

# On the Estimation of the Quantitative Structure-Activity Relationships Descriptor $\sigma_s$ for Aliphatic Compound

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Procedures for the estimation of the novel quantitative structure-activity relationships (QSAR) descriptors  $\sigma_s$ , representing the contributions from both dispersion and repulsion interactions, have been presented for the substituted methane series. After correcting part of the symmetry, the rotation isomer or racemate, observed values  $\sigma_s(\text{obs})$  for  $\text{CH}_2\text{R}_A\text{R}_B$ ,  $\text{CHR}_A\text{R}_B\text{R}_C$ , and  $\text{CR}_A\text{R}_B\text{R}_C\text{R}_D$  can be given by the general equation  $\sigma_s(\text{obs}) = a\Sigma\sigma_s(\text{mono}) + b$ , where  $\sigma_s(\text{mono})$  means the descriptor of the MeR series. The values estimated by this equation agreed with those of the observed ones, and several examples of the large molecule are submitted for the determination of the optimum value.

Furthermore, the hydrophobic substituent constants  $\pi$  of typical  $\alpha$ -amino acids are revealed to be expressed by the linear combination of the descriptor  $\sigma_s$  based on the gas phase and indicator variable, implying the averaged number of hydration.

**Keywords** quantitative structure-activity relationship; substituent entropy constant  $\sigma_s$ ; substituted methane derivative; regression analysis; amino acid

## Introduction

In our previous report,<sup>1)</sup> the method for the estimation of the quantitative structure-activity relationships (QSAR) descriptor  $\sigma_s$  of polysubstituted benzene derivatives having an optional set of substituent groups has been confirmed by means of the statistical way, and the results are fruitful in the evaluation of the phenomena inherent in the van der Waals force, including gas-liquid chromatography (GLC),<sup>2)</sup> etc.

In this work, the authors focused their attention on the estimation of the descriptor  $\sigma_s$  for substituted aliphatic substrates  $\text{CH}_2\text{R}_A\text{R}_B$ ,  $\text{CHR}_A\text{R}_B\text{R}_C$ , and  $\text{CR}_A\text{R}_B\text{R}_C\text{R}_D$ . The successful result after regression analyses enabled us to predict the descriptor  $\sigma_s$  for the aliphatic substrate having an optional set of substituent groups.

## Experimental

**Absolute Entropy  $S_{298}^\circ(\text{g})/\text{e.u.}$**  Data are all cited from the references.<sup>3,4)</sup>

**Regression Analysis** Regression analyses are carried out on NEC PC-9801/M/VX and EPSON PC-286V personal computers using a program package for multi-variate analyses MVA developed by Takagi, *et al.*<sup>5)</sup>

**Definition of the Descriptor  $\sigma_s(\text{obs})$  for Substituted Methane Derivatives** The observed descriptors  $\sigma_s(\text{obs})$  for MeR,<sup>6)</sup>  $\text{CH}_2\text{R}_A\text{R}_B$ ,  $\text{CHR}_A\text{R}_B\text{R}_C$  and  $\text{CR}_A\text{R}_B\text{R}_C\text{R}_D$  are defined as follows;  $\sigma_s(\text{obs}) = \log\{S_{298}^\circ(\text{g})(\text{obs})/S_{298}^\circ(\text{CH}_4)\}$ , where  $S_{298}^\circ(\text{g})(\text{CH}_4) = 44.52$  e.u. is used.

## Results and Discussion

**Correlations between  $\sigma_s$  and  $\Sigma\sigma_s(\text{mono})$  in Substituted Methanes** The successful results<sup>1)</sup> on the estimation of the descriptor  $\sigma_s$  for polysubstituted benzene derivatives having an optional set of substituent as a function of  $\Sigma\sigma_s(\text{mono})$  have prompted us to develop the strategy in the field of aliphatic substrate.

Followed by the definition in our previous work,<sup>6)</sup> the presentation of the QSAR descriptor  $\sigma_s(\text{mono})$  or mono-substituted methane series is given as below;

$$\sigma_s(\text{mono}) = \log\{S_{298}^\circ(\text{g})(\text{MeR})/S_{298}^\circ(\text{g})(\text{CH}_4)\}$$

where  $\text{CH}_4$  is used as reference.

