

Supplementary data for article:

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Supporting information

Consensus based comparison of chromatographic and computational estimated lipophilicity of benzothiepine[3,2-c]pyridine derivatives as potential antifungal drugs

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Table S1. Retention parameters (R_F) of dibenzo[*b,f*]thiepines in typical reversed-phase and microemulsion chromatographic systems.

Comp. No.	Typical reversed-phase TLC															MELC-TLC		
	Organic modifier															MELC1	MELC2	
	Acetone						Methanol					Dioxane						
90%	85%	80%	75%	70%	65%	98%	94%	90%	86%	82%	85%	80%	75%	70%	65%			
1	0.63	0.62	0.57	0.52	0.48	0.41	0.53	0.48	0.46	0.44	0.4	0.84	0.75	0.71	0.62	0.37	0.30	0.43
2	0.73	0.62	0.50	0.35	0.28	0.17	0.56	0.47	0.44	0.36	0.28	0.77	0.67	0.60	0.49	0.25	0.29	0.40
3	0.59	0.50	0.42	0.29	0.23	0.14	0.45	0.36	0.32	0.27	0.18	0.74	0.64	0.53	0.37	0.16	0.27	0.37
4	0.58	0.46	0.40	0.23	0.19	0.08	0.37	0.26	0.23	0.17	0.13	0.76	0.64	0.50	0.32	0.11	0.23	0.33
5	0.64	0.52	0.44	0.26	0.24	0.12	0.44	0.34	0.32	0.24	0.17	0.78	0.66	0.51	0.35	0.14	0.21	0.31
6	0.66	0.53	0.46	0.29	0.24	0.12	0.42	0.32	0.30	0.23	0.14	0.82	0.70	0.53	0.39	0.16	0.21	0.31
7	0.66	0.51	0.45	0.25	0.22	0.09	0.48	0.38	0.35	0.26	0.18	0.78	0.67	0.51	0.37	0.15	0.21	0.30
8	0.66	0.55	0.51	0.35	0.33	0.21	0.50	0.42	0.41	0.35	0.25	0.83	0.76	0.60	0.46	0.23	0.26	0.37
9	0.64	0.50	0.45	0.29	0.27	0.14	0.46	0.38	0.35	0.29	0.20	0.79	0.69	0.53	0.42	0.22	0.22	0.32
10	0.63	0.47	0.41	0.22	0.21	0.08	0.43	0.31	0.28	0.20	0.14	0.76	0.66	0.49	0.34	0.17	0.19	0.30
11	0.63	0.45	0.41	0.24	0.18	0.11	0.42	0.30	0.28	0.20	0.15	0.78	0.67	0.52	0.32	0.19	0.23	0.33
12	0.64	0.47	0.42	0.25	0.21	0.10	0.35	0.24	0.21	0.15	0.12	0.82	0.69	0.52	0.35	0.15	0.23	0.33
13	0.73	0.53	0.46	0.31	0.11	0.11	0.53	0.41	0.39	0.29	0.20	0.88	0.74	0.57	0.37	0.18	0.22	0.32
14	0.67	0.50	0.42	0.27	0.22	0.09	0.42	0.29	0.26	0.18	0.14	0.77	0.68	0.53	0.36	0.16	0.23	0.33
15	0.76	0.56	0.48	0.31	0.25	0.10	0.53	0.42	0.39	0.3	0.24	0.83	0.74	0.57	0.43	0.15	0.24	0.33
16	0.75	0.58	0.56	0.39	0.35	0.23	0.49	0.42	0.39	0.33	0.26	0.76	0.72	0.62	0.52	0.25	0.30	0.39
17	0.71	0.56	0.53	0.33	0.32	0.22	0.54	0.46	0.43	0.39	0.28	0.72	0.70	0.56	0.47	0.21	0.31	0.40

Table S2a. *In silico* estimated logP values and solubility parameter of benzothiepine[3,2-c]pyridine derivatives.

