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Predicting p K_a values from EEM atomic charges

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Abstract

The acid dissociation constant pK_a is a very important molecular property, and there is a strong interest in the development of reliable and fast methods for pK_a prediction. We have evaluated the pK_a prediction capabilities of QSPR models based on empirical atomic charges calculated by the Electronegativity Equalization Method (EEM). Specifically, we collected 18 EEM parameter sets created for 8 different quantum mechanical (QM) charge calculation schemes. Afterwards, we prepared a training set of 74 substituted phenols. Additionally, for each molecule we generated its dissociated form by removing the phenolic hydrogen. For all the molecules in the training set, we then calculated EEM charges using the 18 parameter sets, and the QM charges using the 8 above mentioned charge calculation schemes. For each type of QM and EEM charges, we created one QSPR model employing charges from the non-dissociated molecules (three descriptor QSPR models), and one QSPR model based on charges from both dissociated and non-dissociated molecules (QSPR models with five descriptors). Afterwards, we calculated the quality criteria and evaluated all the QSPR models obtained. We found that QSPR models employing the EEM charges proved as a good approach for the prediction of p K_a (63% of these models had $R^2 > 0.9$, while the best had $R^2 = 0.924$). As expected, QM QSPR models provided more accurate pK_a predictions than the EEM QSPR models but the differences were not significant. Furthermore, a big advantage of the EEM QSPR models is that their descriptors (i.e., EEM atomic charges) can be calculated markedly faster than the QM charge descriptors. Moreover, we found that the EEM QSPR models are not so strongly influenced by the selection of the charge calculation approach as the QM QSPR models. The robustness of the EEM QSPR models was subsequently confirmed by cross-validation. The applicability of EEM QSPR models for other chemical classes was illustrated by a case study focused on carboxylic acids. In summary, EEM QSPR models constitute a fast and accurate pK_a prediction approach that can be used in virtual screening.

Keywords: Dissociation constant, Quantitative structure-property relationship, QSPR, Partial atomic charges, Electronegativity equalization method, EEM, Quantum mechanics, QM

Background

The acid dissociation constant pK_a is an important molecular property, and its values are of interest in pharmaceutical, chemical, biological and environmental research. The pK_a values have found application in many areas, such as the evaluation and optimization of candidate drug molecules [1-3], ADME profiling [4,5], pharmacokinetics [6], understanding of protein-ligand interactions [7,8], etc.. Moreover, the key physicochemical properties

like lipophilicity, solubility, and permeability are all pK_a dependent. For these reasons, pK_a values are important for virtual screening. Therefore, both the research community and pharmaceutical companies are interested in the development of reliable and above all fast methods for pK_a prediction.

Several approaches for pK_a prediction have been developed [8-11], namely LFER (Linear Free Energy Relationships) methods [12,13], database methods, decision tree methods [14], ab initio quantum mechanical calculations [15,16], ANN (artificial neural networks) methods [17] or QSPR (quantitative structure-property relationship) modelling [18-20]. However, pK_a values remain one of the most challenging physicochemical properties to predict.

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A promising approach for pK_a prediction is to use QSPR models which employ partial atomic charges as descriptors [21-24].

The partial atomic charges cannot be determined experimentally and they are also not quantum mechanical observables. For this reason, the rules for determining partial atomic charges depend on their application (e.g. molecular mechanics energy, pK_a etc.), and many different methods have been developed for their calculation. The charge calculation methods can be divided into two main groups, namely quantum mechanical (QM) approaches and empirical approaches.

The quantum mechanical approaches first calculate a molecular wave function by a combination of some theory level (e.g., HF, B3LYP, MP2) and basis set (e.g., STO-3G, 6–31G*), and then partition this wave function among the atoms (i.e., the assignment of a specific part of the molecular electron density to each atom). This partitioning can be done using an orbital-based population analysis, such as MPA (Mulliken population analysis) [25,26], Löwdin population analysis [27] or NPA (natural population analysis) [28]. Other partitioning approaches are based on a wavefunction-dependent physical observable. Such approaches are, for example, AIM (atoms in molecules) [29], Hirshfeld population analysis [30] and electrostatic potential fitting methods like CHELPG [31] or MK (Merz-Singh-Kollman) [32]. Another partitioning method is the mapping of QM atomic charges to reproduce charge-dependent observables (e.g., CM1, CM2, CM3 and CM4) [33].

Empirical approaches determine partial atomic charges without calculating a quantum mechanical wave function for the given molecule. Therefore they are markedly faster than QM approaches. One of the first empirical approaches developed, CHARGE [34], performs a breakdown of the charge transmission by polar atoms into one-bond, two-bond, and three-bond additive contributions. Most of the other empirical approaches have been derived on the basis of the electronegativity equalization principle. One group of these empirical approaches invoke the Laplacian matrix formalism, and result in a redistribution of electronegativity. Such methods are PEOE (partial equalization of orbital electronegativity) [35], GDAC (geometry-dependent atomic charge) [36], KCM (Kirchhoff charge model) [37], DENR (dynamic electronegativity relaxation) [38] or TSEF (topologically symmetric energy function) [38]. The second group of approaches use full equalization of orbital electronegativity, and such approaches are, for example, EEM (electronegativity equalization method) [39], QEq (charge equilibration) [40] or SQE (split charge equilibration) [41]. The empirical atomic charge calculation approaches can also be divided into 'topological' and 'geometrical'. Topological charges are calculated using the 2D structure of the molecule, and they are conformationally independent (i.e., CHARGE,

PEOE, KCM, DENR, and TSEF). Geometrical charges are computed from the 3D structure of the molecule and they consider the influence of conformation (i.e., GDAC, EEM, Qeq, and SQE).

