



Correlation between molecular acidity (pK_a) and vibrational spectroscopy

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Abstract

Molecular acidity is an important physicochemical property, which is often represented by the pK_a value as the measure of acidity strength. However, the accurate calculation and prediction of pK_a values is still an unsolved problem for computational chemistry. In this work, we present for the first time a direct correlation between pK_a values and local vibrational frequencies for 15 different groups of compounds with various substituents. This correlation was derived from a quadratic function of two selected local vibrational frequencies as independent variables used to characterize electronic structure features influencing the molecular acidity. In total, 180 molecules were investigated with this correlation model. For each group of molecules, we found a strong correlation with root mean squared errors and mean absolute errors of less than 0.11 and 0.09 pK_a units, respectively. The correlation between pK_a and local vibrational modes, established in this work, can be generally applied to all compounds whose pK_a values are dominated by electronic substituent effects. In this regard, the new correlation model constitutes a powerful link between the well-known Hammett equation and vibrational spectroscopy. Furthermore, it allows a quick prediction of the pK_a values for new group members with different substituents.

Keywords Vibrational spectroscopy · pK_a value · Linear regression · Local vibrational mode analysis · Hammett equation

Niraj Verma and Yunwen Tao contributed equally to this work.

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Introduction

Molecular acidity, the ability or tendency of an acid to lose a proton, is one of the most fundamental and important physicochemical properties of a molecule, which is essential in many chemical and biological processes [1–17]. Upon dissociation, an acid releases a proton, making the solution acidic. This process is monitored by the equilibrium constant K_a of the corresponding dissociation reaction. For an acid HA, which dissociates into A^- and H^+ , the negative logarithm of K_a is the pK_a of the corresponding acid with $K_a = [H^+][A^-]/[HA]$. For a charged acid BH^+ that dissociates to B and H^+ , the pK_a is the negative logarithm of $K_a = [B][H^+]/[BH^+]$.

Despite the numerous attempts reported in the literature [5, 18–36], the accurate prediction of pK_a values by computational means is still an unsolved problem. A common approach is to use a thermodynamic cycle to calculate the pK_a by utilizing the standard Gibbs free energy change $\Delta G = 2.303 RT pK_a$ [24, 37], applying wave function or density functional theory (DFT) methodologies [18, 38]. This procedure may suffer from errors caused by (i) the setup of the thermodynamic cycle and (ii) the way solvent and proton hydration is treated [39, 40]. Continuum

solvation models based on the quantum mechanical charge density of a solute molecule interacting with a continuum description of the solvent [41], introduced in 2009, are frequently applied for the calculation of pK_a values. Several modifications to this approach were made [42–51], allowing to estimate pK_a values in some cases within an accuracy of ca. 1 pK_a unit. An alternative approach is based on ab initio molecular dynamics (AIMD) simulations [52]. AIMD-based pK_a calculations require the simulation of the dissociation process. The challenge is to run the simulation long enough to capture the final state of the dissociation to be used for the prediction of the pK_a value. Correlating experimentally determined pK_a values with calculated molecular properties has also attracted attention [53–61], as it has been successful in special cases leading to a mean absolute error (MAE) of less than 0.5 pK_a units [1, 62].

The pK_a value is influenced by several factors such as solvent, configuration of the molecule, substituents, etc. Pioneering work by Oszczapowicz and coworkers [62] verified a Hammett-type correlation for a series of substituted amidines, which was extended to substituted benzoic acid by Huang, Liu, and coworkers [63] and to the benzoxaborole pharmacophore by Benkovic and coworkers [64], just to name a few examples. Overall, substituent effects play an important role [65], as they change the electronic structure of a molecule, even if the substitution occurs at a distance from the dissociating proton. Changing the electronic structure of a molecule via a substituent, the stability of the conjugate base is changed, and in this way the pK_a value. We have shown in recent work [66, 67] that the local vibrational modes, introduced by Konkoli and Cremer [68], sensitively reflect all electronic structure changes in a molecule upon

substitution. Therefore, we expect a correlation between the local vibrational frequencies of the conjugate base and the pK_a value of a compound as sketched in Fig. 1.

The main objective of this work was to investigate whether the electronic structure changes resulting from different substituents are captured by some representative local vibrational frequencies of the conjugate base, and if so, a correlation between pK_a values and local vibrational frequencies exists. We tested this hypothesis for 15 different groups of molecules with different substituents R, (180 molecules in total), shown in Fig. 2. We selected five different pairs of representative local vibrational modes ω_1 and ω_2 illustrated in Fig. 3 and as described below. For each group, the local vibrational frequencies ω_1 and ω_2 were quadratically correlated to experimental pK_a values.

The correlation model applied in this work is defined in Eq. 1

$$pK_a = c_1 \times \omega_1 + c_2 \times \omega_2 + c_3 \times \omega_1^2 + c_4 \times \omega_2^2 + c_5 \times \omega_1 \omega_2 + c_6 \quad (1)$$

where c_i , $i \in \{1, \dots, 5, 6\}$ are the correlation constants for each group, which were linearly optimized with regard to experimentally known pK_a values. For any new group member with a different substituent R, the corresponding pK_a can then be calculated via Eq. 1 as long as ω_1 and ω_2 are available.