In the first step, the values of  $\sigma_s(\text{obs})$  derived from the observed absolute entropies  $S_{298}^\circ(\text{g})$  and calculated  $\Sigma\sigma_s(\text{mono})$

for the  $\text{CH}_2\text{R}_A\text{R}_B$  ( $\text{R}_A = \text{R}_B$ ,  $\text{R}_A \neq \text{R}_B$ ),  $\text{CHR}_A\text{R}_B\text{R}_C$  ( $\text{R}_A = \text{R}_B = \text{R}_C$ ,  $\text{R}_A = \text{R}_B \neq \text{R}_C$ ,  $\text{R}_A \neq \text{R}_B \neq \text{R}_C$ ), and  $\text{CR}_A\text{R}_B\text{R}_C\text{R}_D$  ( $\text{R}_A = \text{R}_B = \text{R}_C = \text{R}_D$ ,  $\text{R}_A = \text{R}_B = \text{R}_C \neq \text{R}_D$ ,  $\text{R}_A = \text{R}_B \neq \text{R}_C \neq \text{R}_D$ ,  $\text{R}_A = \text{R}_B \neq \text{R}_C = \text{R}_D$ ) series are arranged in Tables I, II, and III, where additional corrections for  $\sigma_s(\text{obs})$  are carried out, namely,  $n \times R \ln 2$  in the presence of the plane of symmetry,  $R \ln 3$  for *trans* and *gauche* isomer and  $R \ln 2$  for optical antipode. The abbreviations  $n$  and  $R$  denote the number of symmetry and gas constant, respectively. The observed data and those having the correction term

TABLE I.  $\text{CH}_2\text{R}_A\text{R}_B$  and Their Descriptors  $\sigma_s$

	$S_{298}^\circ(\text{obs})$	$\sigma_s(\text{obs})$	Correction	$\sigma_s(\text{cor})$	$\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$
1 $\text{CH}_2\text{F}_2$ <sup>a)</sup>	58.94	0.122	$2 \times R \ln 2$	0.142	0.156	0.148
2 $\text{CH}_2\text{Cl}_2$	64.59	0.162	$2 \times$	0.180	0.198	0.179
3 $\text{CH}_2\text{Br}_2$	70.08	0.197	$2 \times$	0.214	0.244	0.213
4 $\text{CH}_2\text{I}_2$	73.88	0.220	$2 \times$	0.236	0.270	0.233
5 $(\text{Me})_2\text{CH}_2$	64.51	0.161			0.182	0.167
6 $(\text{Et})_2\text{CH}_2$	83.40	0.273			0.322	0.271
7 $(n\text{-Pr})_2\text{CH}_2$	102.27	0.361			0.442	0.361
8 $\text{CH}_2\text{BrCl}$	68.67	0.188	$1 \times$	0.197	0.221	0.196
9 $\text{CH}_2\text{BrF}$	65.97	0.171	$1 \times$	0.180	0.200	0.181
10 $\text{CH}_2\text{BrI}$	73.49	0.218	$1 \times$	0.226	0.257	0.223
11 $\text{CH}_2\text{ClF}$	63.16	0.152	$1 \times$	0.161	0.177	0.163
12 $\text{CH}_2\text{ClI}$	70.78	0.201	$1 \times$	0.210	0.234	0.206
13 $\text{MeCH}_2\text{Br}$	68.71	0.188			0.213	0.190
14 $\text{MeCH}_2\text{Cl}$	65.93	0.171			0.190	0.173
15 $\text{MeCH}_2\text{F}$	63.32	0.153			0.169	0.157
16 $\text{MeCH}_2\text{NH}_2$	68.08	0.184			0.206	0.185
17 $\text{MeCH}_2\text{NO}_2$	75.39	0.229			0.260	0.225
18 $\text{MeCH}_2\text{OH}$	67.54	0.181			0.201	0.181
19 $\text{MeCH}_2\text{SH}$	70.77	0.210			0.227	0.201
20 $\text{MeCH}_2\text{CN}$	68.50	0.187			0.211	0.189
21 $\text{MeCH}_2\text{CHO}$	72.83	0.214			0.243	0.213
22 $(\text{MeCH}_2)_2$	74.12	0.221			0.252	0.219
23 $(\text{ClCH}_2)_2$	73.66	0.219	$1 \times R \ln 3$	0.231	0.271	0.233
24 $(\text{BrCH}_2)_2$	78.81	0.248	$1 \times$	0.260	0.310	0.262
25 $(\text{ICH}_2)_2$	83.30	0.272	$1 \times$	0.283	0.336	0.282
26 $(\text{HOCH}_2)_2$	77.33	0.240	$1 \times$	0.252	0.292	0.249
27 $(\text{NCCCH}_2)_2$	79.04	0.249	$1 \times$	0.261	0.309	0.262
28 $(\text{tert-BuCH}_2)_2$	112.35	0.402			0.494	0.400
29 $(n\text{-Pentyl})_2\text{S}$	144.45	0.513			0.664	0.526
30 $(n\text{-Bu})_2\text{S}$	125.84	0.451			0.566	0.453