Comp. No	miLogP	KOWWIN	AlogPs	AClogP	AlogP	MLOGP	XlogP2	XlogP3	AlogS*
1	2.66	2.70	2.14	2.29	1.80	1.51	1.90	2.34	-2.83
2	3.75	3.73	3.54	3.37	2.95	2.60	3.15	3.14	-4.05
3	4.41	4.37	4.15	3.98	3.62	3.19	3.78	4.04	-4.64
4	5.52	5.49	5.24	5.05	4.47	3.91	5.09	5.04	-6.29
5	5.48	5.49	5.33	4.40	4.43	3.91	5.01	4.59	-5.67
6	5.53	5.57	5.23	4.30	4.42	3.53	4.93	4.56	-5.67
7	5.64	5.69	5.27	4.46	4.64	4.30	5.17	4.69	-5.38
8	5.23	5.04	4.67	4.21	4.31	3.46	4.74	4.31	-4.89
9	4.49	4.62	4.44	4.06	3.41	3.32	3.87	3.82	-4.70
10	5.65	6.02	5.13	5.45	4.78	4.45	5.27	5.20	-5.61
11	5.48	5.49	5.34	4.40	4.43	3.91	5.01	4.59	-5.65
12	5.58	5.57	4.92	4.95	4.46	3.53	5.01	5.01	-6.25
13	5.40	5.57	4.81	5.26	4.66	3.99	4.99	4.92	-4.75
14	5.53	5.57	5.22	4.30	4.42	3.53	4.93	4.56	-5.70
15	5.28	5.04	4.68	4.86	4.35	3.46	4.82	4.76	-4.96
16	3.79	3.81	3.48	3.26	2.94	2.31	3.07	3.39	-4.14
17	2.92	3.93	3.50	3.43	3.16	3.06	3.32	3.52	-3.78

*Data are estimated solubilities

Table S2b. Chromatographic lipophilicity descriptors of benzothiepine[3,2-c]pyridine derivtaves.

Comp. No.	R_M^0_MeOH	b_MeOH	PCI_MeOH	C_0_MeOH	mR_M_MeOH	R_M^0_Acet	b_Acet	PCI_Acet	C_0_Acet	mR_M_Acet	R_M^0_Diox	b_Diox	PCI_Diox	C_0_Diox	mR_M_Diox	R_M_MLC1	R_M_MLC2
1	1.25	-0.01	-3.65	95.06	0.07	1.16	-0.02	-4.96	73.22	-0.07	2.94	-0.04	-5.00	67.77	-0.31	0.37	0.12
2	2.90	-0.03	-2.79	94.64	0.14	3.55	-0.04	-2.08	80.22	0.12	3.39	-0.05	-0.98	72.79	-0.10	0.39	0.17
3	3.33	-0.03	0.33	100.60	0.35	3.18	-0.04	1.02	84.99	0.28	4.28	-0.06	1.59	75.69	0.04	0.44	0.23
4	3.75	-0.04	3.19	105.31	0.54	3.94	-0.05	2.71	85.93	0.39	5.19	-0.07	0.30	76.43	0.10	0.53	0.30
5	3.49	-0.03	0.76	101.07	0.38	3.61	-0.04	0.78	83.85	0.27	4.89	-0.06	1.71	75.63	0.04	0.57	0.34
6	3.79	-0.04	1.45	101.51	0.43	3.68	-0.04	0.23	83.10	0.25	4.95	-0.07	1.17	74.51	-0.03	0.57	0.34
7	3.68	-0.04	-0.13	98.75	0.33	4.09	-0.05	1.12	83.68	0.30	4.74	-0.06	0.86	75.35	0.02	0.57	0.37
8	2.65	-0.03	-1.72	97.71	0.21	2.69	-0.03	-1.67	81.33	0.13	4.34	-0.06	-1.10	72.43	-0.15	0.46	0.24
9	3.10	-0.03	-0.28	99.94	0.31	3.25	-0.04	0.37	83.72	0.24	4.05	-0.05	-0.07	73.96	-0.06	0.54	0.32
10	4.02	-0.04	1.76	101.45	0.45	4.09	-0.05	2.32	84.99	0.36	4.47	-0.06	1.04	75.70	0.04	0.62	0.37
11	3.74	-0.04	1.82	102.46	0.45	3.78	-0.04	2.22	85.33	0.35	4.50	-0.06	-1.88	75.22	0.01	0.53	0.30
12	3.85	-0.04	3.90	106.40	0.59	3.84	-0.05	1.76	84.67	0.32	5.16	-0.07	1.31	74.95	0.00	0.53	0.30
13	3.72	-0.04	-1.18	96.71	0.26	4.66	-0.06	0.72	82.68	0.29	5.49	-0.07	-0.77	73.52	-0.11	0.54	0.32
14	4.01	-0.04	2.24	102.41	0.49	4.07	-0.05	1.29	83.73	0.30	4.61	-0.06	-0.29	75.21	0.01	0.52	0.30
15	3.23	-0.03	-1.52	96.90	0.23	4.33	-0.05	-0.66	81.27	0.20	5.11	-0.07	-2.16	73.87	-0.08	0.49	0.32
16	2.57	-0.03	-1.50	98.56	0.22	2.94	-0.04	-3.17	78.68	0.04	3.36	-0.05	2.63	72.08	-0.14	0.37	0.19
17	2.58	-0.03	-2.68	95.36	0.15	2.92	-0.04	-2.02	80.43	0.11	3.54	-0.05	1.65	73.93	-0.05	0.35	0.17

Table S3. SRD scores for standardized (Std), range scaled (Rng) and Rank (Rnk) lipophilicity data.