The paper is arranged as follows. In Sect. 2 we describe the computational methods used for the calculation of the representative local mode frequencies ω_1 and ω_2 including a short description of the local vibrational mode analysis. In Sect. 3 we discuss the results for a total of 180

Fig. 1 Relationship between pK_a values and local vibrational frequencies for a conjugate base of an acid

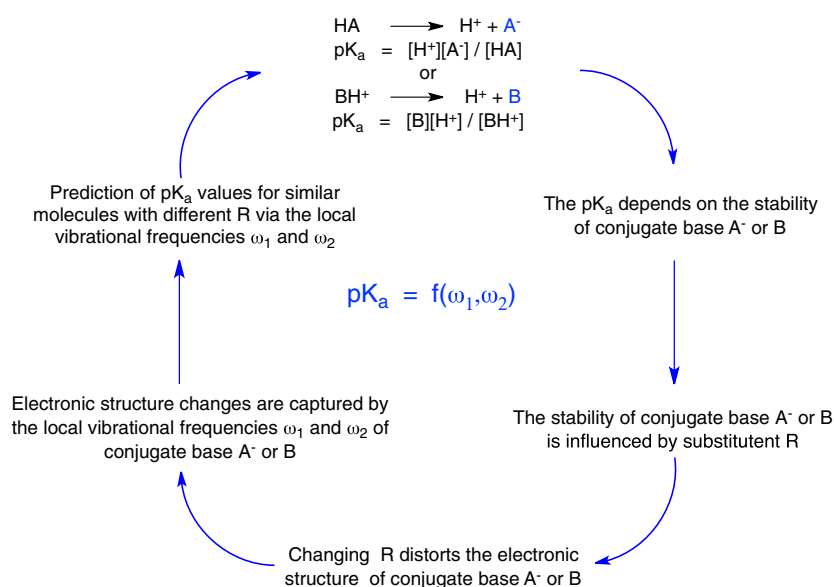
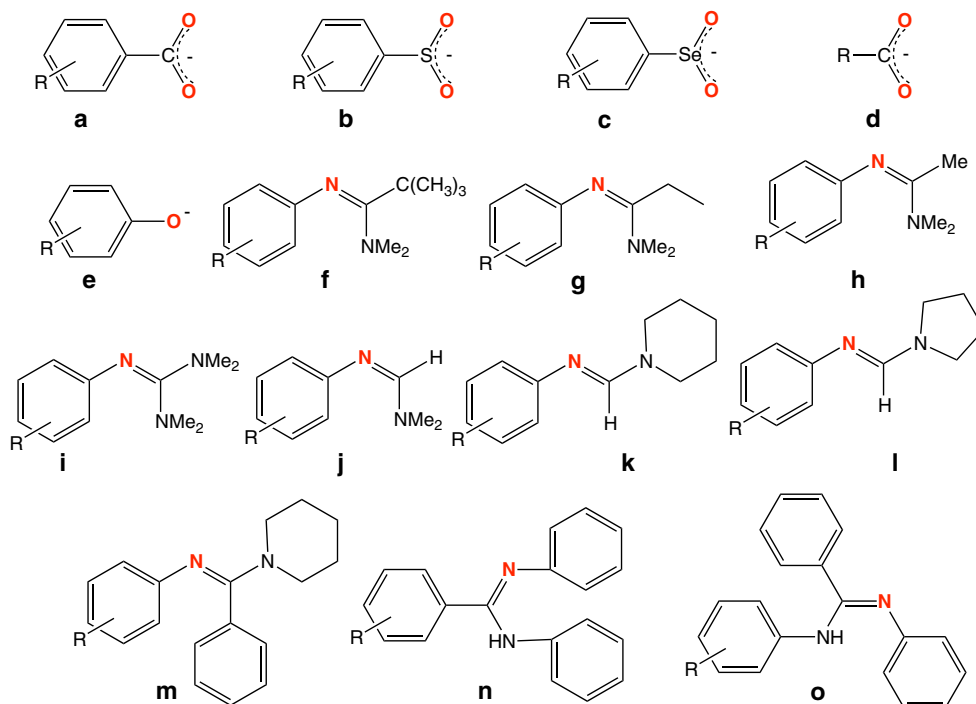


Fig. 2 15 groups (A-O) of molecules investigated in this work, different substituents R are defined in Table 1. The key atoms that are involved in the dissociation reaction and that are used for the definition of the representative local vibrational frequency pairs $p(\omega_1, \omega_2)$ are shown in red



different compounds belonging to one of the groups A-O, summarized in Table 1. We evaluate the accuracy of the calculated pK_a values via the correlation Eq. 1 and applying cross-validation and *t* test (see below) we analyze the predictive nature of these correlations. In the last section, we summarize our results and draw conclusions.

Computational details

Local vibrational modes

The basic equation of vibrational spectroscopy [69] is defined as

$$\mathbf{f}^x \mathbf{L} = \mathbf{M} \mathbf{L} \mathbf{\Lambda}$$

where \mathbf{f}^x is the force constant matrix expressed in Cartesian coordinates. \mathbf{M} is the mass matrix. The diagonal eigenvalue matrix $\mathbf{\Lambda}$ contains N_{vib} vibrational eigenvalues $4\pi^2 c^2 \nu_\mu^2$ (with $\mu = 1, \dots, N_{vib}$ and $N_{vib} = 3N - TR$; $TR = 6$ for nonlinear molecules and 5 for linear molecules) and TR zero eigenvalues corresponding to translations and rotations of the molecule. The harmonic vibrational frequencies ν_μ are given in cm^{-1} and c is the speed of light. The $(3N \times 3N)$ dimensional \mathbf{L} matrix collects the N_{vib} normal vibrational mode vectors \mathbf{I}_μ and TR mode vectors corresponding to translations and rotations.