a) MeF, having  $S_{298}^\circ(\text{g}) = 53.25$  e.u., gives  $\sigma_s(\text{mono}) = 0.078$ , and for  $\text{CH}_2\text{F}_2$ , we get  $\Sigma\sigma_s(\text{mono}) = 0.156$ . Using Eq. 1,  $\text{CH}_2\text{F}_2$  belonged to  $\text{CH}_2\text{R}_A\text{R}_B$ , affords  $\sigma_s(\text{cal}) = 0.148$ . On the other hand,  $\text{CH}_2\text{F}_2$ , having  $S_{298}^\circ(\text{g})(\text{obs}) = 58.94$  e.u. and  $\sigma_s(\text{obs}) = 0.122$ , affords, after correcting the symmetry part  $2 \times R \ln 2 = 2.75$  e.u.,  $S_{298}^\circ(\text{g})(\text{corr.}) = 58.94 + 2.75 = 61.69$  e.u. and  $\sigma_s(\text{corr.}) = 0.142$ .

TABLE II. CHR<sub>A</sub>R<sub>B</sub>R<sub>C</sub> and Their Descriptors  $\sigma_s$ 

	$S_{298}^\circ(\text{g})(\text{obs})$	$\sigma_s(\text{obs})$	Correction	$\sigma_s(\text{cor})$	$\Sigma\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$
1 CHF <sub>3</sub>	62.04	0.144	3 × R ln 2	0.172	0.234	0.182
2 CHCl <sub>3</sub>	70.66	0.201	3 ×	0.225	0.297	0.224
3 CHBr <sub>3</sub>	79.03	0.249	3 ×	0.271	0.366	0.269
4 CHI <sub>3</sub>	85.00	0.281	3 ×	0.301	0.405	0.295
5 CHMe <sub>3</sub>	70.42	0.199			0.273	0.208
6 CHEt <sub>3</sub>	98.35	0.344			0.483	0.346
7 CHBrCl <sub>2</sub>	75.56	0.230	1 ×	0.238	0.326	0.243
8 CHBrF <sub>2</sub>	70.51	0.200	1 ×	0.208	0.278	0.211
9 CHBr <sub>2</sub> Cl	78.31	0.245	1 ×	0.253	0.343	0.254
10 CHBr <sub>2</sub> F	75.70	0.231	1 ×	0.238	0.322	0.240
11 CHClF <sub>2</sub>	67.13	0.178	1 ×	0.187	0.255	0.196
12 CHCl <sub>2</sub> F	69.99	0.196	1 ×	0.205	0.276	0.210
13 Me <sub>2</sub> CHOH	74.03	0.221			0.292	0.220
14 Me <sub>2</sub> CHSH	77.51	0.241			0.318	0.238
15 Me <sub>2</sub> CHNO <sub>2</sub>	83.10	0.271			0.351	0.259
16 Me <sub>2</sub> CHCl	72.70	0.213			0.281	0.213
17 Me <sub>2</sub> CHBr	85.53	0.230			0.304	0.228
18 Me <sub>2</sub> CHF	69.82	0.195			0.260	0.199
19 Me <sub>2</sub> CHI	77.55	0.241			0.317	0.237
20 Me <sub>2</sub> CHCN	74.88	0.226			0.302	0.227
21 MeO-iso-Pr	80.86	0.259			0.338	0.251
22 MeS-iso-Pr	85.87	0.285			0.368	0.270
23 <i>n</i> -PrCHEt <sub>2</sub>	109.51	0.391			0.543	0.386
24 <i>n</i> -Pr <sub>2</sub> CHEt	118.52	0.425			0.603	0.425
25 CHBrClF <sup>a)</sup>	72.88	0.214	-1 ×	0.206	0.298	0.224
26 MeCHClEt <sup>a)</sup>	85.94	0.286	-1 ×	0.279	0.351	0.259
27 MeCHOHEt <sup>a)</sup>	85.81	0.285	-1 ×	0.278	0.362	0.267
28 MeCHSHEt <sup>a)</sup>	87.65	0.294	-1 ×	0.287	0.388	0.284
29 MeCHNH <sub>2</sub> Et <sup>a)</sup>	83.90	0.275	-1 ×	0.268	0.367	0.270
30 MeEtCH- <i>n</i> -Pr <sup>a)</sup>	101.37	0.357	-1 ×	0.351	0.473	0.340
31 ( <i>n</i> -Pr) <sub>3</sub> CH	125.56	0.450			0.663	0.465
32 ( <i>n</i> -Pr) <sub>2</sub> CH-iso-Pr	124.63	0.447			0.641	0.450
33 <i>n</i> -PrEtCH- <i>n</i> -Bu <sup>a)</sup>	129.12	0.462	-1 ×	0.458	0.655	0.459

a) Racemic compound.