The prediction of pK_a using QSPR models which employ QM atomic charges was described in several studies [21-24], which have analyzed the precision of this approach and compared the quality of QSPR models based on different QM charge calculation schemes. All these studies show that QM charges are successful descriptors for p K_a prediction, as the QSPR models based on QM atomic charges are able to calculate pK_a with high accuracy. The weak point of QM charges is that their calculation is very slow, as the computational complexity is at least $\theta(E^4)$, where E is the number of electrons in the molecule. Therefore, pK_a prediction by QSPR models based on QM charges cannot be applied in virtual screening, as it is not feasible to compute QM atomic charges for hundreds of thousands of compounds in a reasonable time. This issue can be avoided if empirical charges are used instead of QM charges. A few studies were published, which give QSPR models for predicting pK_a using topological empirical charges as descriptors (specifically PEOE charges) [22,42,43]. But these models provided relatively weak predictions.

The geometrical charges seem to be more promissing descriptors, because they are able to take into consideration the influence of the molecule's conformation on the atomic charges. The conformation of the atoms surrounding the dissociating hydrogens strongly influences the dissociation process, and also the atomic charges.

The EEM method is a geometrical empirical charge calculation approach which can be useful for pK_a prediction by QSPR. This approach calculates charges using the following equation system:

$$\begin{pmatrix} B_{1} & \frac{\kappa}{R_{1,2}} & \dots & \frac{\kappa}{R_{1,N}} & -1 \\ \frac{\kappa}{R_{2,1}} & B_{2} & \dots & \frac{\kappa}{R_{2,N}} & -1 \\ \vdots & \vdots & \ddots & \vdots & \vdots \\ \frac{\kappa}{R_{N,1}} & \frac{\kappa}{R_{N,2}} & \dots & B_{N} & -1 \\ 1 & 1 & \dots & 1 & 0 \end{pmatrix} \begin{pmatrix} q_{1} \\ q_{2} \\ \vdots \\ q_{N} \\ \overline{\chi} \end{pmatrix} = \begin{pmatrix} -A_{1} \\ -A_{2} \\ \vdots \\ -A_{N} \\ Q \end{pmatrix}$$
(1)

where q_i is the charge of atom i; $R_{i,j}$ is the distance between atoms i and j; Q is the total charge of the molecule; N is the number of atoms in the molecule; $\overline{\chi}$ is the molecular electronegativity, and A_i , B_i and κ are empirical parameters. These parameters are obtained by a parameterization process, which uses QM atomic charges to calculate a set of parameters for which EEM best reproduces these QM charges. EEM is very popular, and despite the fact that it was developed more than twenty years ago, new

parameterizations [39,44-50] and modifications [47,51,52] of EEM are still under development. Its accuracy is comparable to the QM charge calculation approach for which it was parameterized. Additionally, EEM is very fast, as its computational complexity is $\theta(N^3)$, where N is the number of atoms in the molecule.

Therefore, in the present study, we focus on pK_a prediction using QSPR models which employ EEM charges. Specifically, we created and evaluated QSPR models based on EEM charges computed using 18 EEM parameter sets. We also compared these QSPR models with corresponding QSPR models which employ QM charges computed by the same charge calculation schemes used for EEM parameterization.

Methods

EEM parameter sets

In our study, we used all EEM parameters published till now. Specifically, we found 18 different EEM parameters sets, published in 8 different articles [39,44-50]. The parameters cover two QM theory levels (HF and B3LYP), two basis sets (STO-3G and 6-31G*) and six population analyses (MPA, NPA, Hirshfeld, MK, CHELPG, AIM). Unfortunately, only some combinations of QM theory levels, basis sets and population analyses are available. On the other hand, more parameter sets were published for some combinations (i.e., 6 parameter sets for HF/STO-3G/MPA). All the parameter sets include parameters for C, O, N and H. Some sets include also parameters for S, P, halogens and metals. Most of the sets do not include parameters for C and N bonded by triple bond. Summary information about all these parameter sets is given in Table 1.

EEM charge calculation

The EEM charges were calculated by the program EEM SOLVER [53] using each of the 18 EEM parameter sets.

QM charge calculation

We calculated QM atomic charges for all the combinations of QM theory level, basis set and population analysis for which we have EEM parameters (see Table 1). Specifically, atomic charges were calculated for these eight QM approaches: HF/STO-3G/MPA, HF/6–31G*/MK, and B3LYP/6–31G* with MPA, NPA, Hirshfeld, MK, CHELPG and AIM). The QM charge calculations were carried out using Gaussian09 [54]. In the case of AIM population analysis, the output from Gaussian09 was further processed by the software package AIMAll [55].

Data set for phenols

There are two main ways to create a QSPR model for a feature to be predicted. The first is to create as general a model as possible, with the risk that the accuracy of such a model may not be high. The second approach is to develop more models, each of them being dedicated to a certain class of compounds. Here we took the second approach, following a similar methodology as in previous studies [21-24]. Specifically, we focus on substituted phenols, because they are the most common test set molecules employed in the evaluation of novel p K_a prediction approaches [21-24,56-58]. Our data set contains the 3D structures of 74 distinct phenol molecules. This data set is of high structural diversity and it covers molecules with p K_a values from 0.38 to 11.1. The molecules were obtained from the NCI Open Database Compounds [59] and their 3D structures were generated by CORINA 2.6 [60], without any further geometry optimization. Our data set is a subset of the phenol data set used in our previous work related to pK_a prediction from QM atomic charges [24]. The subset is made up of phenols which contain only C, O, N and H, and none of the molecules contain triple bonds. This limitation is necessary, because the EEM parameters of all 18 studied EEM parameter sets are available only for such molecules (see Table 1). For each phenol molecule from our data set, we also prepared the structure of the dissociated form, where the hydrogen is missing from the phenolic OH group. This dissociated molecule was created by removing the hydrogen from the original structure without subsequent geometry optimization. The list of the molecules, including their names, NCS numbers, CAS numbers and experimental pK_a values, can be found in the (Additional file 1: Table S1a). The SDF files with the 3D structures of molecules and their dissociated forms are also in the (Additional file 2: Molecules).