Expressing Eq. 2 in internal coordinates \mathbf{q} leads to the Wilson GF formalism with [69, 70]:

$$\mathbf{F}^q \mathbf{D} = \mathbf{G}^{-1} \mathbf{D} \mathbf{\Lambda} \quad (2) \quad (3)$$

Fig. 3 Five different categories I, II, III, IV, and V of representative local vibrational frequency pairs $p(\omega_1, \omega_2)$ used in this work

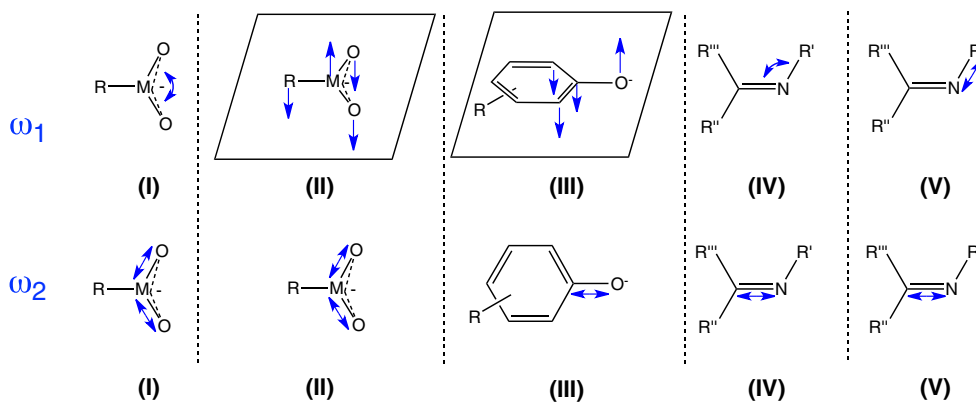


Table 1 Substituent R, local vibrational frequencies ω_1 and ω_2 used in Eq. 1, experimental and predicted pK_a values for all molecules of groups A-O, (180 in total). Diff is the difference between the experimental and predicted pK_a value

Group	Substituent R	ω_1 (cm ⁻¹)	ω_2 (cm ⁻¹)	pK _a (Experimental)	pK _a (Predicted)	Diff
Group A (category I)						
A-1	H	1560	899	4.20	4.23	0.03
A-2	m-Br	1568	895	3.81	3.80	-0.01
A-3	m-CN	1571	896	3.60	3.59	-0.01
A-4	m-Cl	1567	896	3.84	3.81	-0.03
A-5	m-F	1566	897	3.86	3.90	0.04
A-6	m-Me	1560	899	4.25	4.27	0.02
A-7	m-NO ₂	1572	895	3.46	3.42	-0.04
A-8	m-OH	1563	898	4.08	4.09	0.01
A-9	m-OMe	1563	898	4.10	4.07	-0.03
A-10	p-Br	1565	898	3.96	3.94	-0.02
A-11	p-CN	1570	896	3.55	3.58	0.03
A-12	p-Cl	1565	898	4.00	3.99	-0.01
A-13	p-F	1562	899	4.15	4.15	0.00
A-14	p-Me	1559	899	4.37	4.39	0.02
A-15	p-NMe ₂	1555	899	4.98	4.99	0.01
A-16	p-NO ₂	1572	895	3.43	3.49	0.06
A-17	p-OH	1558	899	4.47	4.45	-0.02
A-18	p-OMe	1558	899	4.50	4.46	-0.04
Group B (category I)						
B-1	H	1078	534	2.97	2.96	-0.02
B-2	m-Cl	1087	528	2.26	2.26	0.01
B-3	m-NO ₂	1084	516	1.67	1.67	0.00
B-4	p-Br	1083	530	2.51	2.44	-0.07
B-5	p-Cl	1082	532	2.51	2.58	0.07
B-6	p-Me	1078	536	3.12	3.13	0.01
B-7	p-NO ₂	1089	501	1.43	1.43	0.00
B-8	p-OMe	1078	543	3.84	3.83	-0.01
Group C (category II)						
C-1	H	819	367	4.79	4.72	-0.07
C-2	m-Br	823	366	4.43	4.45	0.02
C-3	m-Cl	823	366	4.47	4.46	-0.01
C-4	m-F	821	360	4.34	4.30	-0.04
C-5	m-Me	818	367	4.80	4.81	0.01
C-6	m-NO ₂	827	357	4.07	4.03	-0.04
C-7	m-OMe	819	361	4.65	4.68	0.03
C-8	p-Br	821	366	4.50	4.48	-0.02
C-9	p-Cl	820	367	4.48	4.51	0.03
C-10	p-F	819	369	4.50	4.47	-0.03
C-11	p-Me	818	368	4.88	4.85	-0.03
C-12	p-NO ₂	824	359	4.00	4.07	0.07
C-13	p-OMe	817	369	4.65	4.73	0.08
Group D (category II)						
D-1	ArCH ₂ CH ₂	1571	616	4.66	4.66	0.00
D-2	CH ₃	1568	512	4.76	4.74	-0.02
D-3	C ₂ H ₅	1569	550	4.87	4.84	-0.03
D-4	HC≡CCH ₂	1591	530	3.32	3.32	0.00
D-5	iPr	1566	595	4.86	4.94	0.08

Table 1 (continued)

