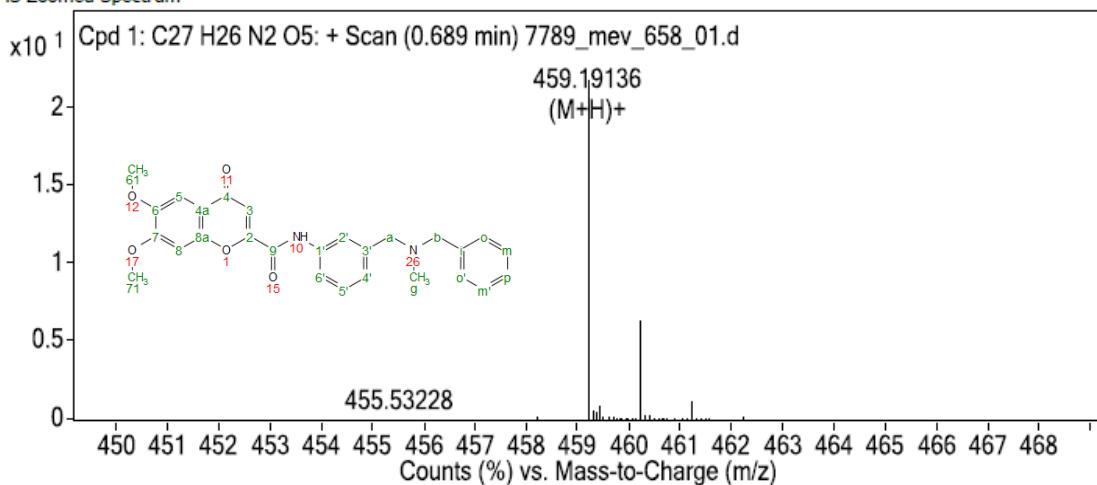


Figure S12. HRMS for compound **6**

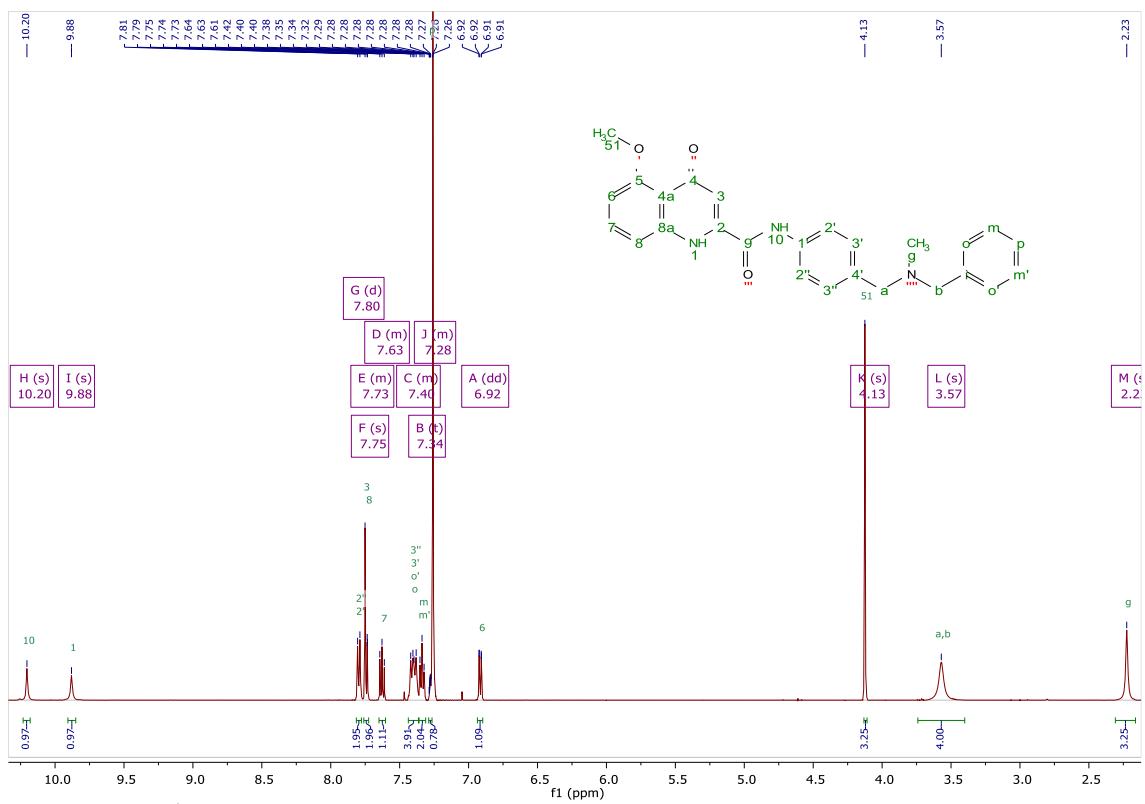


Figure S13. ^1H NMR of 7 in CDCl_3

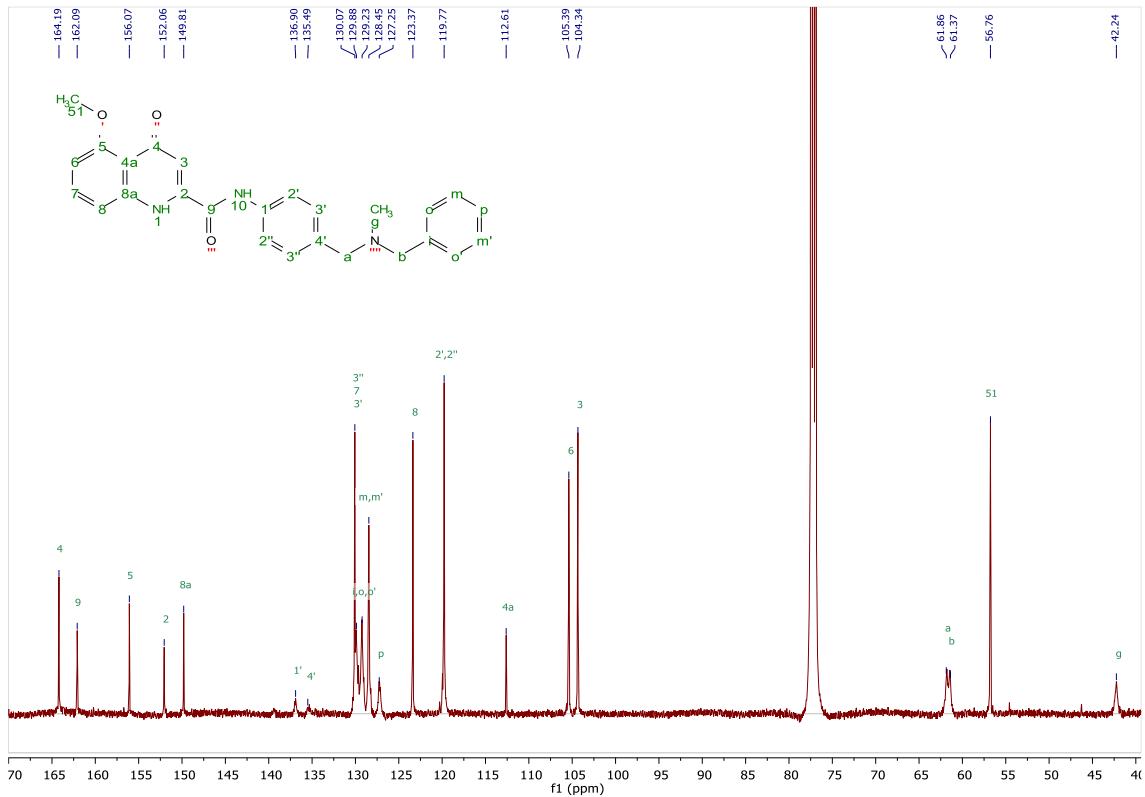


Figure S14. ^{13}C NMR of 7 in CDCl_3

Data File	7618_mev_454_f1_01.d	Sample Name	mev_454_f1
Sample Type	Sample	Position	Vial 2
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Some Ions Missed
DA Method	defecto.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C26 H25 N3 O3	0.607	427.19001	618453	C26 H25 N3 O3	427.18959	0.98

Compound Label	RT	Algorithm	Mass
Cpd 1: C26 H25 N3 O3	0.607	Find By Formula	427.19001