Std		Rng		Rnk	
Variable	SRD	Variable	SRD	Variable	SRD
XlogP2	13.89	XlogP2	13.89	milogP	13.89
<i>PCI</i> _Acet	13.89	<i>PCI</i> _Acet	13.89	XlogP2	13.89
milogP	15.28	milogP	15.28	AlogP	15.97
<i>mR_M</i> _Acet	15.28	<i>mR_M</i> _Acet	15.28	XlogP3	16.67
AlogP	15.97	AlogP	15.97	<i>PCI</i> _Acetone	16.67
XlogP3	19.44	XlogP3	19.44	KOWWIN	18.06
<i>R_M⁰</i> _MeOH	19.44	<i>R_M⁰</i> _MeOH	19.44	<i>mR_M</i> _Acetone	18.06
mlogP	20.14	mlogP	20.14	<i>R_M⁰</i> _MeOH	19.44
KOWWIN	20.83	KOWWIN	20.83	mlogP	20.14
AClogP	20.83	AClogP	20.83	AClogP	20.83
AlogpS	20.83	AlogpS	20.83	AlogpS	22.22
AlogPs	23.61	AlogPs	23.61	<i>mR_M</i> _MeOH	22.22
<i>b</i> _MeOH	23.61	<i>b</i> _MeOH	23.61	<i>b</i> _MeOH	23.61
<i>mR_M</i> _MeOH	23.61	<i>mR_M</i> _MeOH	23.61	<i>PCI</i> _MeOH	23.61
<i>C₀</i> _Acet	23.61	<i>C₀</i> _Acet	23.61	AlogPs	25.00
<i>C₀</i> _Diox	23.61	<i>C₀</i> _Diox	23.61	<i>C₀</i> _Acet	26.39
<i>PCI</i> _MeOH	25.00	<i>PCI</i> _MeOH	25.00	<i>C₀</i> _Diox	26.39
<i>mR_M</i> _Diox	25.00	<i>mR_M</i> _Diox	25.00	<i>mR_M</i> _Diox	26.39
<i>R_M⁰</i> _Diox	26.39	<i>R_M⁰</i> _Diox	26.39	<i>R_M⁰</i> _Diox	27.78
<i>C₀</i> _MeOH	31.94	<i>C₀</i> _MeOH	31.94	<i>C₀</i> _MeOH	30.56
<i>R_M⁰</i> _Acet	31.94	<i>R_M⁰</i> _Acet	31.94	<i>R_M⁰</i> _Acet	31.94
<i>b</i> _Diox	33.33	<i>b</i> _Diox	33.33	<i>b</i> _Diox	31.94
<i>R_M</i> _MELC1	33.33	<i>R_M</i> _MELC1	33.33	<i>R_M</i> _MELC1	31.94
<i>b</i> _Acet	34.72	<i>b</i> _Acet	34.72	<i>R_M</i> _MELC2	34.03
<i>R_M</i> _MELC2	36.11	<i>R_M</i> _MELC2	36.11	<i>b</i> _Acet	34.72
<i>PCI</i> _Diox	59.72	<i>PCI</i> _Diox	59.72	<i>PCI</i> _Diox	59.72