Group	Substituent R	ω_1 (cm ⁻¹)	ω_2 (cm ⁻¹)	pK _a (Experimental)	pK _a (Predicted)	Diff
D-6	n-C ₄ H ₉	1570	565	4.84	4.84	0.00
D-7	n-C ₅ H ₁₁	1570	436	4.88	4.87	-0.01
D-8	n-C ₆ H ₁₃	1570	440	4.89	4.91	0.02
D-9	n-C ₉ H ₁₉	1570	549	4.89	4.85	-0.04
D-10	n-Pr	1569	546	4.76	4.84	0.08
D-11	tBu	1565	602	5.05	4.99	-0.06
Group E (category III)						
E-1	H	1535	731	9.98	10.02	0.04
E-2	m-Br	1556	724	9.03	8.99	-0.04
E-3	m-CH ₃	1536	730	10.08	9.97	-0.11
E-4	m-CN	1558	721	8.61	8.74	0.13
E-5	m-Cl	1553	727	9.02	9.16	0.14
E-6	m-F	1548	734	9.28	9.31	0.03
E-7	m-NH ₂	1537	735	9.87	9.84	-0.03
E-8	m-NO ₂	1559	716	8.40	8.32	-0.08
E-9	m-OCH ₃	1542	733	9.65	9.64	-0.01
E-10	m-OH	1544	732	9.44	9.57	0.13
E-11	p-Br	1550	726	9.36	9.27	-0.09
E-12	p-CH ₃	1531	727	10.14	10.11	-0.03
E-13	p-CN	1583	729	7.95	7.81	-0.14
E-14	p-Cl	1546	726	9.38	9.45	0.07
E-15	p-F	1526	724	9.95	10.15	0.20
E-16	p-NH ₂	1531	726	10.30	10.05	-0.25
E-17	p-NO ₂	1599	726	7.15	7.21	0.06
E-18	p-OCH ₃	1519	720	10.21	10.07	-0.14
E-19	p-OH	1514	718	9.96	10.07	0.11
Group F (category V)						
F-1	H	1572	1188	8.03	8.09	0.06
F-2	m-Br	1562	1193	6.81	6.77	-0.04
F-3	m-CH ₃	1574	1187	8.24	8.21	-0.03
F-4	m-Cl	1563	1192	7.01	6.97	-0.04
F-5	m-NO ₂	1553	1196	5.86	5.88	0.02
F-6	m-OCH ₃	1571	1187	7.91	7.86	-0.05
F-7	m-OC ₂ H ₅	1573	1187	7.93	8.11	0.18
F-8	p-Br	1566	1192	7.20	7.27	0.07
F-9	p-CH ₃	1573	1188	8.43	8.28	-0.15
F-10	p-Cl	1567	1193	7.19	7.24	0.05
F-11	p-I	1566	1193	7.17	7.11	-0.06
F-12	p-NO ₂	1550	1199	5.18	5.18	0.00
F-13	p-OCH ₃	1575	1187	8.62	8.56	-0.06
F-14	p-OC ₂ H ₅	1576	1186	8.63	8.67	0.04
Group G (category V)						
G-1	H	1579	1192	8.30	8.19	-0.11
G-2	m-Br	1571	1197	7.13	7.13	0.00
G-3	m-CH ₃	1581	1190	8.38	8.44	0.06
G-4	m-Cl	1572	1197	7.15	7.21	0.06
G-5	m-NO ₂	1559	1202	6.06	6.05	-0.01
G-6	m-OCH ₃	1579	1192	8.13	8.17	0.04

Table 1 (continued)

Group	Substituent R	ω_1 (cm ⁻¹)	ω_2 (cm ⁻¹)	pK _a (Experimental)	pK _a (Predicted)	Diff
G-7	m-OC ₂ H ₅	1580	1192	8.16	8.29	0.13
G-8	p-Br	1575	1198	7.43	7.46	0.03
G-9	p-CH ₃	1580	1190	8.44	8.36	-0.08
G-10	p-Cl	1575	1197	7.55	7.48	-0.07
G-11	p-I	1574	1198	7.36	7.33	-0.03
G-12	p-NO ₂	1564	1205	5.65	5.66	0.01
G-13	p-OCH ₃	1579	1186	8.94	9.00	0.06
G-14	p-OC ₂ H ₅	1580	1187	8.89	8.80	-0.09
Group H (category V)						
H-1	H	1578	1187	8.32	8.36	0.04
H-2	m-Br	1568	1194	7.19	7.19	0.00
H-3	m-CH ₃	1579	1185	8.41	8.56	0.15
H-4	m-Cl	1569	1193	7.25	7.26	0.01
H-5	m-OCH ₃	1577	1188	8.22	8.13	-0.09
H-6	m-OC ₂ H ₅	1578	1188	8.26	8.31	0.05
H-7	p-Br	1572	1192	7.55	7.54	-0.01
H-8	p-CH ₃	1580	1186	8.65	8.53	-0.12
H-9	p-Cl	1573	1192	7.65	7.66	0.01
H-10	p-NO ₂	1558	1203	5.69	5.69	0.00
H-11	p-OCH ₃	1580	1183	8.96	8.91	-0.05
H-12	p-OC ₂ H ₅	1581	1183	8.90	8.91	0.01
Group I (category IV)						
I-1	H	1535	567	11.52	11.55	0.03
I-2	m-CH ₃	1535	564	11.74	11.71	-0.03
I-3	m-Cl	1520	564	10.55	10.52	-0.03
I-4	m-OCH ₃	1531	564	11.44	11.47	0.03
I-5	m-OC ₂ H ₅	1537	558	11.38	11.38	0.00
I-6	p-CH ₃	1538	565	11.94	11.85	-0.09
I-7	p-Cl	1526	564	10.98	11.04	0.06
I-8	p-OCH ₃	1544	570	12.16	12.16	0.00
I-9	p-OC ₂ H ₅	1543	565	12.08	12.13	0.05
Group J (category IV)						
J-1	H	1620	524	7.45	7.31	-0.14
J-2	m-Br	1614	518	6.45	6.42	-0.03
J-3	m-CH ₃	1621	528	7.63	7.65	0.02
J-4	m-Cl	1615	521	6.50	6.62	0.12
J-5	m-OCH ₃	1621	523	7.45	7.51	0.06
J-6	m-OC ₂ H ₅	1605	600	7.45	7.47	0.02
J-7	p-Br	1614	523	6.69	6.61	-0.08
J-8	p-CH ₃	1608	598	7.75	7.73	-0.02
J-9	p-Cl	1615	526	6.84	6.88	0.04
J-10	p-NO ₂	1584	590	5.25	5.25	0.00
J-11	p-OCH ₃	1621	539	7.91	7.95	0.04
J-12	p-OC ₂ H ₅	1620	538	7.83	7.78	-0.05
Group K (category IV)						
K-1	H	1601	595	8.69	8.60	-0.09
K-2	m-Br	1590	600	7.70	7.70	0.00
K-3	m-CH ₃	1602	596	8.74	8.83	0.09