afford excellent linear relations with  $\Sigma\sigma_s(\text{mono})$  as given below;

$$\sigma_s(\text{CH}_2\text{R}_A\text{R}_B) = 0.745(0.011)\Sigma\sigma_s(\text{mono}) + 0.032(0.003) \quad (1)$$

$$n = 30, r = 0.999, s = 0.003$$

$$\sigma_s(\text{CHR}_A\text{R}_B\text{R}_C) = 0.658(0.025)\Sigma\sigma_s(\text{mono}) + 0.028(0.010) \quad (2)$$

$$n = 32, r = 0.995, s = 0.008$$

$$\sigma_s(\text{CR}_A\text{R}_B\text{R}_C\text{R}_D) = 0.684(0.022)\Sigma\sigma_s(\text{mono}) - 0.013(0.010) \quad (3)$$

$$n = 40, r = 0.995, s = 0.005$$

And, the calculated values of  $\sigma_s(\text{cal})$  given by Eqs. 1—3 are all aligned in the extreme right of the Tables.

Using Eq. 1, we are able to estimate the descriptor for important CH<sub>2</sub>R<sub>A</sub>R<sub>B</sub> type substrate, as given in Table IVa.

**Optimum of Descriptor  $\sigma_s(\text{cal})$  and  $\Sigma\sigma_s(\text{mono})$  as Fragment Sum** In contrast to benzene derivatives, aliphatic substrate composed of a variety of binding fragments does not specify the optimal  $\Sigma\sigma_s(\text{mono})$ . This situation is approved for several kinds of examples given in Chart 1, where the fragment tentatively specified by the optional reference position affords good agreement among observed and calculated descriptors, and the following conclusions are obtained;

- 1) The average of  $\Sigma\sigma_s(\text{mono})$  is desirable.
- 2) The disparity of the size of the fragment should be kept out.
- 3) Under conditions 1) and 2), an optional choice of fragment affords that the deviation not exceed 5% from  $S_{298}^\circ(\text{g})(\text{obs})$ .

From the above view point, the authors selected and

TABLE III. CR<sub>A</sub>R<sub>B</sub>R<sub>C</sub>R<sub>D</sub> and Their Descriptors  $\sigma_s$ 