Data set for carboxylic acids

An aspect which is very important for the applicability of the pK_a prediction approach is its transferability to other chemical classes. In this work, we provide a case study showing the performance of the approach on carboxylic acids, which are also very common testing molecules for pK_a prediction approaches [19-21,43]. The data set contains 71 distinct molecules of carboxylic acids and their dissociated forms. The 3D structures of these molecules were obtained in the same way as for the phenols. The list of the molecules, including their names, NCS numbers, CAS numbers and experimental pK_a values can be found in the (Additional file 3: Table S1b). The SDF files with the 3D structures of the molecules and their dissociated forms are also included in the (Additional file 2: Molecules).

pK_a values

The experimental pK_a values were taken from the Physprop database [61].

QM theory level	PA	EEM parameter	Published by	Year of	Elements included
+ basis set		set name		publication	
HF/STO-3G	MPA	Svob2007_cbeg2	Svobodova et al. [44]	2007	C, O, N, H, S
		Svob2007_cmet2	Svobodova et al. [44]	2007	C, O, N, H, S, Fe, Zn
		Svob2007_chal2	Svobodova et al. [44]	2007	C, O, N, H, S, Br, Cl, F, I
		Svob2007_hm2	Svobodova et al. [44]	2007	C, O, N, H, S, F, Cl, Br, I, Fe, Zn
		Baek1991	Baekelandt et al. [45]	1991	C, O, N, H, P, Al, Si
		Mort1986	Mortier et al. [39]	1986	C, O, N, H
HF/6-31G*	MK	Jir2008_hf	Jirouskova et al. [46]	2008	C, O, N, H, S, F, Cl, Br, I, Zn
B3LYP/6-31G*	MPA	Chaves2006	Chaves et al. [47]	2006	C, O, N, H, F
		Bult2002_mul	Bultinck et al. [48]	2002	C, O, N, H, F
	NPA	Ouy2009	Ouyang et al. [49]	2009	C, O, N, H, F
		Ouy2009_elem	Ouyang et al. [49]	2009	C, O, N, H, F
		Ouy2009_elemF	Ouyang et al. [49]	2009	C, O, N, H, F
		Bult2002_npa	Bultinck et al. [48]	2002	C, O, N, H, F
	Hir.	Bult2002_hir	Bultinck et al. [48]	2002	C, O, N, H, F
	MK	Jir2008_mk	Jirouskova et al. [46]	2008	C, O, N, H, S, F, Cl, Br, I, Zn
		Bult2002_mk	Bultinck et al. [48]	2002	C, O, N, H, F
	CHELPG	Bult2002_che	Bultinck et al. [48]	2002	C, O, N, H, F
	AIM	Bult2004_aim	Bultinck et al. [50]	2004	C, O, N, H, F

Table 1 Summary information about the EEM parameter sets used in the present study

Descriptors and QSPR models for phenols

Our descriptors were atomic charges. We analyzed two types of QSPR models, namely QSPR models with three descriptors (3d QSPR models) and QSPR models with five descriptors (5d QSPR models).

The 3d QSPR models used those descriptors which proved to be the most relevant for pK_a prediction in our previous study [24]. Therefore these descriptors were the atomic charge of the hydrogen atom from the phenolic OH group (q_H) , the charge on the oxygen atom from the phenolic OH group (q_O) , and the charge on the carbon atom binding the phenolic OH group (q_{C1}) . These descriptors were used to establish the QSPR models by the general equation:

$$pK_a = p_H \cdot q_H + p_O \cdot q_O + p_{C1} \cdot q_{C1} + p \tag{2}$$

where p_H , p_O , p_{C1} and p are parameters of the QSPR model (i.e., constants derived by multiple linear regression). The 5d QSPR models employ the above mentioned descriptors q_H , q_O and q_{C1} and additionally also the charge on the phenoxide O^- from the dissociated molecule (q_{OD}) , and the charge on the carbon atom binding this oxygen (q_{C1D}) . Using the charges from the dissociated molecules for p_{K_a} prediction was inspired by the work of Dixon et al. [19]. The equation of the 5d QSPR models is therefore:

$$pK_a = p'_H \cdot q_H + p'_O \cdot q_O + p'_{C1} \cdot q_{C1} + p'_{OD} \cdot q_{OD} + p'_{C1D} \cdot q_{C1D} + p'$$

where p'_H , p'_O , p'_{C1} , p'_{OD} , p'_{C1D} and p' are parameters of the OSPR model.

Descriptors and QSPR models for carboxylic acids

The descriptors were again atomic charges and, similarly as for phenols, two types of QSPR models were developed and evaluated. Specifically, QSPR models with four descriptors (4d QSPR models) and QSPR models with seven descriptors (7d QSPR models). The 4d QSPR models used similar descriptors as the 3d models for phenols the atomic charge of the hydrogen atom from the COOH group (q_H), the charge on the hydrogen bound oxygen atom from the COOH group (q_O), and the charge on the carbon atom binding the COOH group (q_{C1}). Additionally, also the charge of the second carboxyl oxygen (q_{O2}) is included. These 4d QSPR models are represented by the equation:

$$pK_a = p_H \cdot q_H + p_O \cdot q_O + p_{O2} \cdot q_{O2} + p_{C1} \cdot q_{C1} + p \quad (4)$$

where p_H , p_O , p_{O2} , p_{C1} and p are parameters of the QSPR model. The 7d QSPR models employ also charges from the dissociated forms, namely the charge on the carboxyl oxygens (q_{OD} , q_{O2D}) and the charge on the carboxylic carbon atom (q_{C1D}). The equation of the 7d QSPR models is therefore:

$$pK_{a} = p'_{H} \cdot q_{H} + p'_{O} \cdot q_{O} + p'_{O2} \cdot q_{O2} + p'_{C1} \cdot q_{C1} + p'_{OD} \cdot q_{OD} + p'_{O2D} \cdot q_{O2D} + p'_{C1D} \cdot q_{C1D} + p'$$
(5)

where p'_H , p'_O , p'_{O2} , p'_{C1} , p'_{OD} , p'_{O2D} , p'_{C1D} and p' are parameters of the QSPR model.