Table 1 (continued)

Group	Substituent R	ω_1 (cm ⁻¹)	ω_2 (cm ⁻¹)	pK _a (Experimental)	pK _a (Predicted)	Diff
K-4	m-Cl	1592	594	7.83	7.84	0.01
K-5	m-OCH ₃	1599	598	8.47	8.52	0.05
K-6	p-Br	1593	597	7.95	7.95	0.00
K-7	p-CH ₃	1603	601	9.10	9.10	0.00
K-8	p-Cl	1595	599	8.09	8.06	-0.03
K-9	p-OCH ₃	1604	597	9.38	9.34	-0.04
K-10	p-OC ₂ H ₅	1606	592	9.10	9.11	0.01
Group L (category V)						
L-1	H	1600	1194	9.31	9.41	0.10
L-2	m-Br	1593	1200	8.46	8.40	-0.06
L-3	m-CH ₃	1601	1193	9.53	9.56	0.03
L-4	m-Cl	1593	1201	8.52	8.59	0.07
L-5	m-OCH ₃	1599	1192	9.22	9.22	0.00
L-6	p-Br	1594	1201	8.75	8.76	0.01
L-7	p-CH ₃	1602	1193	9.75	9.61	-0.14
L-8	p-Cl	1595	1200	8.85	8.82	-0.03
L-9	p-OCH ₃	1605	1191	9.97	9.98	0.01
L-10	p-OC ₂ H ₅	1605	1192	9.70	9.71	0.01
Group M (category V)						
M-1	H	1564	1189	7.45	7.51	0.06
M-2	m-Br	1552	1195	6.38	6.38	0.00
M-3	m-CH ₃	1564	1187	7.72	7.66	-0.06
M-4	m-Cl	1554	1195	6.45	6.46	0.01
M-5	m-OCH ₃	1563	1189	7.35	7.43	0.08
M-6	m-OC ₂ H ₅	1564	1189	7.58	7.56	-0.02
M-7	p-Br	1557	1195	6.65	6.67	0.02
M-8	p-CH ₃	1565	1187	7.90	7.83	-0.07
M-9	p-Cl	1558	1194	6.77	6.73	-0.04
M-10	p-OCH ₃	1567	1185	8.19	8.21	0.02
M-11	p-OC ₂ H ₅	1567	1185	8.17	8.19	0.02
Group N (category V)						
N-1	H	1593	1187	14.65	14.56	-0.09
N-2	m-Br	1596	1186	13.81	13.85	0.04
N-3	m-NO ₂	1596	1186	13.38	13.37	-0.01
N-4	p-Br	1594	1187	14.05	14.10	0.05
N-5	p-CH ₃	1591	1187	14.92	14.96	0.04
N-6	p-NO ₂	1596	1185	13.19	13.17	-0.02
N-7	p-OCH ₃	1588	1188	15.27	15.27	0.00
Group O (category V)						
O-1	H	1594	1173	6.15	6.16	0.01
O-2	m-Br	1591	1178	5.25	5.26	0.01
O-3	m-CH ₃	1596	1171	6.36	6.51	0.15
O-4	m-Cl	1591	1178	5.23	5.28	0.05
O-5	m-NO ₂	1589	1184	4.24	4.29	0.05
O-6	m-OCH ₃	1598	1173	6.04	6.01	-0.03
O-7	p-Br	1590	1178	5.42	5.38	-0.04
O-8	p-CH ₃	1595	1172	6.52	6.39	-0.13

Table 1 (continued)

Group	Substituent R	ω_1 (cm ⁻¹)	ω_2 (cm ⁻¹)	pK _a (Experimental)	pK _a (Predicted)	Diff
O-9	p-Cl	1590	1178	5.46	5.46	0.00
O-10	p-F	1591	1174	5.90	6.08	0.18
O-11	p-I	1590	1179	5.43	5.26	-0.17
O-12	p-OCH ₃	1593	1170	6.78	6.70	-0.08

where

$$\mathbf{G} = \mathbf{B}\mathbf{M}^{-1}\mathbf{B}^\dagger \quad (4)$$

\mathbf{F}^Q is the force constant matrix expressed in terms of internal coordinates \mathbf{q} . Each normal mode vector \mathbf{d}_μ represents a column vector of the ($N_{\text{vib}} \times N_{\text{vib}}$) dimensional \mathbf{D} matrix, which is defined as the product $\mathbf{B}\mathbf{L}$. The rectangular ($N_{\text{vib}} \times 3N$) dimensional \mathbf{B} matrix contains the first derivatives of the internal coordinates with regard to the Cartesian coordinates, thus connecting both coordinate systems, and the ($N_{\text{vib}} \times N_{\text{vib}}$) dimensional matrix \mathbf{G} is the Wilson \mathbf{G} matrix. [69, 70].

The normal mode frequencies ν_μ are coupled caused by electronic coupling (off-diagonal elements of the \mathbf{F}^Q matrix, which can be eliminated via the Wilson GF formalism) and mass-coupling (caused by the off-diagonal elements of the Wilson \mathbf{G} matrix) [69, 70]. However, for our correlation, we need local vibrational modes focusing on the influence of the substituent R on the electronic structure of the conjugated base. Konkoli and Cremer [68] solved this problem by introducing the concept of local vibrational modes. They solved the mass-decoupled Euler–Lagrange equation by setting all the atomic masses to zero except those of the molecular fragment (e.g., bond, angle, or dihedral, etc.) carrying out a localized vibration. As shown in their original work, the change in the local displacement of a specific internal coordinate is equivalent to an adiabatic relaxation of the molecule [68]. For any molecular fragment associated with an internal coordinate q_n , the corresponding local mode vector \mathbf{a}_n is given by

$$\mathbf{a}_n = \frac{\mathbf{K}^{-1}\mathbf{d}_n^\dagger}{\mathbf{d}_n^\dagger\mathbf{K}^{-1}\mathbf{d}_n^\dagger} \quad (5)$$

where \mathbf{d}_n contrary to \mathbf{d}_μ is a row vector of matrix \mathbf{D} . Matrix \mathbf{K} is the diagonal matrix of force constants k_Q , expressed in normal coordinates Q_μ with

$$\mathbf{F}^Q = \mathbf{K} = \mathbf{L}^\dagger \mathbf{f}^x \mathbf{L} \quad (6)$$

resulting from the Wilson GF formalism shown in Eqs. 3 and 4.