	$S_{298}^\circ(\text{obs})$	$\sigma_s(\text{obs})$	Correction	$\sigma_s(\text{cor})$	$\Sigma\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$
1 ClF <sub>2</sub> CCN	76.08	0.233	1 × R ln 2	0.241	0.375	0.243
2 Cl <sub>2</sub> FCCN	79.33	0.251	1 ×	0.258	0.396	0.257
3 Cl <sub>3</sub> CCN	79.38	0.251	3 ×	0.273	0.417	0.272
4 F <sub>3</sub> CCN	71.27	0.204	3 ×	0.229	0.354	0.229
5 Me <sub>4</sub> C	73.23	0.216	6 ×	0.263	0.364	0.236
6 Et <sub>4</sub> C	110.31	0.394	6 ×	0.425	0.644	0.427
7 CF <sub>4</sub>	62.50	0.147	6 ×	0.201	0.312	0.200
8 CCl <sub>4</sub>	74.12	0.221	6 ×	0.267	0.396	0.257
9 CBr <sub>4</sub>	85.53	0.284	6 ×	0.324	0.488	0.320
10 Cl <sub>4</sub>	93.60	0.323	6 ×	0.359	0.540	0.356
11 CBrClF <sub>2</sub>	76.14	0.233	1 ×	0.241	0.377	0.244
12 CBrCl <sub>2</sub> F	78.87	0.248	1 ×	0.256	0.398	0.259
13 CBrCl <sub>3</sub>	79.55	0.252	3 ×	0.274	0.419	0.273
14 CBrF <sub>3</sub>	71.19	0.204	3 ×	0.228	0.356	0.230
15 CBr <sub>2</sub> ClF	81.89	0.265	1 ×	0.272	0.421	0.275
16 CBr <sub>2</sub> Cl <sub>2</sub>	83.23	0.272	2 ×	0.286	0.442	0.289
17 CBr <sub>2</sub> F <sub>2</sub>	77.66	0.242	2 ×	0.257	0.400	0.260
18 CBr <sub>3</sub> Cl	85.36	0.283	3 ×	0.303	0.465	0.305
19 CBr <sub>3</sub> F	82.65	0.269	3 ×	0.290	0.444	0.290
20 CCl <sub>3</sub> F	68.28	0.186	3 ×	0.211	0.333	0.214
21 CCl <sub>2</sub> F <sub>2</sub>	71.91	0.208	2 ×	0.225	0.354	0.229
22 CClF <sub>3</sub>	74.13	0.221	3 ×	0.245	0.375	0.243
23 CF <sub>3</sub> I	73.50	0.218	3 ×	0.241	0.369	0.239
24 CF <sub>2</sub> I <sub>2</sub>	82.78	0.269	2 ×	0.284	0.426	0.278
25 CCl <sub>3</sub> I	81.34	0.262	3 ×	0.282	0.432	0.282
26 Me <sub>3</sub> CCl	77.00	0.238			0.372	0.241
27 Me <sub>2</sub> CCl <sub>2</sub>	77.92	0.243			0.380	0.247
28 Me <sub>2</sub> CBr	79.34	0.251			0.395	0.257
29 Me <sub>3</sub> CI	81.79	0.264			0.408	0.266
30 Me <sub>2</sub> COH	77.98	0.243			0.383	0.249
31 Me <sub>2</sub> CSH	80.79	0.259			0.409	0.266
32 EtMe <sub>2</sub> COH	87.68	0.294			0.453	0.296
33 EtMe <sub>2</sub> CSH	92.48	0.317			0.479	0.314
34 EtCMe <sub>3</sub>	85.62	0.284			0.434	0.283
35 <i>n</i> -PrCMe <sub>3</sub>	93.90	0.324			0.494	0.324
36 Me <sub>2</sub> C( <i>n</i> -Pr) <sub>2</sub>	113.87	0.408			0.624	0.413
37 Et <sub>2</sub> CMe <sub>2</sub>	95.53	0.332			0.504	0.331
38 iso-PrCMe <sub>3</sub>	91.61	0.313			0.472	0.309
39 <i>tert</i> -BuO(iso-Pr)	99.89	0.351			0.536	0.366
40 EtMe <sub>2</sub> C(iso-Pr)	103.14	0.365			0.542	0.357

TABLE IVa. CH<sub>2</sub>R<sub>A</sub>R<sub>B</sub> Series and Their  $\Sigma\sigma_s(\text{mono})$ ,  $\sigma_s(\text{cal})$  and  $S_{298}^\circ(\text{g})(\text{e.u.})$  (cal)

R <sub>A</sub>	R <sub>B</sub>	$\Sigma\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$	$S_{298}^\circ(\text{g})(\text{cal})$
Ph	COOH	0.417	0.342	97.85
Me	COOH	0.272	0.234	76.31
Me	CONH <sub>2</sub>	0.282	0.242	77.65
Ph	Ph	0.472	0.383	107.54
Ph	CH <sub>2</sub> Ph	0.523	0.421	117.37
OPh	Ph	0.523	0.421	117.37
NH <sub>2</sub>	COOH	0.296	0.252	79.53

TABLE IVb. R<sub>A</sub>CH<sub>2</sub>CH<sub>2</sub>R<sub>B</sub> Series and Their  $\Sigma\sigma_s(\text{mono})$ ,  $\sigma_s(\text{cal})$  and  $S_{298}^\circ(\text{g})(\text{e.u.})$  (cal)

R <sub>A</sub>	R <sub>B</sub>	$\Sigma\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$	$S_{298}^\circ(\text{g})(\text{cal})$	$S_{298}^\circ(\text{g})(\text{obs})^{3,4)}$
Ph	COOH	0.469	0.381	107.04	
Me	COOH	0.353	0.295	87.71	
Me	CONH <sub>2</sub>	0.343	0.287	86.21	
Me	F	0.242	0.212	72.54	72.71
Me	Cl	0.262	0.227	75.09	75.27
Me	Br	0.280	0.240	77.37	79.08
Me	I	0.294	0.251	79.35	80.32
Ph	CH <sub>2</sub> Ph	0.572	0.458	127.72	

estimated the optimal  $\Sigma\sigma_s(\text{mono})$  and  $\sigma_s(\text{cal})$  for substrates summarized in Table IVb

