

**A COMPARATIVE STUDY OF MOLECULAR  
LIPOPHILICITY INDICES OF SOME FORMYL-  
AND ACETYLPIRIDINE-3-  
THIOSEMICARBAZONE DERIVATIVES  
ESTIMATED BY RP- HPTLC AND CALCULATED  
*LOG P* VALUES**



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**Cristina Onișor** and **Costel Sârbu**

**Babeș-Bolyai University, Faculty of Chemistry and  
Chemical Engineering, Cluj-Napoca, Romania**

***costelsrb@yahoo.co.uk***



# QSAR/QSPR/QSRR

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➤ **QSAR** - Quantitative Structure-Activity Relations

- relates molecular structure to biological activity

➤ **QSPR** - Quantitative Structure-Property Relations

- correlates molecular information with different properties

➤ **QSRR** - Quantitative Structure-Retention Relations

- explains chromatographic retention by molecular information



# Lipophilicity

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- **Log P** estimated by direct equilibration method or by calculation according to different mathematical models

- Octanol - water partition coefficient (Log P)

$$P = C_o / C_w$$

- The “**flask shaking**” method has some disadvantages:

- it is tedious

- time consuming

- may be applied in a limited range on the lipophilicity scale



# Estimation of Lipophilicity by RPTLC

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- RPTLC method based on the assumed linear relationship between the molecular parameter  $R_M$  and  $\text{Log}P$

$$R_M = \log (1/ R_f - 1)$$

- $R_M$  (related to molecular lipophilicity) depends linearly on the concentration of the organic modifier of the mobile phase

$$R_M = R_{M_0} + bC$$

- $\varphi_0 = R_{M_0}/b$

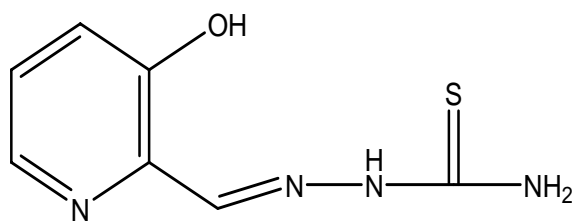


# Principal Component Analysis

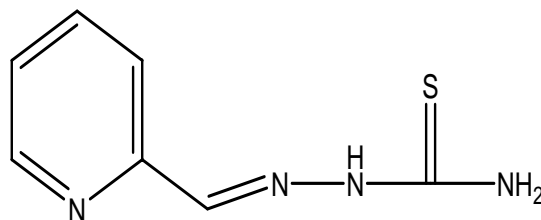
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- Principal component analysis (**PCA**) is a favorite tool in chemometrics for data compression and information extraction.
- **PCA** finds linear combinations of the original measurement variables that describe the significant variations in the data.
- **PCA** represents in an economic way the location of the compounds in a reduced coordinate system describing the data set with maximum possible information.
- **PCA** gives both coordinates (**scores**) of the studied compounds and the **loadings** of variables on the principal components.

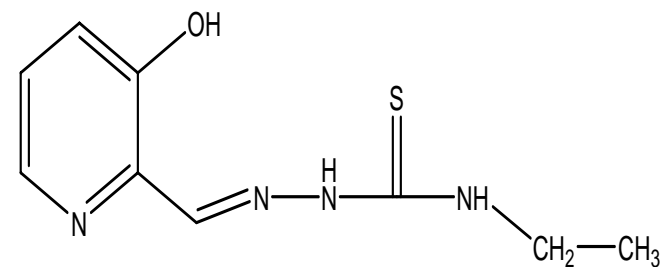
# Structure of the studied compounds



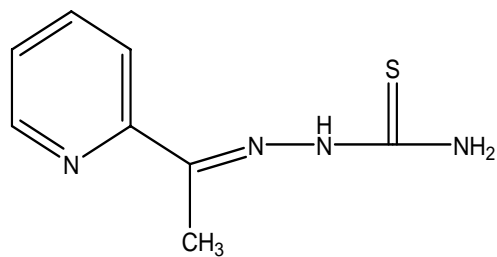
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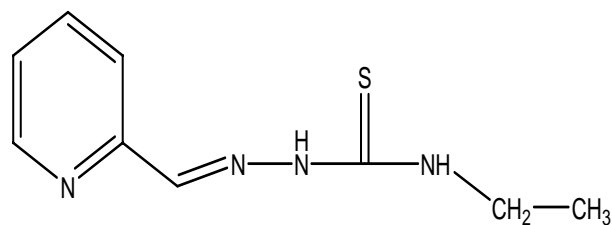
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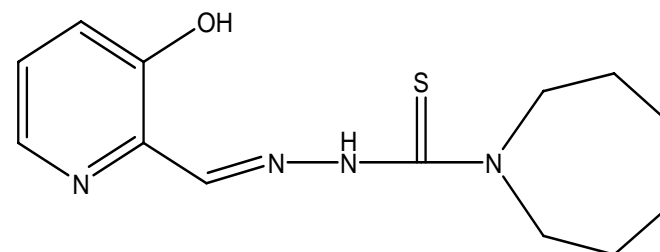
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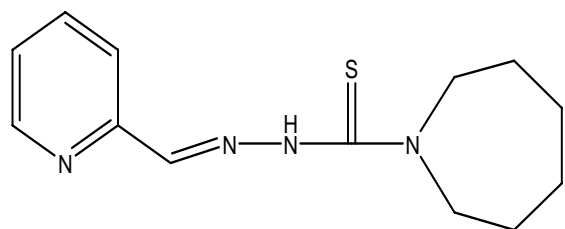