The local mode force constant k_n^a corresponding to local mode \mathbf{a}_n is obtained by

$$k_n^a = \mathbf{a}_n^\dagger \mathbf{K} \mathbf{a}_n \quad (7)$$

The local vibrational frequency ω_n^a corresponding to local mode \mathbf{a}_n is obtained by

$$(\omega_n^a)^2 = \frac{1}{4\pi^2 c^2} k_n^a G_{nn} \quad (8)$$

in which G_{nn} is a diagonal element of the Wilson \mathbf{G} -matrix and corresponds to the reduced mass of the local mode \mathbf{a}_n [68].

Local vibrational modes have successfully been applied to quantify weak chemical interactions such as hydrogen bonding [71–76], halogen bonding [77–79], pnictogen bonding [80–82], chalcogen bonding [83] and tetrel bonding [84], and to derive new chemical descriptors such as a new aromaticity index [85–87] or a generalized Tolman electronic parameter [88–90], as well as for the derivation of a generalized Badger Rule [91] and several others new concepts [91–97].

Geometry optimizations and normal mode calculations for all molecules investigated in this work were performed in the gas phase with the ω B97XD density functional [98] using the Gaussian 16 quantum chemistry program [99, 100]. For members of groups B, D, E, F, G, H, I, J, K, L, M, N, and O (see Fig. 2) Dunning's cc-pVTZ basis set [101–103] was applied, while for members of group A and C (see Fig. 2) Pople's 6-31++G(d,p) basis set [104–106] turned out to be the best choice. Experimental pK_a values were taken from ref [1, 62]. There are no reliable experimental pK_a values for sulphinic acid available. Therefore, we used computed pK_a values instead [107]. The local mode analysis was performed with the COLOGNE2017 program package [108].

Results and discussion

Correlation of pK_a and local vibrational frequencies

Vibrational frequencies are second-order response properties [109] and therefore they are sensitive to any changes in the electronic structure of a molecule. First it was important to determine the local vibrational modes that best reflect the influence of the substituent R on the dissociation reaction for each molecule investigated in this work.

Depending on the molecule, there exist two or three vibrational modes near the dissociating proton that will dominate the pK_a value. As there are some redundancies between stretching, bending, and pyramidalization modes at a specific molecular site with regard to the electronic structure change during the vibration, one has to remove either one in order to obtain a robust model. Testing ${}^3C_2 = 3$ different combinations, we identified the set of parameters that performs best and leads to the desired model. This led to the five different categories of local vibrational mode pairs $p(\omega_1, \omega_2)$ shown in Fig. 3 used in this study. For category I molecules, a pair of local bending and stretching vibrations turned out to be most sensitive; for category II and category III molecules, a pair of local pyramidalization and stretching vibrations; for category IV molecules, a pair of local bending and stretching vibrations, and for category V molecules, two local stretching vibrations. The two local stretching vibrations for category I and II molecules were averaged. Only one of the substituent (R or R' or R'' or R''') was changed at a time. The 15 groups A-O of molecules used for the correlation comprise the conjugate bases of (A) benzoic acid; (B) benzene sulphinic acid; (C) benzeneselenic acids; (D) alkyl carboxylic acids; (E) phenols; (F) protonated N^1, N^1 -dimethyltertiarybutylamide; (G) protonated N^1, N^1 -dimethylethylamide; (H) protonated N^1, N^1 -dimethylethylamine; (I) protonated N^2 -alkyl-tetramethylguanidine; (J) protonated N^1, N^1 -dimethylformamide; (K) protonated N^1, N^1 -(pentamethylene-1,5)-formamide; (L) protonated N^1, N^1 -(butamethylene-1,4)-formamide; (M) protonated N^1, N^1 -(pentamethylene-1,5)-benzamide; (N) protonated N, N' -diphenyl-benzamide; (O) protonated N, N' -diphenyl-benzamide, shown in Fig. 2.

In the following, the results of our investigation are discussed for each individual group, referring to the data in Table 1, which includes for each individual compound the substituent R, the local vibrational mode frequencies ω_1 and ω_2 used in Eq. 1, experimental pK_a and predicted pK_a values, as well as the difference *DIFF* between experimental and predicted pK_a values. The correlation protocol for groups A-O is presented in Table 2 including for each group the number of molecules, the root mean square error (RMSE), the mean average error (MAE), the mean average derivation (MAD), and the R^2 value, as a statistical measure of how close the data are to the fitted regression line.

Group A consists of 18 different compounds, which all fall into the category I of local mode frequency pairs $p(\omega_1, \omega_2)$, as shown in Table 1. The substituents vary from electron donating groups (EDG) to electron withdrawing groups (EWG), which accordingly change the electronic structure of the parent compound ($R = H$) over a wide range, as such influencing the pK_a value by

stabilizing/destabilizing the corresponding conjugate base. We obtained a strong correlation with R^2 greater than 0.99 and a MAE of 0.025 pK_a units for this group, reflecting the fact that the pK_a values of group A molecules are dominated by substituent effects. It has to be noted that the error of the experimental data is as high as ± 0.09 pK_a units as reported in ref [1], which may amplify the predicted error due to quadratic fitting (QFE). **Group B** consists of eight different compounds and shows a MAD of 0.578. Despite this large MAD value, we observed a strong correlation with R^2 of 1.0 and a MAE of 0.02. A similarly strong correlation was found for **group C** and **group D** molecules, again reflecting that the pK_a values of these groups members are dominated by substituent effects. **Group E** shows the two largest deviations between calculated and experimental pK_a values of all molecules investigated in this work, namely for $R = p\text{-F}$ and $R = p\text{-NH}_2$. Apart from the QFE, another reason for these deviations might be the model chemistry used. The optimal level of theory and basis set for each molecule varies according to the type of substituent R involved. We tested different levels of theory and basis sets, which led to somewhat different outliers, (see supporting information for a detailed report).