**Amino Acids and Their Descriptors  $\sigma_s$ .** Successful results

1)		$S_{298}^{\circ}(\text{g}) = 124.47 \text{ e.u.}$	$\sigma_s = 0.447$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}(\text{g})$ (cal)
	C-3		$0.161 + 0.371 = 0.532$	0.428	119.37
	C-4		$0.091 + 0.161 + 2 \times 0.221 = 0.694$	<u>0.462</u>	<u>128.90</u>
			average	<u>0.445</u>	<u>124.14</u>

2)		$S_{298}^{\circ}(\text{g}) = 110.32 \text{ e.u.}$	$\sigma_s = 0.394$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}(\text{g})$ (cal)
	C-3		$0.091 + 0.161 + 0.273 = 0.525$	0.373	105.20
	C-4		$0.266 + 0.221 = 0.487$	0.395	110.50
	C-5		$0.161 + 0.309 = 0.470$	<u>0.382</u>	<u>107.33</u>
			average	0.383	107.68
	when the correction of racemate is added, we get	$R \ln 2 = 1.38 \text{ e.u.}$		<u>0.389</u>	<u>109.05</u>

3)		$S_{298}^{\circ}(\text{g}) = 101.37 \text{ e.u.}$	$\sigma_s = 0.357$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}(\text{g})$ (cal)
	C-3		$0.091 + 0.161 + 0.221 = 0.473$	0.339	97.23
	C-4		$0.161 + 0.266 = 0.427$	<u>0.350</u>	<u>99.69</u>
			average	0.345	98.46
	when the correction of racemate is added, we get	$R \ln 2 = 1.38 \text{ e.u.}$		<u>0.351</u>	<u>99.84</u>

4)		$S_{298}^{\circ}(\text{g}) = 111.34 \text{ e.u.}$	$\sigma_s = 0.398$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}(\text{g})$ (cal)
	C-2	$3 \times \text{-C-} + \text{-C-C-C-C-}$	$0.273 + 0.310 = 0.583$	0.386	108.22
	C-3		$0.091 + 0.210 + 0.221 = 0.522$	0.371	104.72
	C-4		$0.161 + 0.313 = 0.474$	<u>0.385</u>	<u>108.07</u>
			average	0.381	107.00
	when the correction of racemate is added, we get	$R \ln 2 = 1.38 \text{ e.u.}$		<u>0.386</u>	<u>108.38</u>

5)		$S_{298}^{\circ}(\text{g}) = 122.90 \text{ e.u.}$	$\sigma_s = 0.441$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}(\text{g})$ (cal)
	C-3		$0.091 + 0.199 + 0.310 = 0.600$	0.423	117.86
	C-4		$0.266 + 0.293 = 0.559$	0.448	125.03
	C-5		$0.199 + 0.347 = 0.546$	<u>0.439</u>	<u>122.27</u>
			average	0.437	121.68
	when the correction of racemate is added, we get	$R \ln 2 = 1.38 \text{ e.u.}$		<u>0.442</u>	<u>123.06</u>

Chart 1. (Continued)

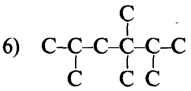
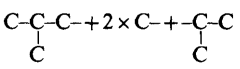
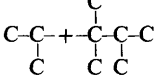
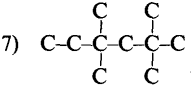
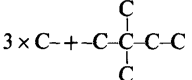
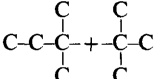
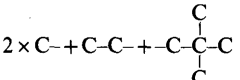
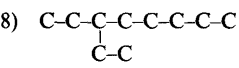
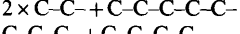

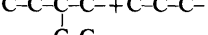
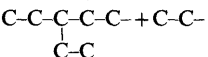
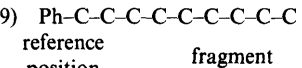
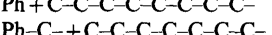
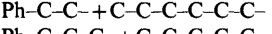
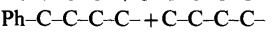
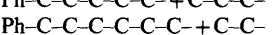