MS Zoomed Spectrum

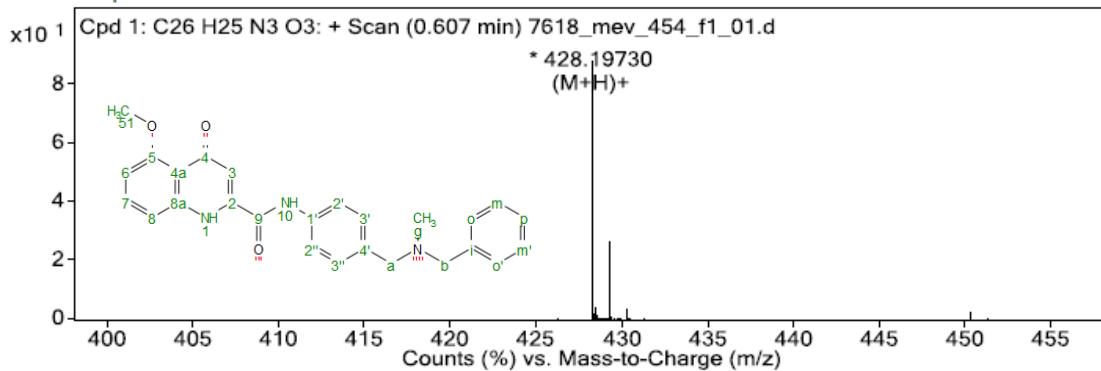


Figure S15. HRMS for Compound 7

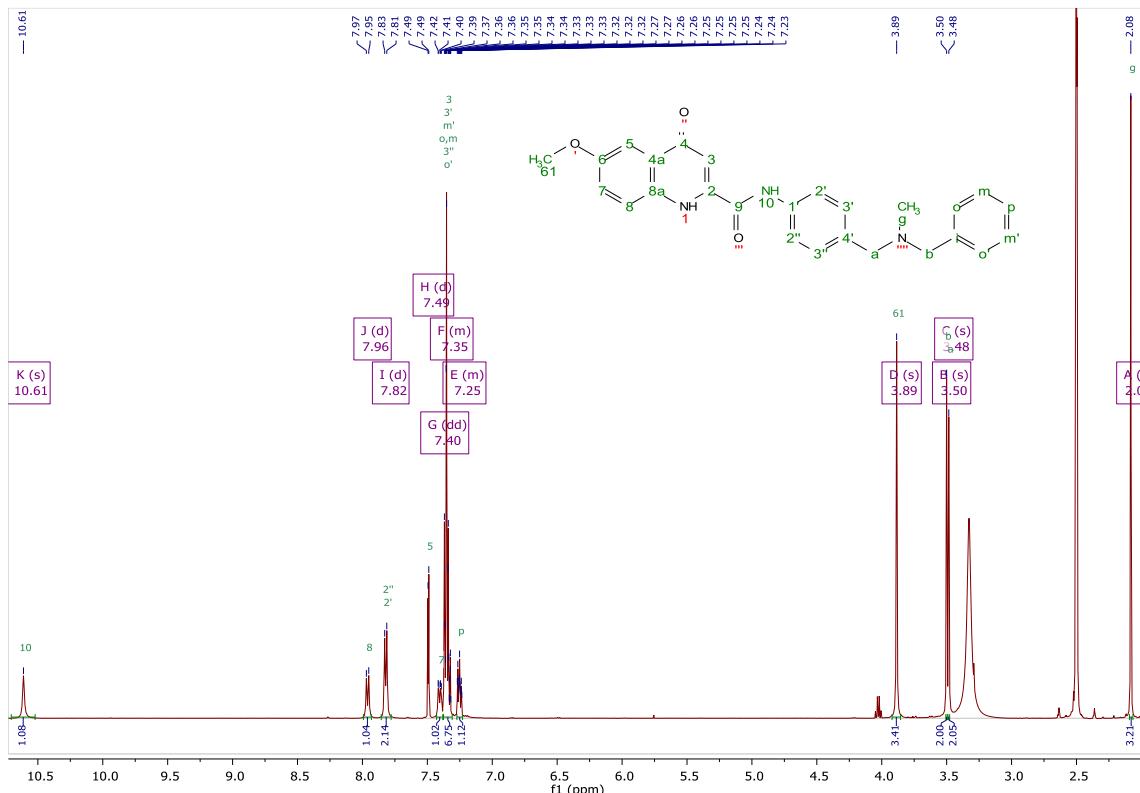


Figure S16. ¹H NMR of 8 in DMSO-d6

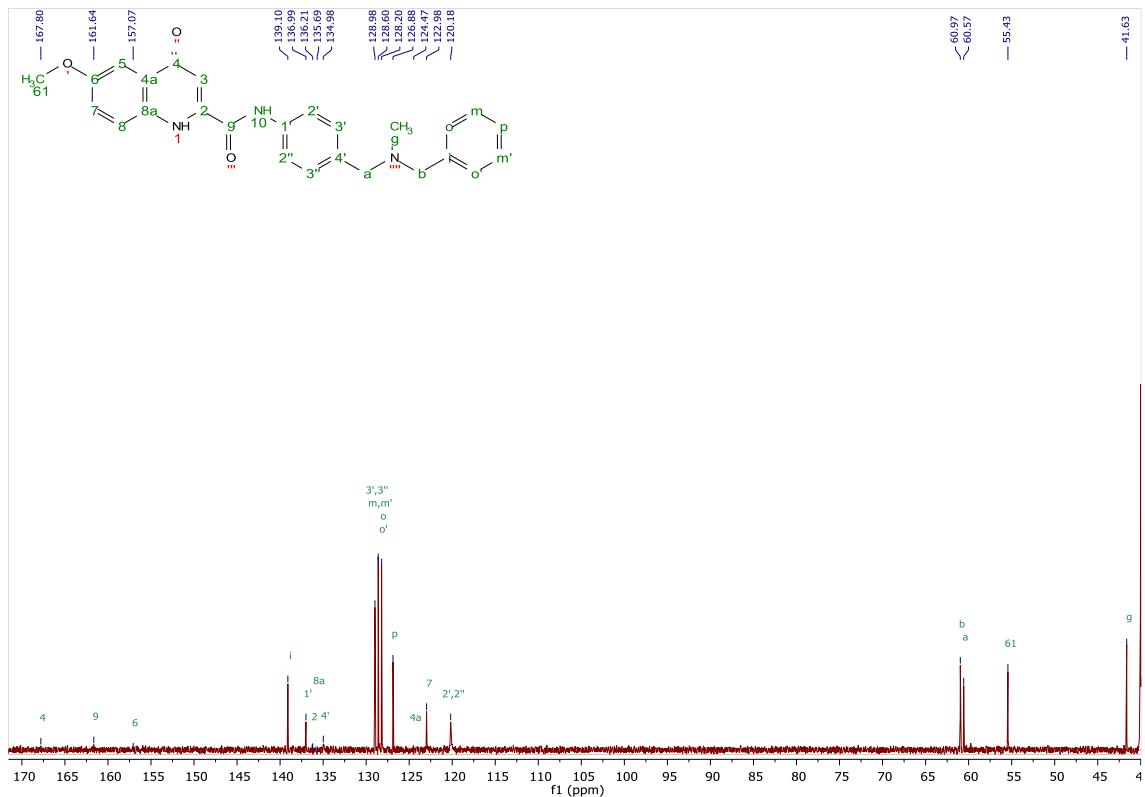


Figure S17. ^{13}C NMR of **8** in DMSO-d_6

Data File	6355_mev_456_01.d	Sample Name	mev_456
Sample Type	Sample	Position	Vial 15
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Success
DA Method	01_busqueda.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C26 H25 O3 N3	0.56	427.1905	344225	C26 H25 O3 N3	427.1896	2.13

Compound Label	RT	Algorithm	Mass
Cpd 1: C26 H25 O3 N3	0.56	Find By Formula	427.1905