Group F, G and **H** members are characterized by a substituent R'' at the C atom of the $C=N$ bond with a decreasing number of C atoms, e.g., **group F**: $-\text{C}(\text{CH}_3)_3$, **group G**: $-\text{C}_2\text{H}_5$ and **group H** $-\text{CH}_3$, respectively. The number of carbon atoms in R'' determine changes in the electronic structure caused by inductive effects, which is reflected by the local $C=N$ double bond stretching frequency. Largest inductive effects were found for **group G** and **group F** molecules, in particular for molecules G-7 and F-7, substituent $m\text{-OC}_2\text{H}_5$, the outliers of these groups. In **group F** also the $-\text{CH}_3$ functional group F-9 led to an outlier, caused by an increased resonance effect, which we also found for molecule H-8 of **Group H**. **Group I** molecules show a similar deviations as found for **group H** members, due to resonance effects. **Group J** and **group K** show the largest outliers for the parent compound ($R = H$), molecules J-1 and K-1, respectively. As all other substituents in these two groups are either strong EDG or EWG functional groups, $R = H$ deviates from the general group pattern. In **Group L** and **group M** the $p\text{-CH}_3$ functional group leads again to the largest outliers. Due to hyperconjugation, the $p\text{-CH}_3$ substituent increases the electron density at the *para* position, and in this way strongly influences the electronic structure. **Group O** has several outliers. Nevertheless, we observe a strong correlation with R^2 of almost 0.98. The outliers in this group can be attributed to both the QFE and the model chemistry used.

The electronic structure is affected by resonance, inductive effects or by hyperconjugation. It is remarkable to see that even changes in substituents at distance from

Table 2 Correlation protocol for each group A-O

Group	Number of molecules	RMSE ^a	MAE ^b	MAD ^c	R ² ^d
Group A	18	0.028	0.025	0.311	0.995
Group B	8	0.035	0.022	0.578	0.998
Group C	13	0.044	0.038	0.192	0.970
Group D	11	0.041	0.031	0.261	0.991
Group E	19	0.114	0.096	0.637	0.981
Group F	14	0.077	0.060	0.812	0.994
Group G	12	0.067	0.056	0.779	0.995
Group H	12	0.066	0.045	0.712	0.994
Group I	9	0.043	0.035	0.398	0.992
Group J	12	0.065	0.052	0.628	0.992
Group K	10	0.047	0.033	0.497	0.993
Group L	10	0.062	0.046	0.449	0.985
Group M	11	0.043	0.036	0.557	0.995
Group N	7	0.044	0.035	0.656	0.996
Group O	12	0.098	0.075	0.560	0.979

^bMAE = $\frac{1}{n} \sum |y_j^e - y_j^p|$, ^aRMSE = $\sqrt{\frac{1}{n} \sum (y_j^e - y_j^p)^2}$, ^cMAD = $\frac{1}{n} \sum |y_j^e - \bar{y}_j^e|$ \bar{y}_j^e equals the pK_a; superscript e denotes experimental, p predicted; \bar{y}_j^e is the mean of the experimental pK_a values, j equals the number of molecules in a group.

^dR² is the statistical measure for how close the data points are to the fitted regression line

the protonation center can be captured by the two local vibrational reference frequencies ω_1 , and ω_2 involving the protonation center. It is also noteworthy that although all predicted pK_a values were derived from gas phase calculations, while the experimental pK_a values were measured in solution, the local vibrational modes calculated in the gas phase capture already the important electronic structure changes influencing the pK_a values. This allows for a quick check of the pK_a values for new group members with different substituents R.

General trends

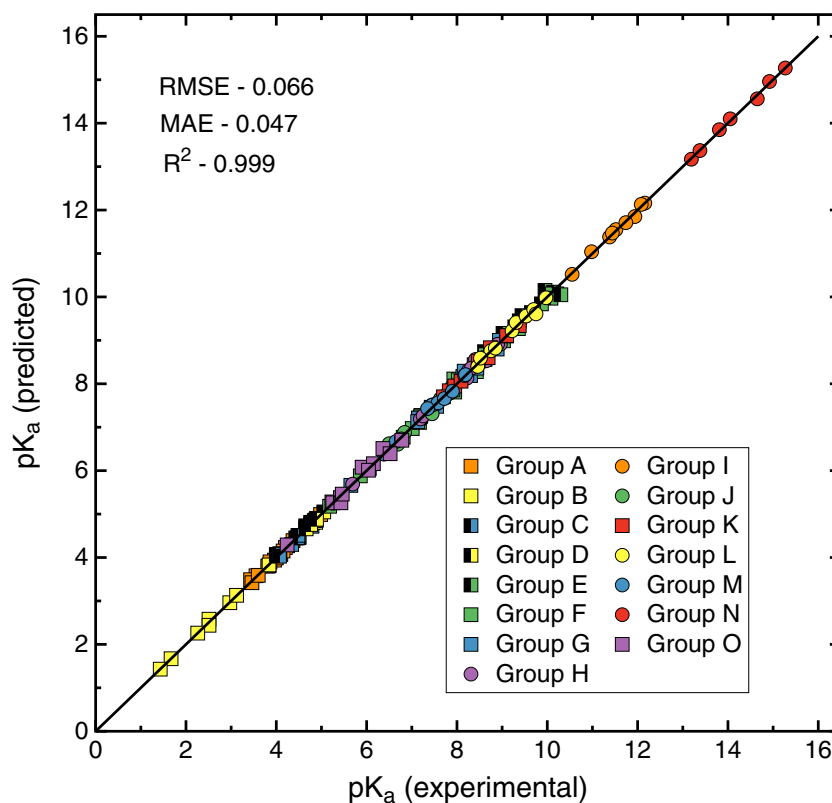
As reflected by the data in Table 2, each group shows a very strong correlation with an R² value greater than 0.97 and a MAE of less than 0.09. By inspection of the correlation constants in Eq. 1 (see supporting information for further details), one can easily see that there is however no relation between the different groups, and hence an overall correlation is not meaningful. For each particular group, the changes of the electronic structure upon substitution leading to a change in the molecular acidity are dominated by the substituent. Since the local vibrational modes can capture the sensitive changes in the electronic structure of conjugate base, we obtain a significant correlation between pK_a values and local vibrations. Figure 4 shows the predicted and experimental pK_a values. The results for MAE, RMSE, and R², which are 0.05, 0.07, and 0.999, respectively, confirm our hypothesis that there exists a strong correlation