QSPR model parameterization

The parameterization of the QSPR models was done by multiple linear regression (MLR) using the software tool QSPR Designer [62].

Results and discussion

QM and EEM QSPR models for phenols

We prepared one 3d QSPR model and one 5d QSPR model using atomic charges calculated by each of the above mentioned 18 EEM parameter sets. These models are denoted 3d or 5d EEM QSPR models. Additionally, we created one 3d and one 5d QSPR model using atomic charges calculated by each of the corresponding 8 QM charge calculation approaches (denoted as 3d or 5d QM QSPR models). The data set of 74 phenol molecules was used for the parameterization of the QSPR models, and the obtained models were validated for all molecules in the data set.

The parameterization of the 3d EEM QSPR models showed that several molecules in the data set perform as outliers. For this reason, we created also EEM QSPR models without outliers (i.e., EEM QSPR models for which parameterization was done using a data set that excluded the previously observed outliers). These models are denoted 3d EEM QSPR WO models. We classified as outliers 10% of the molecules from our data set, which had the highest Cook's square distance. Therefore the 3d EEM QSPR WO models were parameterized using 67 molecules, and their validation was also done on the data set excluding outliers.

The quality of the QSPR models, i.e. the correlation between experimental pK_a and the pK_a calculated by each model, was evaluated using the squared Pearson correlation coefficient (R^2) , root mean square error (RMSE), and average absolute pK_a error $(\overline{\Delta})$, while the statistical criteria were the standard deviation of the estimation (s) and Fisher's statistics of the regression (F).

Table 2 contains the quality criteria (R^2 , RMSE, $\overline{\Delta}$) and statistical criteria (s and F) for all the QSPR models analyzed. All these models are statistically significant at p=0.01. Since our data sets contained 74 and 67 molecules, respectively, the appropriate F value to consider was that for 60 samples. Thus, the 3d QSPR models are statistically significant (at p=0.01) when F>4.126 and the 5d QSPR models when F>3.339. Figure 1 summarizes the R^2 of all QSPR models for ease of visual comparison, and Tables 3 and 4 provide a comparison of the models from specific points of view. The parameters of the QSPR models are summarized in the (Additional file 4: Table S2) and all charge descriptors and pK_a values are contained in the (Additional file 5: Table S6). The most relevant graphs of

correlation between experimental and calculated pK_a are visualized in Figure 2.

Prediction of p K_a using EEM charges

The key question we wanted to answer in this paper is whether EEM QSPR models are applicable for pK_a prediction. For this purpose we selected a set of phenol molecules and generated QSPR models which used EEM atomic charges as descriptors. We then evaluated the accuracy of these models by comparing the predicted pK_a values with the experimental ones. The results (see Tables 2 and 3, Figure 1) clearly show that QSPR models based on EEM charges are indeed able to predict the pK_a of phenols with very good accuracy. Namely, 63% of the EEM QSPR models evaluated in this study were able to predict p K_a with $R^2 > 0.9$. The average R^2 for all 54 EEM QSPR models considered was 0.9, while the best EEM QSPR model reached $R^2 = 0.924$. Our findings thus suggest that EEM atomic charges may prove as efficient QSPR descriptors for pK_a prediction. The only drawback of EEM is that EEM parameters are currently not available for some types of atoms. Nevertheless, EEM parameterization is still a topic of research, therefore more general parameter sets are being developed.

Improvement of EEM QSPR models by removing outliers

The quality of 3d EEM QSPR models can be markedly increased by removing the outliers. In this case, the models have average $R^2 = 0.911$ and 83% of them have $R^2 > 0.9$. The disadvantage of these models is that they are not able to cover the complete data set (i.e., 10% of molecules must be excluded as outliers).

On the other hand, the outliers are similar for all EEM QSPR models. For example, while 16 molecules from our data set are outliers for at least one parameter set, 10 out of these 16 molecules are outliers for five or more parameter sets. From the chemical point of view, most of the outliers contain one or more nitro groups. This may be related to reported lower accuracy of EEM for these groups [48]. In general one limitation of the 3d EEM QSPR models is that they are very sensitive to the quality of EEM charges. Therefore, if the EEM charges are inaccurate for certain compounds or class of compounds, the 3d QSPR models based on these EEM charges will have lower performance for these compounds or class of compounds. In addition, a lower experimental accuracy of these p K_a values may also be a reason for low performance in some cases. A table containing information about outlier molecules is given in the (Additional file 6: Table S3).