MS Zoomed Spectrum

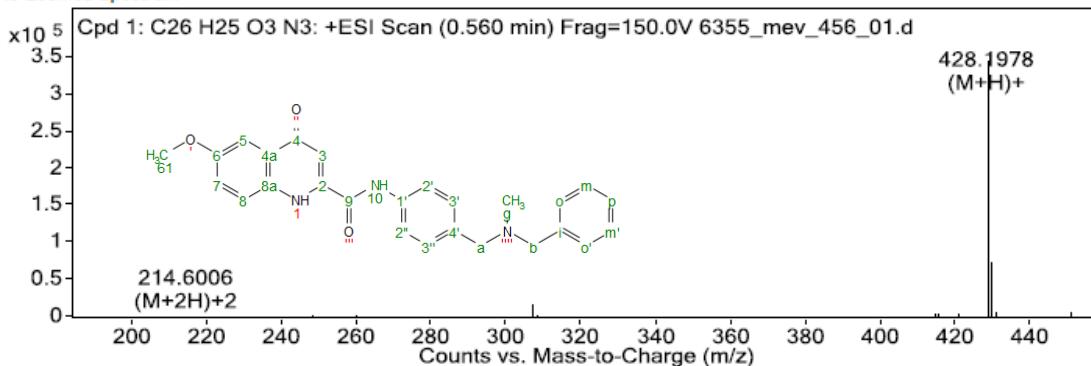


Figure S18. HRMS for compound **8**

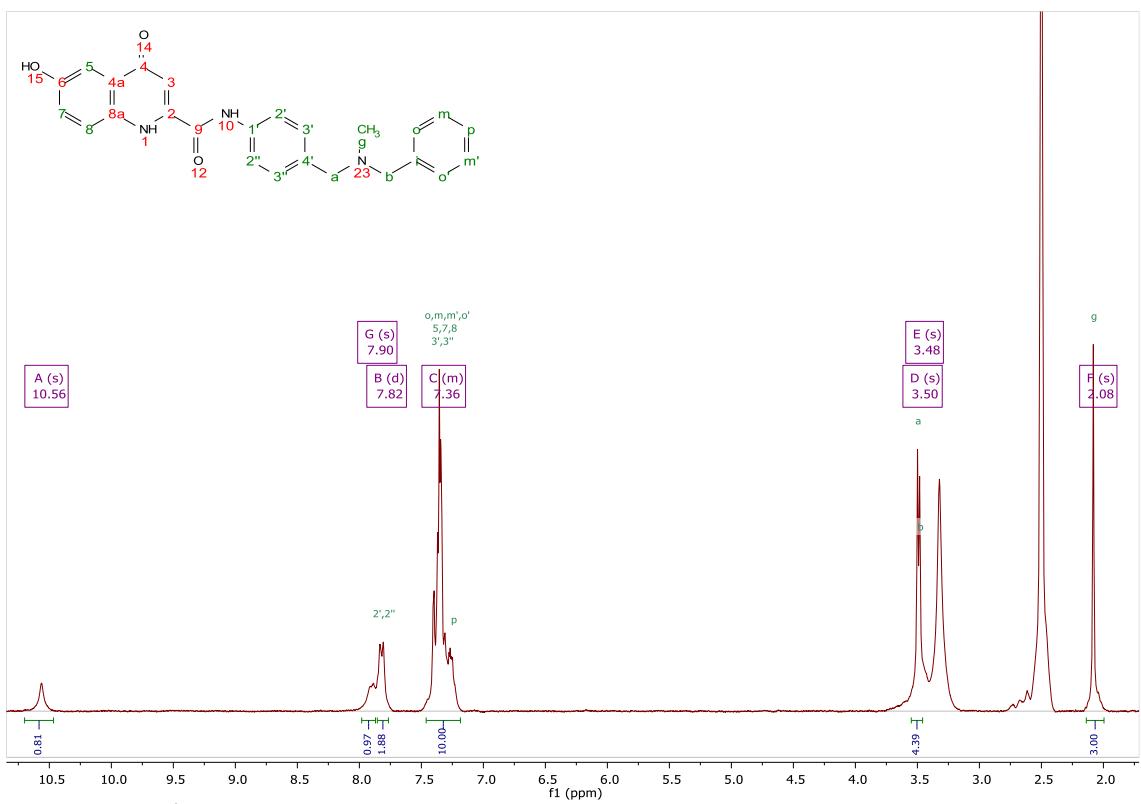


Figure S19. ¹H NMR of **9** in DMSO-d₆

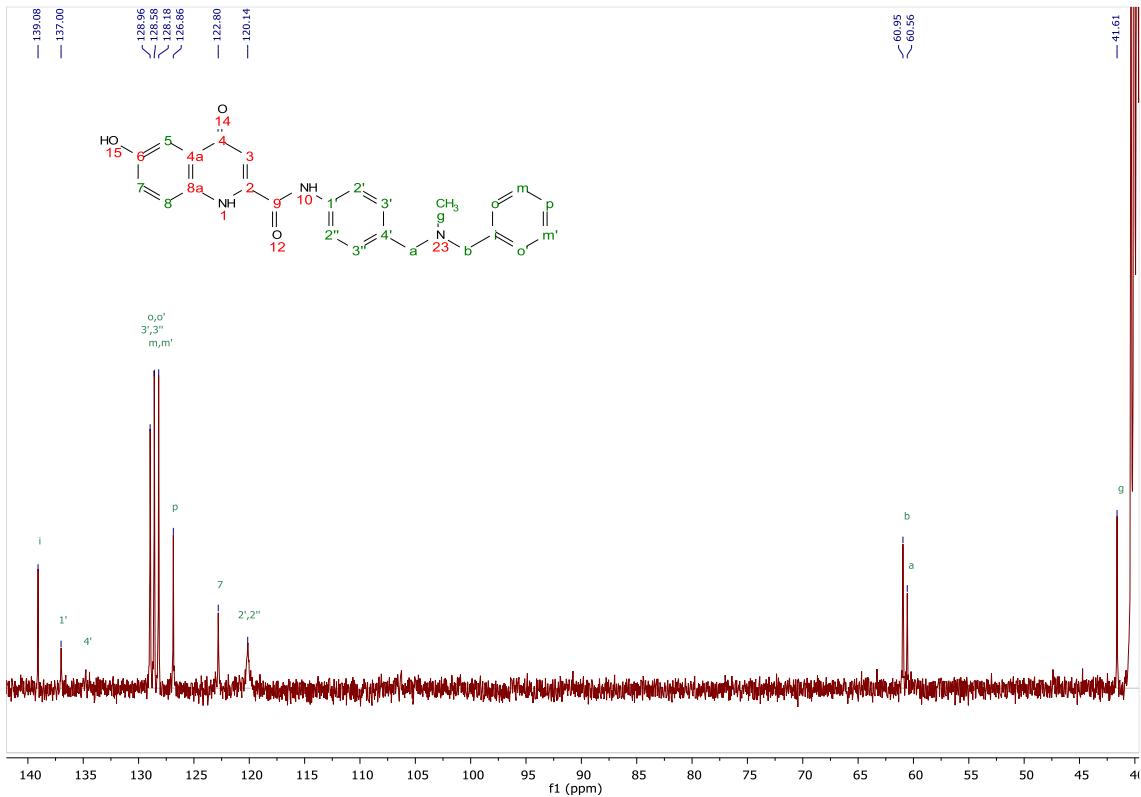


Figure S20. ¹³C NMR of **9** in DMSO-d₆

Data File	6354_mev_458_01.d	Sample Name	mev_458
Sample Type	Sample	Position	Vial 14
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Success
DA Method	01_busqueda.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C25 H23 O3 N3	0.883	413.1751	349781	C25 H23 O3 N3	413.1739	2.84

Compound Label	RT	Algorithm	Mass
Cpd 1: C25 H23 O3 N3	0.883	Find By Formula	413.1751