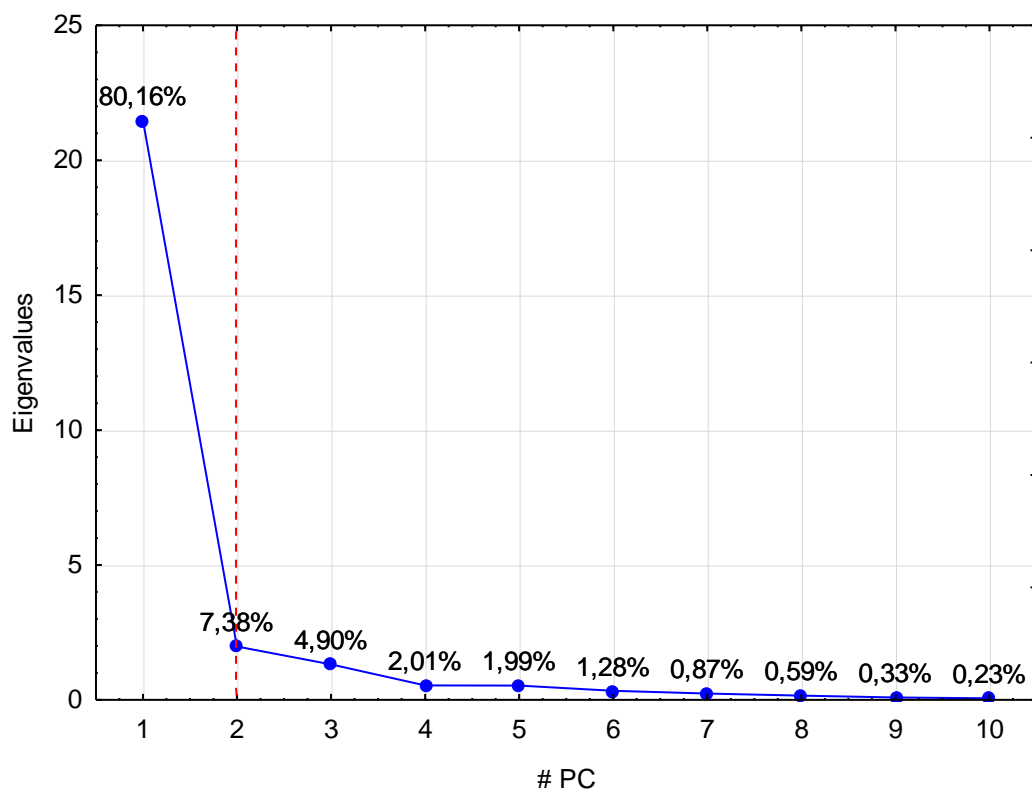


Figure S1 Eigenvalues scree plot showing distribution of data variability by principal components. Following the “broken stick” rule the first two principal components can be selected to efficiently describe retention data structure.

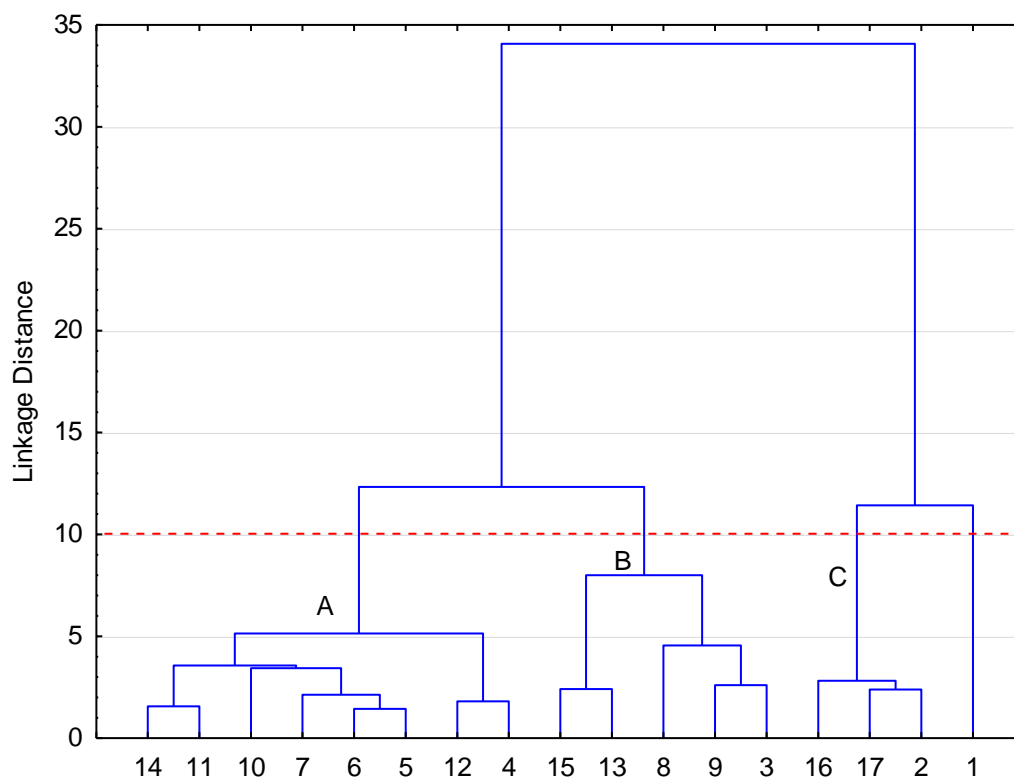


Figure S2 HCA of retention data of compounds. Three clusters (A, B and C) can be observed at linkage distance of 10 units and one outlier (compound no. 1). For numeration and chemical structures of compounds see Table 1 in the manuscript