Improvement of EEM QSPR models by adding descriptors

Our first EEM QSPR models contained three descriptors (3d), namely atomic charges originating from the non-dissociated molecule. Nonetheless, in our study we found

Table 2 Quality criteria and statistical criteria for all the QSPR models analyzed in the present study and focused on phenols

QM theory level	PA	EEM parameter	QSPR model	R^2	RMSE	$\overline{\Delta}$	S	F
+ basis set		set name						
HF/STO-3G	MPA	-	3d QM	0.9515	0.490	0.388	0.504	458
		-	5d QM	0.9657	0.412	0.310	0.430	358
		Svob2007_cbeg2	3d EEM	0.8671	0.812	0.571	0.835	152
		-	3d EEM WO	0.9239	0.482	0.382	0.497	255
			5d EEM	0.9179	0.638	0.481	0.666	152
		Svob2007_cmet2	3d EEM	0.8663	0.814	0.577	0.837	151
			3d EEM WO	0.9239	0.482	0.386	0.497	255
			5d EEM	0.9189	0.634	0.476	0.661	154
		Svob2007_chal2	3d EEM	0.8737	0.792	0.554	0.814	161
			3d EEM WO	0.9127	0.483	0.387	0.498	220
			5d EEM	0.9203	0.629	0.473	0.656	157
			3d EEM	0.8671	0.812	0.578	0.835	152
			3d EEM WO	0.9241	0.481	0.387	0.496	256
			5d EEM	0.9179	0.638	0.478	0.666	152
		 Baek1991	3d EEM	0.9099	0.669	0.531	0.688	236
			3d EEM WO	0.9166	0.531	0.423	0.548	231
			5d EEM	0.9195	0.632	0.493	0.659	155
		 Mort1986	3d EEM	0.8860	0.752	0.577	0.773	181
		orersee	3d EEM WO	0.9151	0.520	0.405	0.536	226
			5d EEM	0.9142	0.652	0.524	0.680	145
HF/6-31G*	MK		3d QM	0.8405	0.890	0.727	0.915	123
111/0 310	IVIIX		5d QM	0.8865	0.750	0.641	0.782	106
		 Jir2008_hf	3d EEM	0.8612	0.730	0.582	0.762	145
		3112000±111	3d EEM WO	0.9182	0.500	0.394	0.516	236
			5d EEM	0.9154	0.648	0.488	0.676	147
D2LVD/C 21C*	MDA							
B3LYP/6-31G*	MPA		3d QM	0.9671	0.404	0.317	0.415	686
			5d QM	0.9724	0.370	0.274	0.386	479
		Chaves 2006	3d EEM	0.891	0.735	0.570	0.756	191
			3d EEM WO	0.9198	0.505	0.398	0.521	241
		D. 1+2002	5d EEM	0.9192	0.633	0.489	0.660	155
		Bult2002_mul	3d EEM	0.8876	0.747	0.589	0.768	184
			3d EEM WO	0.9151	0.520	0.416	0.536	226
			5d EEM	0.9158	0.646	0.504	0.674	148
B3LYP/6-31G*	NPA	-	3d QM	0.9590	0.451	0.349	0.464	546
		-	5d QM	0.9680	0.399	0.295	0.416	411
		Ouy2009	3d EEM	0.8731	0.793	0.541	0.815	161
			3d EEM WO	0.9043	0.505	0.379	0.521	198
			5d EEM	0.9094	0.670	0.503	0.699	137
		Ouy2009_elem	3d EEM	0.8727	0.795	0.546	0.817	160
			3d EEM WO	0.9113	0.487	0.382	0.502	216
			5d EEM	0.9132	0.656	0.495	0.684	143
		Ouy2009_elemF	3d EEM	0.8848	0.756	0.519	0.777	179
			3d EEM WO	0.9012	0.512	0.386	0.528	192
			5d EEM	0.8866	0.750	0.520	0.782	106

Table 2 Quality criteria and statistical criteria for all the QSPR models analyzed in the present study and focused on phenols (continued)

	Bult2002_npa	3d EEM	0.9044	0.689	0.532	0.708	221
		3d EEM WO	0.9098	0.523	0.405	0.539	212
		5d EEM	0.9180	0.638	0.488	0.666	152
Hir.	-	3d QM	0.9042	0.689	0.503	0.708	220
	-	5d QM	0.9477	0.509	0.356	0.531	246
	Bult2002_hir	3d EEM	0.8415	0.887	0.636	0.912	124
		3d EEM WO	0.8838	0.579	0.414	0.597	160
		5d EEM	0.9050	0.687	0.522	0.717	130
MK	-	3d QM	0.8447	0.878	0.705	0.903	127
	-	5d QM	0.8960	0.718	0.594	0.749	117
	Jir2008_dft	3d EEM	0.8696	0.804	0.555	0.827	156
		3d EEM WO	0.9224	0.487	0.371	0.502	250
		5d EEM	0.9148	0.650	0.489	0.678	146
	Bult2002_mk	3d EEM	0.8639	0.822	0.610	0.845	148
		3d EEM WO	0.9053	0.519	0.384	0.535	201
		5d EEM	0.9131	0.657	0.508	0.685	143
Chel.		3d QM	0.8528	0.854	0.712	0.878	135
	-	5d QM	0.9087	0.673	0.552	0.702	135
	Bult2002_che	3d EEM	0.8695	0.805	0.597	0.828	155
		3d EEM WO	0.8863	0.588	0.436	0.606	164
		5d EEM	0.9057	0.684	0.540	0.714	131
AIM	-	3d QM	0.9609	0.440	0.332	0.452	573
	-	5d QM	0.9677	0.400	0.285	0.417	407
	Bult2004_aim	3d EEM	0.8646	0.819	0.619	0.842	149
		3d EEM WO	0.8972	0.590	0.438	0.608	183
		5d EEM	0.9017	0.698	0.571	0.728	125

that using two additional charge descriptors from the dissociated molecule can markedly improve the predictive power of the EEM QSPR models. Tables 2 and 3, Figure 1 show that these new 5d EEM QSPR models provide better pK_a prediction than their corresponding 3d EEM QSPR models. Specifically, adding the descriptors derived from the dissociated molecules increased the average R^2 value for the EEM QSPR models from 0.876 to 0.913.