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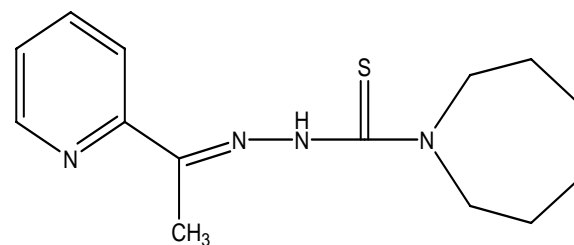


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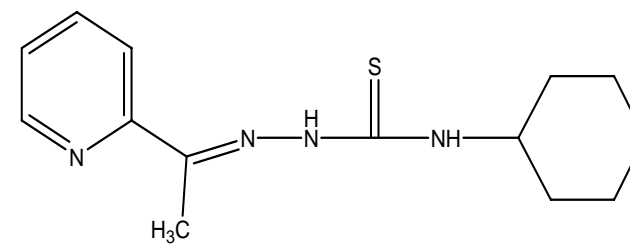
# Structure of the studied compounds



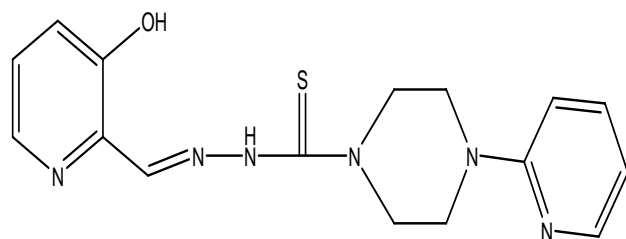
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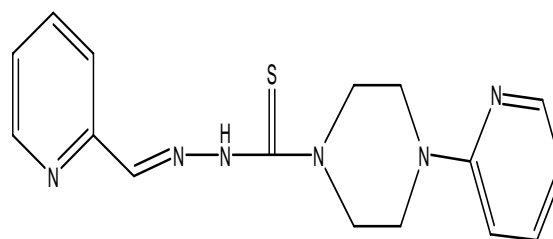
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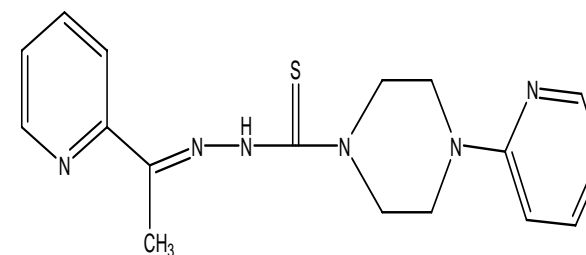
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11



12




# Experimental

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- ***Stationary phase***: C<sub>18</sub> silica gel bonded plates
- RP-C<sub>18</sub>/UV<sub>254</sub> (20x20 cm)
- RP-C<sub>18</sub>W/UV<sub>254</sub> (10x20 cm) Merck (Darmstadt, Germany)
- ***Mobile phase***: methanol-water (25-45 %; 15-35 v/v)
- **Colored zones** appeared on a colorless background
- **fluorescent blue-orange** zones under UV lamp  
( $\lambda = 365 \text{ nm}$ )




# Lipophilicity indices based on retention data (RP-C<sub>18</sub> silica gel)



<b>Cpd</b>	<b>R<sub>M0</sub></b>	<b><i>b</i></b>	<b>φ<sub>0</sub></b>	<b>PC1<sub>RF</sub></b>	<b>Mean<sub>RF</sub></b>	<b>PC1<sub>RM</sub></b>	<b>Mean<sub>RM</sub></b>
<b>1</b>	0.922	-0.029	-31.893	-0.434	0.549	1.153	-0.090
<b>2</b>	1.332	-0.046	-29.275	-0.600	0.621	1.486	-0.262
<b>3</b>	2.160	-0.060	-36.063	-0.277	0.473	0.726	0.061
<b>4</b>	1.611	-0.050	-32.020	-0.495	0.570	1.227	-0.152
<b>5</b>	2.297	-0.062	-37.107	-0.217	0.445	0.568	0.129
<b>6</b>	3.407	-0.080	-42.485	0.180	0.273	-0.518	0.597
<b>7</b>	3.195	-0.071	-44.874	0.295	0.227	-0.715	0.702
<b>8</b>	2.951	-0.065	-45.747	0.276	0.224	-0.662	0.693
<b>9</b>	3.862	-0.086	-44.856	0.355	0.199	-1.083	0.846
<b>10</b>	3.465	-0.076	-45.593	0.352	0.197	-0.949	0.802
<b>11</b>	3.080	-0.068	-45.100	0.288	0.225	-0.671	0.687
<b>12</b>	2.850	-0.063	-45.244	0.278	0.235	-0.562	0.644