MS Zoomed Spectrum

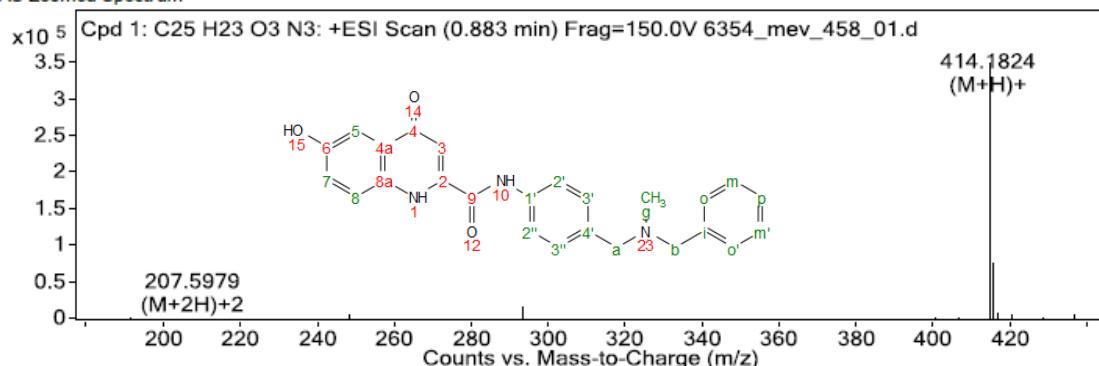


Figure S21. HRMS for compound 9

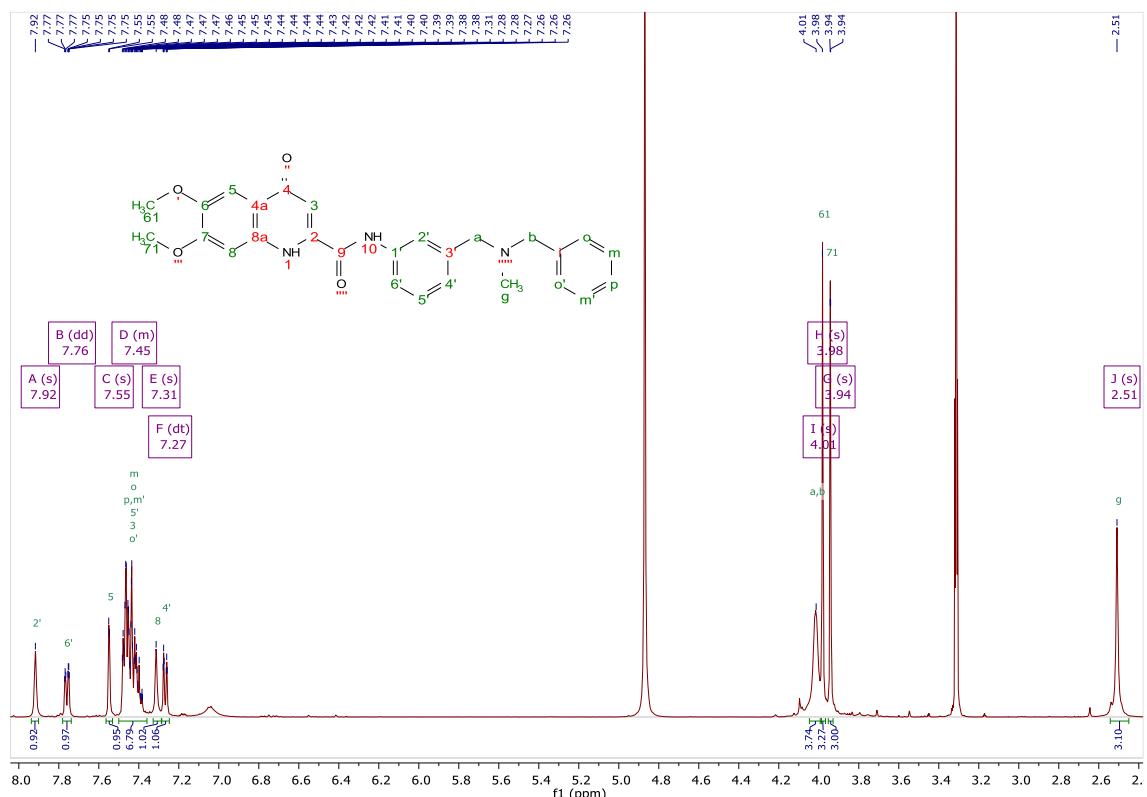


Figure S22. ¹H NMR of 13 in MeOD

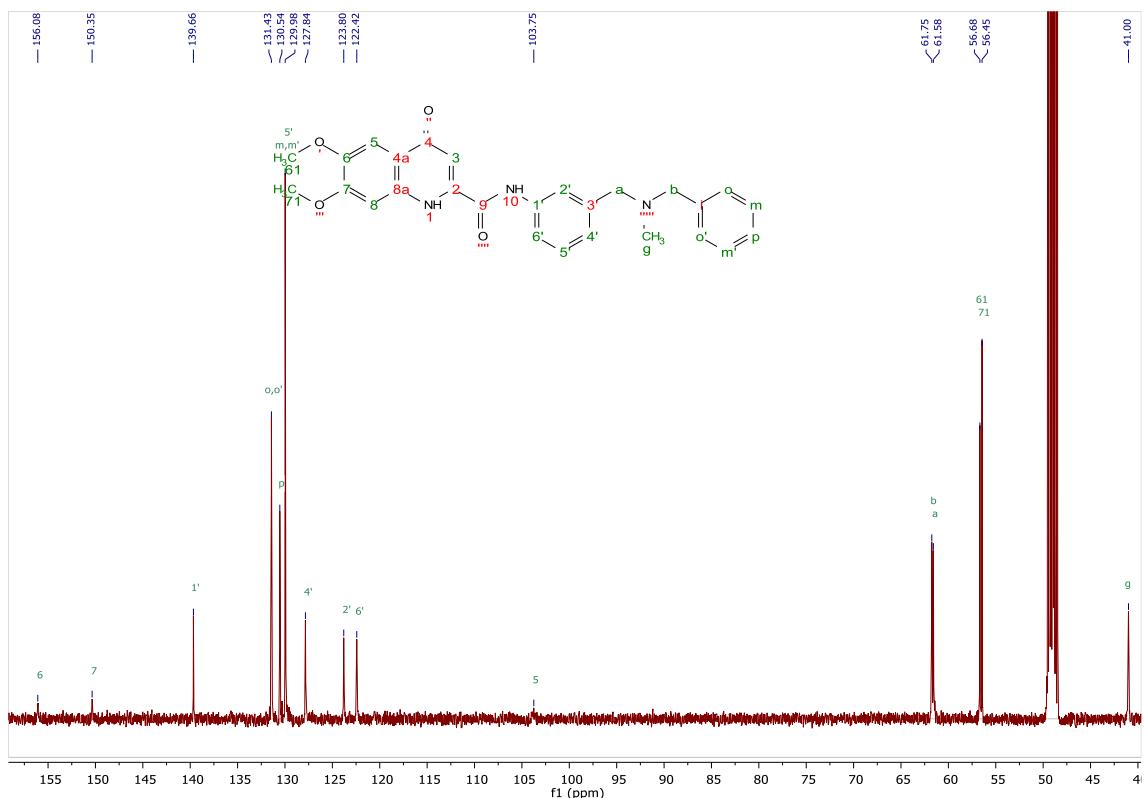


Figure S23. ^{13}C NMR of **13** in MeOD

Data File	7788_mev_657_01.d	Sample Name	mev_657
Sample Type	Sample	Position	Vial 2
Instrument Name	Instrument 1	User Name	
Acq Method	ESI_ACN_75_pos.m	IRM Calibration Status	Success
DA Method	defecto.m	Comment	

Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)
Cpd 1: C27 H27 N3 O4	0.707	457.20024	180025	C27 H27 N3 O4	457.20016	0.17

Compound Label	RT	Algorithm	Mass