Comparison of EEM QSPR models and QM QSPR models

Another important question is how accurate the EEM QSPR models are in comparison with QM QSPR models. Table 2 and Figure 1 show that QM QSPR models provide, in most cases, more precise predictions. This is confirmed also by the average R^2 values from Table 3. This is not surprising, since EEM is an empirical method which just mimics the QM approach for which it was parameterized. An interesting fact is that the differences in accuracy between QM QSPR models and EEM QSPR models are not substantial. For example, 5d EEM QSPR models have average $R^2 = 0.913$, while 5d QM QSPR models

have average $R^2 = 0.951$. We also note that adding more descriptors to a QM QSPR model brings less improvement than adding more descriptors to an EEM QSPR model.

Influence of theory level and basis set

EEM parameters are available only for a relatively small number of theory levels (HF and B3LYP) and basis sets (STO-3G and 6–31G*). Therefore we can not perform such a deep analysis of theory level and basis set influence on pK_a prediction capability from EEM atomic charges, as was done for QM QSPR models by Gross et al. [22] or Svobodova et al. [24]. We can only compare the models employing HF/STO-3G and B3LYP/6–31G* charges, as these are the only combinations for which EEM parameters are available for the same population analysis (MPA). Therefore we can study only the influence of the combination of theory level / basis set, and not the isolated influence of the theory level or basis set. Our analysis revealed that B3LYP/6–31G* charges provide slightly more accurate QM QSPR models than HF/STO-3G charges (see

QM theory leve	el PA	EEM parameter		R ² of	QSPR mode	el	
+ basis set		set name	3d EEM	3d EEM WO	D 5d EEM	3d QM	5d QM
HF/STO-3G	MPA	Svob2007_cbeg2	0.8671	0.9239	0.9179	0.9515	0.9657
		Svob2007_cmet2	0.8663	0.9239	0.9189		
		Svob2007_chal2	0.8737	0.9127	0.9203		
		Svob2007_hm2	0.8671	0.9241	0.9179		
	İ	Baek1991	0.9099	0.9166	0.9195		
		Mort1986	0.8860	0.9151	0.9142		
HF/6-31G*	MK	Jir2008_hf	0.8696	0.9182	0.9154	0.8405	0.8865
B3LY P/6-31G	MPA	Chaves 2006	0.8910	0.9198	0.9192	0.9671	0.9724
		Bult2002_mul	0.8876	0.9151	0.9158		
	NPA	Ouy2009	0.8731	0.9043	0.9094	0.9590	0.9680
		Ouy2009_elem	0.8727	0.9113	0.9132		
	İ	Ouy2009_elemF	0.8848	0.9012	0.8866		
		Bult2002_npa	0.9044	0.9098	0.9180		
	Hir.	Bult2002_hir	0.8415	0.8838	0.9050	0.9042	0.9477
	MK	Jir2008_mk	0.8696	0.9224	0.9148	0.8447	0.8960
		Bult2002_mk	0.8639	0.9053	0.9131		
	Chel.	Bult2002_che	0.8695	0.8863	0.9057	0.8528	0.9087
	AIM	Bult2004_aim	0.8646	0.8972	0.9017	0.9609	0.9677
							_
Legend		, ,	good		acceptable	weak	_
\mathbb{R}^2	0.95 – 0.9	97 0.92 – 0.95 0	.91 – 0.92	0.9 – 0.91	0.85 – 0.9	0.8 - 0.85)
\mathbf{r} e 1 \mathbb{R}^2 for the correlation bety	ween calcı	ulated and experir	nental p K_a				

Table 3 Average R^2 between experimental and predicted p K_a for all QSPR models of a certain type and percentages of QSPR models whose R^2 values are in a certain interval

'R model	3d EEM	3d EEM WO	5d EEM	3d QM	5d QM
erage R ²	0.876	0.911	0.913	0.929	0.951
$R^2 > 0.9$	11%	83%	94%	78%	83%
$0.9 \ge R^2 > 0.85$	83%	17%	6%	6%	17%
$0.85 \ge R^2 > 0.8$	6%	0%	0%	17%	0%
R model		EEM based models		QM base	d models
erage R ²		0.900		0.9	940
$R^2 > 0.9$		63%		8	1%
$0.9 \ge R^2 > 0.85$		35%		13	3%
$0.85 \ge R^2 > 0.8$		2%		6	%
	erage R^2	erage R^2 0.876 $ \begin{array}{cccc} R^2 > 0.9 & 11\% \\ \hline 0.9 \ge R^2 > 0.85 & 83\% \\ \hline 0.85 \ge R^2 > 0.8 & 6\% \\ \end{array} $ R model $ \begin{array}{cccc} R^2 > 0.9 & \\ \hline 0.9 \ge R^2 > 0.85 & \\ \end{array} $	R model EEM based models $R^2 > 0.9$ 63% $R^2 > 0.85$ 83% $R^2 > 0.85$ 83% $R^2 > 0.85$ 6% $R^2 > 0.85$ 6% $R^2 > 0.9$ 63% $R^2 > 0.85$ 35%	erage R^2 0.876 0.911 0.913 $ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	erage R^2 0.876 0.911 0.913 0.929 $R^2 > 0.9$ 11% 83% 94% 78% $0.9 \ge R^2 > 0.85$ 83% 17% 6% 6% $0.85 \ge R^2 > 0.8$ 6% 0% 0% 17% R model EEM based models QM base erage R^2 0.900 0.9 $R^2 > 0.9$ 63% 81 $0.9 \ge R^2 > 0.85$ 35% 13