# Lipophilicity indices based on retention data

(RP-C<sub>18</sub>W silica gel)



Cpd	R <sub>M0</sub>	<i>b</i>	φ <sub>0</sub>	PC1 <sub>RF</sub>	Mean <sub>RF</sub>	PC1 <sub>RM</sub>	Mean <sub>RM</sub>
1	1.468	-0.056	-26.254	-0.523	0.469	1.414	0.069
2	1.345	-0.053	-25.615	-0.547	0.488	1.480	0.031
3	2.198	-0.071	-30.777	-0.241	0.334	0.646	1.069
4	1.519	-0.056	-27.269	-0.464	0.446	1.273	0.125
5	2.074	-0.067	-30.994	-0.222	0.333	0.657	0.399
6	2.801	-0.066	-42.313	0.363	0.099	-1.018	1.144
7	2.493	-0.059	-42.534	0.317	0.120	-0.767	1.025
8	2.371	-0.066	-35.703	0.084	0.214	-0.054	0.708
9	3.300	-0.090	-36.872	0.234	0.143	-0.820	1.062
10	2.169	-0.045	-48.296	0.372	0.100	-0.777	1.044
11	3.075	-0.081	-37.781	0.255	0.137	-0.779	1.039
12	2.755	-0.061	-45.544	0.372	0.095	-1.255	1.243



# Computed different *Log P*

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- **Computer programs based on atom contributions:**
  - SciQSAR: *LogP<sup>1</sup>*
  - SciLogP: *LogP<sub>c</sub>*
  - Chem3D Ultra 8.0: *LogP<sup>2</sup>*, *PartCoeff*
  - XLOGP: *XLOGP*
- **Computer programs based on atom/fragment contributions:**
  - KOWWIN: *KOWWIN*
- **Computer programs based on fragmental contributions:**
  - cLogP: *cLogP*
- **Computer programs based on atom-type electrotopological-state indices and neural network modeling:**
  - ALOGPS: *ALOGPs*, *AB/LogP*, *miLogP*, *AvLogP*, *COSMOFrag*
  - IAlogP: *IAlogP*



## Values of the computed *Log P*

Cpd	LogP <sup>1</sup>	LogP <sub>c</sub>	LogP <sup>2</sup>	Part Coeff	ALOGPs	AB/LogP	COSMO Frag	miLogP	KOWWIN	XLOGP	AvLogP
1	0.953	0.253	0.395	1.375	1.010	1.690	-0.130	0.830	1.310	0.080	0.800
2	1.279	2.382	0.785	1.088	1.090	1.090	0.130	0.710	0.790	0.490	0.720
3	1.273	2.765	1.254	1.513	1.690	3.050	1.340	1.580	2.270	1.020	1.820
4	1.468	2.345	0.350	1.324	1.320	0.790	0.790	0.630	1.950	0.630	1.020
5	1.530	1.755	1.643	1.113	1.720	2.450	1.690	1.460	1.750	1.430	1.750
6	2.624	2.804	2.443	2.769	2.780	3.110	1.920	2.860	3.840	1.960	2.750
7	2.459	3.694	2.832	2.482	2.830	2.510	2.220	2.740	3.320	2.370	2.670
8	2.948	2.388	2.397	2.278	2.940	2.600	2.880	2.660	4.480	2.510	3.010
9	2.967	3.037	2.418	3.241	3.120	3.150	3.980	2.900	4.690	2.980	3.470
10	1.361	2.669	2.502	1.408	2.370	3.210	2.550	2.140	3.160	2.030	2.580
11	2.929	2.545	2.892	1.121	2.430	2.610	2.800	2.020	2.640	2.430	2.490
12	2.825	2.262	2.457	0.917	2.540	2.700	3.450	1.930	3.800	2.580	2.830





# Conclusions

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- The lipophilic character of of formyl- and acetylpyridine-3-thiosemicarbazone derivatives has been investigated using **TLC retention data and various calculated  $\log P$**
- Statistically significant correlations were found between lipophilicity indices,  **$R_{M0}$ ,  $\phi_0$ , scores corresponding to PC1** and a new scale based on the **mean of retention indices**
- The **scores corresponding to PC1** and the **mean of retention indices** appeared to be the best solution for the lipophilicity scale resulted from the retention data
- **LogP<sup>2</sup>, AvLogP and XLOGP** have been appearing to be the most appropriate for chromatography

# QSAR/QSPR/QSRR and the Bright Future of TLC

The responsibility for  
**change** ...  
lies within us. We must  
begin with ourselves,  
teaching ourselves not  
to close our minds  
prematurely to the novel,  
the surprising, the  
seemingly radical.

Alvin Toffler



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# Thank you for your attention!