Cpd 1: C27 H27 N3 O4	0.707	Find By Formula	457.20024

MS Zoomed Spectrum

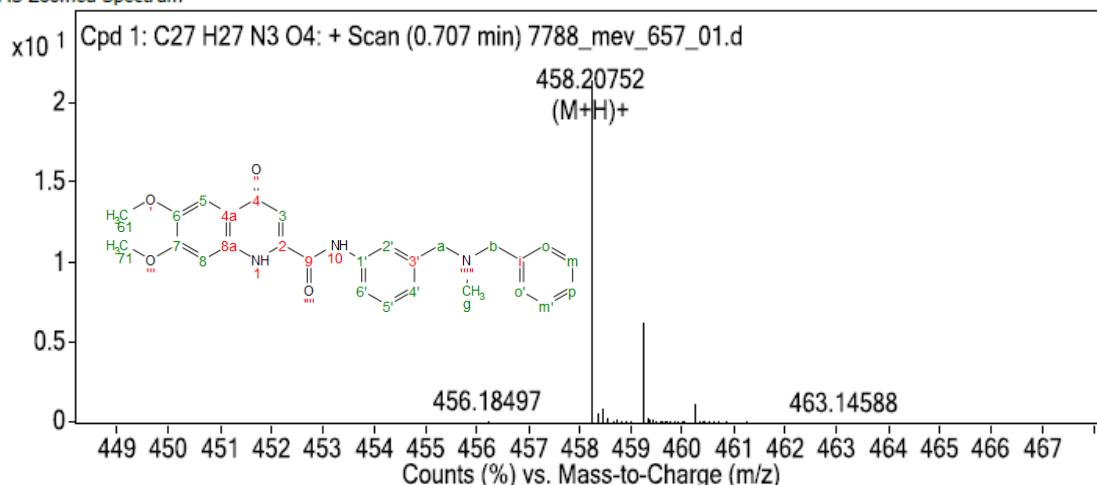


Figure S24. HRMS for compound **13**

References

- [1] T. Sterling, J.J. Irwin, ZINC 15--Ligand Discovery for Everyone, *J. Chem. Inf. Model.* 55 (2015) 2324-2337.
- [2] A. Daina, O. Michelin, V. Zoete, SwissADME: a free web tool to evaluate pharmacokinetics, drug-likeness and medicinal chemistry friendliness of small molecules, *Sci. Rep.* 7 (2017) 42717.
- [3] C.A. Marchant, K.A. Briggs, A. Long, In silico tools for sharing data and knowledge on toxicity and metabolism: derek for windows, meteor, and vitic, *Toxicol. Mech. Methods* 18 (2008) 177-187.