Table 4 Average R^2 between experimental and predicted p K_a for all QSPR models using atomic charges calculated by a specific combination of theory level and basis set, or by a specific population analysis

QSPF	R model	3d EEM	3d EEM WO	5d EEM	3d QM	5d QM
Theory level	HF/STO-3G	0.878	0.919	0.918	0.952	0.966
and basis set *	B3LYP/6-31G*	0.889	0.917	0.918	0.967	0.972
Population	MPA	0.889	0.917	0.918	0.967	0.972
analysis **	NPA	0.884	0.907	0.907	0.959	0.968
	Hirshfeld	0.842	0.884	0.905	0.904	0.948
	MK	0.867	0.914	0.914	0.845	0.896
	CHELPG	0.870	0.886	0.906	0.853	0.909
	AIM	0.865	0.897	0.902	0.961	0.968

^{*}Only QSPR models employing MPA were included in this analysis.
**Only QSPR models using B3LYP/6–31G* were included in this analysis.

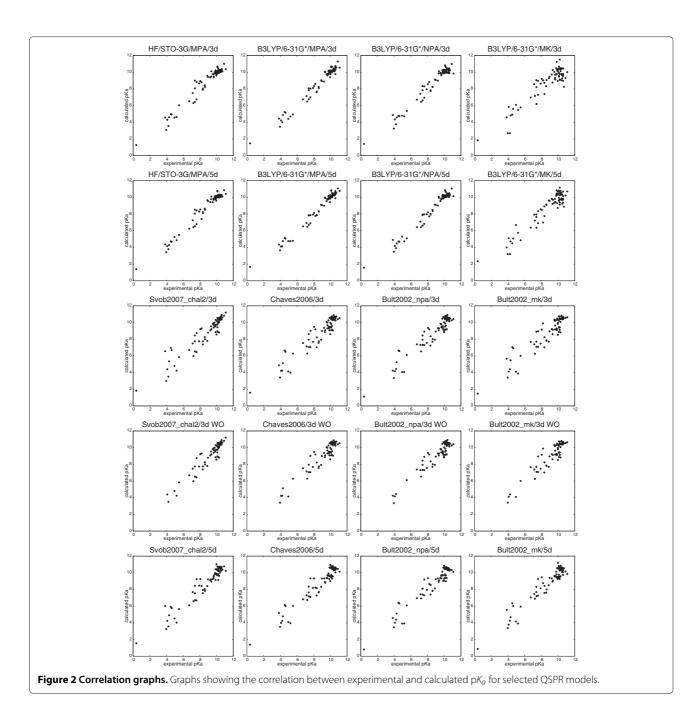


Table 4). This is in agreement with our previous findings [24], and it can be explained by the fact that $6-31G^*$ is a more robust basis set than STO-3G. However, the difference is not marked in the case of EEM QSPR models.

Influence of population analysis

Eleven EEM parameter sets were published for B3LYP/6–31G* with six different population analyses (see Table 1). Therefore it is straightforward to analyze the influence of the population analysis on the predictive power of the resulting QSPR models (see Table 4). We found that MPA

and NPA provide the best QM models, while MK and CHELPG (PAs based on fitting the atomic charges to the molecular electrostatic potential) provide weak QM models. Our results are in agreement with previous studies [22,24]. QM QSPR models based on AIM predict pK_a with accuracy comparable to MPA and NPA. In the case of EEM QSPR models, we did indeed find that MPA provided the best models, but most of the other population analyses gave comparable results. This confirms previous observations that the EEM approach is not able to faithfully mimic MK charges [63]. On the other hand,

Table 5 Comparison between the performance of the QSPR models developed here, and previously developed models

	Theory							Number of	
Method	level	PA	Basis set	Descriptors	R ²	s	F	molecules	Source
QM	B3LYP	NPA	6-311G**	90н	0.789	1.300	48	15	Kreye and Seybold [23] ^a
	B3LYP	NPA	6-311G**	90	0.731	1.500	38	15	Kreye and Seybold [23] ^a
	B3LYP	NPA	6-31+G*	9он	0.880	0.970	95	15	Kreye and Seybold [23] ^b
	B3LYP	NPA	6-31+G*	90	0.865	1.000	38	15	Kreye and Seybold [23] ^b
	B3LYP	NPA	6-311G(d,p)	90-	0.911	0.252	173	19	Gross and Seybold [22]
	B3LYP	NPA	6-311G(d,p)	9н	0.887	0.283	134	19	Gross and Seybold [22]
	B3LYP	NPA	6-31G*	9н, 90, 9с1	0.961	0.440	986	124	Svobodova and Geidl [24]
	B3LYP	NPA	6-311G	9 _H , 9 _O , 9 _{C1}	0.962	0.435	1013	124	Svobodova and Geidl [24]
	B3LYP	NPA	6-31G*	9н, 90, 9с1	0.959	0.464	545	74	This work
	B3LYP	NPA	6-31G*	9н, 90, 9с1,	0.968	0.410	705	74	This work
				90D, 9C1D					
EEM	B3LYP	NPA	6-31G*	9н, 90, 9с1,	0.918	0.656	261	74	This work ^c
				90D, 9C1D					
QM	B3LYP	MPA	6-311G(d,p)	9н	0.913	0.248	179	19	Gross and Seybold [22]
	B3LYP	MPA	6–311G(d,p)	90-	0.894	0.274	144	19	Gross and Seybold [22]
	B3LYP	MPA	6–311G	9н, 90, 9c1	0.938	0.556	605	124	Svobodova and Geidl [24]
	B3LYP	MPA	6-31G*	9н, 90, 9с1	0.959	0.450	936	124	Svobodova and Geidl [24]
	B3LYP	MPA	6-31G*	9 _H , 9 _O , 9 _{C1}	0.967	0.415	685	74	This work
	B3LYP	MPA	6-31G*	9н, 90, 9с1,	0.972	0.380	822	74	This work
				90D, 9C1D					
EEM	B3LYP	MPA	6-31G*	9н, 90, 9с1,	0.919	0.651	265	74	This work ^d
				9 _{OD} , 9 _{C1D}					
QM	B3LYP	MK	6-311G(d,p)	9 _H	0.344	0.682	9	19	Gross and Seybold [22]
	B3LYP	MK	6-311G(d,p)	90-	0.692	0.467	38	19	Gross and Seybold [22]
	B3LYP	MK	6–311G	9H, 9O, 9C1	0.822	0.941	185	124	Svobodova and Geidl [24]
	B3LYP	MK	6–31G*	9H, 90, 9C1	0.808	0.978	168	124	Svobodova and Geidl [24]
	B3LYP	MK	6-31G*	9H, 9O, 9C1	0.845	0.902	126	74	This work
	B3LYP	MK	6-31G*	9H, 90, 9C1	0.896	0.739	201	74	This work
				90D, 9C1D					
EEM	B3LYP	MK	6-31G*	9H, 90, 9C1	0.915	0.669	250	74	This work ^e
	DULII	77113	0 510	90D, 9C1D	0.213	0.009	250	, ,	THIS WOIN