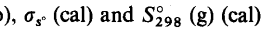

6)		$S_{298}^{\circ}$ (g) = 118.25 e.u.	$\sigma_s = 0.424$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}$ (g) (cal)
	C-3		$0.266 + 0.182 + 0.199 = 0.647$	0.430	119.70
	C-4		$0.199 + 0.313 = 0.512$	<u>0.413</u>	<u>115.34</u>
			average	<u>0.422</u>	<u>117.52</u>
7)		$S_{298}^{\circ}$ (g) = 115.91 e.u.	$\sigma_s = 0.416$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}$ (g) (cal)
	C-2		$0.273 + 0.332 = 0.605$	0.401	112.04
	C-3		$0.284 + 0.210 = 0.494$	0.400	111.84
	C-4		$0.182 + 0.161 + 0.284 = 0.627$	<u>0.416</u>	<u>115.99</u>
			average	0.406	113.29
8)		$S_{298}^{\circ}$ (g) = 127.74 e.u.	$\sigma_s = 0.458$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}$ (g) (cal)
	C-3		$0.322 + 0.319 = 0.641$	0.450	125.41
	C-4		$0.309 + 0.273 = 0.582$	0.466	130.06
	C-5		$0.344 + 0.221 = 0.565$	0.453	126.32
	C-6		$0.391 + 0.161 = 0.552$	<u>0.443</u>	<u>123.54</u>
			average	0.453	126.33
9)		$S_{298}^{\circ}$ (g) = 151.71 e.u.	$\delta_s = 0.532$		
	reference position	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)	$S_{298}^{\circ}$ (g) (cal)
	C-1		$0.236 + 0.434 = 0.670$	0.531	151.25
	C-2		$0.287 + 0.399 = 0.686$	0.543	155.46
	C-3		$0.333 + 0.361 = 0.694$	0.549	157.61
	C-4		$0.373 + 0.319 = 0.692$	0.548	157.07
	C-5		$0.410 + 0.273 = 0.683$	0.541	154.66
	C-6		$0.444 + 0.221 = 0.665$	0.527	149.95
	C-7		$0.476 + 0.161 = 0.637$	<u>0.507</u>	<u>142.93</u>
			average	<u>0.535</u>	<u>152.70</u>

Chart 1. Optimal  $\Sigma\sigma_s$  (mono),  $\sigma_s$  (cal) and  $S_{298}^{\circ}$  (g) (cal)

hitherto obtained have prompted us to estimate the descriptor  $\sigma_s$  of amino acids. With the exception of glycine, they are all classified in the  $\text{CHR}_A\text{R}_B\text{R}_C$  group. Estimations are carried out by two ways:

1) substrate as  $\text{CHR}_A\text{R}_B\text{R}_C$  type

2) substrate as  $\text{CH}_2\text{R}_A\text{R}_B$  type

where  $\sigma_s[-\text{CH}(\text{NH}_2)\text{COOH}] = 0.285$  is employed. The results are all summarized in Table V, where the descriptor  $\sigma_s(\text{R})$  given by the substituent R on *tert*-carbon becomes linear with  $\sigma_s(\text{cal})$  inherent to the substrate.

For instance, lysine  $\text{H}_2\text{N}-\text{CH}_2-\text{CH}_2-\text{CH}_2-\text{CH}_2-\text{CH}(\text{NH}_2)\text{COOH}$  gives descriptor  $\sigma_s$  by two ways 1) and 2) as

mentioned below;

1) estimation of  $\sigma_s$  for *n*-pentyl amine classified to  $\text{CH}_2\text{R}_A\text{R}_B$  is carried out under the conditions given below;

	fragment	$\Sigma\sigma_s$ (mono)	$\sigma_s$ (cal)
a.	$\text{N}-\text{C}+\text{C}-\text{C}$	$0.184 + 0.221 = 0.405$	0.334
b.	$\text{N}-\text{C}-\text{C}+\text{C}-\text{C}$	$0.241 + 0.161 = 0.402$	0.331

Mathematical average gives  $\sigma_s(\text{C}-\text{C}-\text{C}-\text{C}-\text{N}) = 0.333$ .

Then, lysine affords  $\Sigma\sigma_s(\text{mono}) = 0.333 + 0.115 + 0.181 = 0.629$ , and, using Eq. 2, we get the descriptor  $\sigma_s(\text{cal}) = 0.442$ .