between pK_a and local vibrational frequencies. In order to check if this correlation results predominantly from an electronic effect, we also correlated local mode force constants (Eq. 7) with experimental pK_a values, because the local mode force constants are free from any mass effects. We observed similar results as for the local vibrational frequencies, (detailed information can be found in the supporting information).

Validation of the correlation

In order to test the predictive power of our new correlation model, we performed a k-fold cross-validation for molecules similar to group E members, including 33 compounds of singly and doubly substituted benzenes, (shown in the supporting information). Cross-validation is a reliable means to check the efficiency of predicting models. In a k-fold cross-validation, data is randomly distributed to k splits as shown in Fig. 5. A model is then trained for k-1 splits and tested for the left out split. Any extra data point(s) is randomly assigned to one of the splits. This is repeated k times and the final error (MAE or RMSE) is averaged to get k-fold cross-validation error. We performed a four-fold cross-validation for this group and obtained a cross-validation MAE of 0.05, which equals the actual MAE for this group. This proves that our model is predictive in nature and can be used to predict pK_a values for any new group member with a different substituent R for any group of molecules, provided we have access to sufficient

Fig. 4 Correlation between experimental and predicted pK_a values



experimental pK_a values and the representative local mode pairs $p(\omega_1, \omega_2)$ for a variety of substituted molecules of this group, so that Eq. 1 can be solved.

The t test is a measure of statistical significance between two different data distributions, which can be computed as

$$t_{value} = \frac{\bar{x}_1 - \bar{x}_2}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}} \quad (9)$$

where, \bar{x} , s and n are mean, standard deviation and variance of the data. Subscript 1 and 2 denotes the first data set and the second data set, respectively.

We calculated t_{value} for all pairs of coefficients (Group X coefficients (data set 1) with Group Y coefficients (data set 2), where $X, Y \in [A, B...O]$). The degrees of freedom can be calculated as

$$df = l_1 + l_2 - 2 \quad (10)$$

where, l_1, l_2 are the number of data points in data set 1 and data set 2, respectively. With the degrees of freedom of 10 (6+6-2) and 95% confidence, we keep the null hypothesis (the coefficients are statistically similar) if the t_{value} is less than 2.228 (two tailed t -distribution table). The maximum t_{value} was 1.5465, which confirms the null hypothesis, i.e.,



Fig. 5 k-fold cross-validation. E is the error in each iterations

there is no statistically significant difference between each of the two models.

Conclusions and outlook

Our investigation shows that the changes in electronic structure caused by different substituent R are captured by local vibrational frequencies related to the conjugate base. We verified that there exists a relationship between the local vibrational modes and the pK_a of a molecule provided substituent effects play the major role in determining the pK_a value. Fifteen different groups of molecules with various substituents (a total of 180 molecules) were investigated and showed strong correlations. The slopes and intercepts obtained in the correlation model were different for each group, which reflects that pK_a values are system dependent. However, within a group, the system dependency of pK_a value is removed and we can predict the pK_a value of any new group members with different substituents R.

Using cross-validation and *t* test, we showed that our model is predictive in nature and offers a lot of potential for the pK_a prediction for other groups of molecules, which is currently under investigation, including the following two examples. (i) Huge efforts are devoted to capture CO and CO₂ and to store these gases in an efficient manner [110–112]. Amine solutions play an important role in this regard [113–115]. The pK_a values of these amines are important indicators for their adsorption efficiency [116, 117]. A wide range of pK_a values have been collected for amine species [61, 117–120], therefore we can apply our correlation model to estimate the pK_a values of novel amine substituted amines with a potentially higher adsorption efficiency. (ii) The ionic form of a weak acid/base varies across a range of pH values. This is important in physiological systems, in which the ionization state affects the rate of diffusion across membranes. The pK_a influences permeability, protein binding, solubility, etc., which in turn affects absorption, metabolism, excretion, etc., of potential drug candidates [121–126]. Due to this connection, the pK_a plays an important role in engineering optimal pharmacokinetic characteristics. A large number of drug candidates have already been studied [121, 122] leading to a wealth of experimental pK_a values, serving as the basis for our correlation model.

In summary, our new correlation model constitutes a powerful link between the well-known Hammett equation and vibrational spectroscopy. From a practical perspective, one needs only to calculate the local vibrational modes in the gas phase. The correlation with experimental pK_a values removes the system dependency and leads to an equation that allows a quick check of the pK_a value of

a new compound with a different substituent R, avoiding any extensive pK_a calculation. This makes our approach the perfect tool for the engineering of compounds with a specific molecular acidity regulated by substituent effects.

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