

Figure S1. <sup>1</sup>H NMR spectrum of compound 3a.







Figure S3. <sup>1</sup>H NMR spectrum of compound 3c.



Figure S4. <sup>1</sup>H NMR spectrum of compound 3d.



Figure S5. <sup>1</sup>H NMR spectrum of compound 3e.



Figure S6. <sup>13</sup>C NMR spectrum of compound 3e.



Figure S7. IR spectrum of compound 3e.

## Elemental Composition Report

Single Mass Analysis Tolerance = 10.0 mDa / DBE: min = -2.0, max = 1000.0 Element prediction: Off Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions 60 formula(e) evaluated with 3 results within limits (up to 19 closest results for each mass) Elements Used: C: 0-100 H: 0-200 N: 3-3 O: 0-30 24-Jun-2016 jhm-24jun16-77 284 (5.252) Cn (Cen,7, 50.00, Ar); Sm (SG, 3x5.00); Sb (12,5.00 )

100 % 0	338.2 338.3 0 338.50	339.00	339.2 339.3 				341.00	41.3	342.50	343.3 343.50 m/z		
Minimum:		10.0	10.0	-2.0								
Maximum:		10.0	10.0	1000.0							~	
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula			( L.L	_N [	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	
338.1506	338.1505 338.1563 338.1411	0.1 -5.7 9.5	0.3 -16.9 28.1	11.5 2.5 -1.5	n/a n/a n/a	C19 H20 C12 H24 C8 H24	N3 03 N3 08 N3 011		N N		ОН	
									[M+H]+	= 338	8.149918 Da	

Figure S8. HRMS of compound 3e.

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TOF MS ES+ 6.54e+003



Figure S9. <sup>1</sup>H NMR spectrum of compound 3f.



**Figure S10.** <sup>1</sup>H NMR spectrum of compound **3g**.



Figure S11. <sup>1</sup>H NMR spectrum of compound 3h.



Figure S12. <sup>1</sup>H NMR spectrum of compound 3i.



Figure S13. <sup>1</sup>H NMR spectrum of compound 3j.



Figure S14. <sup>1</sup>H NMR spectrum of compound 4.



Figure S15. <sup>1</sup>H NMR spectrum of compound 5.

## **Supplementary figure S16.** Evaluation of –cide/-static effect

A. Fungicide effect. S. cerevisiae (W303) exposed to Schiff base 5



B. Bacteriostatic and fungistatic effect. a. *Escherichia coli* exposed to Schiff base **3d**; b. *Enterococcus faecalis* exposed to Schiff base **3a**; c. *Candida albicans* exposed to Schiff base **3b**; d. *Candida albicans* exposed to Schiff base **3f**; e. *Aspergillus niger* exposed to Schiff base **3f**; f. *Staphylococcus aureus* exposed to Schiff base **3f**.

(2% DMSO)



Compound	MW	HBA	HBD	nrotb	PSA	м	т	RE	Ι	cLogP	cLogS	DL	DS	IARC <sup>b</sup>
Cyclophosphamide	261.09	4	1	4	51.38	high	high	high	none	0.73	-1.90	-10.32	0.10	1
Methotrexate	454.45	13	5	9	210.54	none	high	none	none	-1.23	-3.77	-7.59	0.22	3
Fluorouracil	130.08	4	2	0	58.20	high	high	high	high	-0.84	-1.77	-4.50	0.06	3
Doxorubicin	543.52	12	6	5	206.07	none	none	none	high	0.17	-4.51	6.65	0.33	NR
Procarbazine	221.3	4	3	5	53.16	high	high	high	none	0.24	-2.41	3.86	0.20	2A
Prednisolone	360.45	5	3	2	94.83	none	none	none	none	1.14	-2.96	4.10	0.85	NR
Bleomycin	1415.57	38	20	36	683.55	none	none	none	none	-9.96	-2.70	5.30	0.48	2B
Etoposide	588.56	13	3	5	160.83	none	none	none	none	0.67	-3.95	-1.98	0.30	1
Cisplatin	300.05	2	2	0	55.28	none	none	none	none	0.00	-1.93	-3.18	0.49	2A
Epirubicin	543.52	12	6	5	206.07	none	none	none	high	0.17	-4.51	6.65	0.33	NR
Capecitabine	359.35	9	3	6	120.69	high	none	low	low	0.47	-3.37	-21.33	0.16	NR
Folinic acid	473.44	14	7	9	215.55	none	none	none	none	-2.21	-4.08	-6.77	0.34	NR
Temozolomide	194.15	8	1	1	105.94	high	high	high	none	-0.50	-1.90	-1.80	0.12	NR
Anastrozole	293.37	5	0	4	78.29	none	high	low	none	2.61	-4.17	-4.16	0.19	NR

Supplemental Table S1. Theoretical prediction of ADME properties<sup>a</sup> of some drug used in chemotherapy.

<sup>a</sup> MW: molecular weight; HBA: number of hydrogen bond acceptor; HBD: number of hydrogen bond dador; nrotb: number of rotatable bonds; PSA: polar surface area; M: mutagenicity; T; tumorigenicity; RE: reproductive effect; I: irritant effect; cLogP: logarithm of compound partition coefficient between n-octanol and water; cLogS: logarithm of compound aqueous solubility; DL: drug-likeness; DS: drug score. VR: voriconazole; CP: ciprofloxacin.

<sup>b</sup> Group of classification of IARC; NR = no reported.