2)  $\sigma_s(\text{cal}) = 0.285$  for alanine means  $\sigma_s[\text{CH}(\text{NH}_2)\text{COOH}] = 0.285$ . Using Eq. 2, we get

TABLE V. Amino Acids R-CH(NH<sub>2</sub>)COOH and Their QSAR Descriptors

	R	$\sigma_s(R)$	$\Sigma\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$	$\pi^7)$	I
Gly	H	0	0.296	0.252	0	0
Ala	Me	0.091	0.387	0.282	0.31	0
Val	iso-Pr	0.199	0.495	0.354	1.22	0
Leu	iso-Bu	0.266	0.562	0.398	1.70	0
Ile	sec-Bu	0.266	0.562	0.398	1.80	0
Ser	HOCH <sub>2</sub> -	0.182	0.478	0.343	-0.04	1
Thr	MeCH(OH)-	0.221	0.517	0.369	0.26	1
Asp	HOOC-CH <sub>2</sub> -	0.235	0.531	0.378	-0.77	2
Asn	H <sub>2</sub> NOC-CH <sub>2</sub> -	0.235	0.531	0.378	-0.60	2
Glu	HOOC-CH <sub>2</sub> CH <sub>2</sub> -	0.285	0.591	0.417	-0.64	2
Gln	H <sub>2</sub> NOC-CH <sub>2</sub> CH <sub>2</sub> -	0.285	0.591	0.417	-0.62	2
Cys	HS-CH <sub>2</sub> -	0.201	0.497	0.355	1.54	1
Met	Me-S-CH <sub>2</sub> CH <sub>2</sub> -	0.300	0.596	0.421	1.23	1
Lys	H <sub>2</sub> N-(CH <sub>2</sub> ) <sub>4</sub> -	0.330	0.626	0.445	-0.99	3
Hyl	H <sub>2</sub> N-CH <sub>2</sub> CHCH <sub>2</sub> CH <sub>2</sub> - OH	0.365	0.666	0.467		
Phe	Ph-CH <sub>2</sub> -	0.287	0.583	0.412	0.79	1
Tyr	HO-C <sub>6</sub> H <sub>4</sub> -CH <sub>2</sub> -	0.315	0.611	0.430	0.96	1
tert-Leu	tert-Bu	0.210	0.506	0.361		
$\gamma$ -Me-Leu	Neopentyl	0.284	0.580	0.410		
nor-Val	n-Pr	0.221	0.517	0.370		
nor-Leu	n-Bu	0.273	0.569	0.403		
homo-Ala	Et	0.161	0.457	0.322		

homo-alanine R = Et  $\Sigma\sigma_s(\text{mono})=0.376$   $\sigma_s(\text{cal})=0.312$

nor-valine R = n-Pr  $\Sigma\sigma_s(\text{mono})=0.446$   $\sigma_s(\text{cal})=0.364$

The estimation is carried out under the conditions given below;

fragment	$\Sigma\sigma_s(\text{mono})$	$\sigma_s(\text{cal})$
a. N-C-C-C + 0.285	0.286 + 0.285 = 0.571	0.457
b. N-C-C + 0.312	0.241 + 0.312 = 0.553	0.444
c. N-C + 0.364	0.184 + 0.364 = 0.548	0.440

The mathematical average gives  $\sigma_s = 0.447$ .

Finally, the descriptor  $\sigma_s = 0.445$  is estimated for lysine.

Furthermore, the  $\pi$ -values of amino acids determined by Fauchère and Pliska<sup>7)</sup> were found to be expressed by the linear combination of the independent variable  $\sigma_s(R)$  or

$\sigma_s(\text{cal})$  and hydrogen bonding index ( $I=0, 1, 2$ ) as shown by Eq. 4.

$$\pi = 11.656(6.382)\sigma_s(\text{cal}) - 1.124(0.342)I + 2.789(2.184) \quad (4)$$

$$n = 16, r = 0.909, s = 0.442, F = 26.25$$

This result suggests that the hydrophobic substituent constant  $\pi$  could be expressed by the sum of contributions from the dispersion, repulsion, and averaged hydration number.

### Conclusion

A logical estimation of the QSAR descriptor  $\sigma_s$  for  $\text{CH}_2\text{R}_A\text{R}_B$ ,  $\text{CHR}_A\text{R}_B\text{R}_C$  and  $\text{CR}_A\text{R}_B\text{R}_C\text{R}_D$  could be carried out by means of the regression analyses using  $\Sigma\sigma_s(\text{mono})$  as an independent variable. The term  $\Sigma\sigma_s(\text{mono})$ , composed of the sum of fragments, should be taken as the maximum or mean value for the branched- or straight-chain substrate, and corrections in the presence of the plane of symmetry, *trans* and *gauche* isomers, and optical antipode should be taken into account.

This method could be extended for an aliphatic substrate having a substituted phenyl ring in the molecular framework, and successful evidence is observed in the evaluation of the  $\pi$ -value of amino acids.

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