^aWith solvent model SM5.4.

this drawback of EEM allowed the EEM QSPR models employing MK charges to predict pK_a more accurately than the corresponding QM QSPR models.

Influence of the EEM parameter set

Two or more EEM parameter sets are available in literature for four combinations of theory level, basis set and population analysis (see Table 1). We found that the quality of EEM QSPR models employing the same types of

charges slightly varies when using EEM parameters coming from different studies (see Table 2 and Figure 1). Even EEM parameters from the same study, but obtained by different approaches, lead to QSPR models of slightly different quality. In any case, these differences are minimal.

Comparison with previous work

QM QSPR models for pK_a prediction in phenols, similar to those presented in this paper (i.e., employing similar

^bWith solvent model SM8.

^cEEM parameter set Bult2002 npa.

dEEM parameter set Chaves 2006.

^eEEM parameter set Jir2008 mk.

Table 6 Comparison of the quality criteria and statistical criteria for the training set, test set and complete set for some selected charge calculation approaches

5d EEM QSPR model employing Svob2007_chal2 EEM parameters:

Complete set:

R^2	RMSE	S	F	Number of molecules
0.920	0.629	0.647	269	74

Cross-validation:

Cross-			Training set					Test set		
validation					Number of					Number of
step	R^2	RMSE	S	F	molecules	R^2	RMSE	S	F	molecules
1	0.9283	0.5211	0.5498	137	59	0.9202	1.0754	1.3884	21	15
2	0.9210	0.6538	0.6899	124	59	0.9029	0.5394	0.6963	17	15
3	0.9191	0.6442	0.6796	120	59	0.9275	0.5823	0.7517	23	15
4	0.9207	0.6244	0.6588	123	59	0.9271	0.6878	0.8880	23	15
5	0.9274	0.6302	0.6643	138	60	0.9008	0.6678	0.8834	15	14

5d EEM QSPR model employing Ouy2009_elemF EEM parameters:

Complete set:

R^2	RMSE	S	F	Number of molecules
0.8866	0.7501	0.7825	106	74

Cross-validation:

Cross-			Training set					Test set		
validation					Number of					Number of
step	R^2	RMSE	S	F	molecules	R^2	RMSE	S	F	molecules
1	0.8936	0.6349	0.6698	89	59	0.8704	1.2857	1.6598	12	15
2	0.8953	0.7526	0.7940	91	59	0.8018	0.7802	1.0072	7	15
3	0.8908	0.7481	0.7893	86	59	0.8647	0.7983	1.0306	12	15
4	0.8821	0.7614	0.8033	79	59	0.9154	0.7481	0.9658	19	15
5	0.8956	0.7557	0.7966	93	60	0.8089	0.8396	1.1107	7	14

charges) were previously published by Gross and Seybold [22], Kreye and Seybold [23] and Svobodova and Geidl [24]. Table 5 shows a comparison between these models and the models developed in this study. Our work is the first which presents QSPR models for pK_a prediction based on EEM charges. Therefore, we can not provide a comparison between EEM QSPR models, but we can compare against QSPR models based on QM charges only. It is seen therein that our 3d QM QSPR models show markedly higher R^2 and F values than the models published by Gross and Seybold and Kreye and Seybold (even if some of these models employ higher basis sets) and comparable R^2 and F values as models published by Svobodova and Geidl. Moreover, our 5d QM QSPR models outperform the models from Svobodova and Geidl. Our best EEM QSPR models (i.e., 5d EEM QSPR models) provide even better results than QM QSPR models from Gross and Seybold and Kreye and Seybold. These EEM QSPR models are not as accurate as the QM QSPR models published by Svobodova and Geidl or those developed

in this work, but the loss of accuracy is not too high (R^2 values are still > 0.91).

Cross-validation

Our results show that 5d EEM QSPR models provide a fast and accurate approach for pK_a prediction. Nonetheless, the robustness of these models should be proved. Therefore, all the 5d EEM QSPR models (i.e., 18 models) were tested by cross-validation. For comparison, also the cross-validation of all 5d QM QSPR models (i.e., 8 models) was done. The k-fold cross-validation procedure was used [64,65], where k = 5. Specifically, the set of phenol molecules was divided into five parts (each contained 20% of the molecules). The division was done randomly, and included stratification by pK_a value. Afterwards, five cross validation steps were performed. In the first step, the first part was selected as a test set, and the remaining four parts were taken together as the training set. The test and training sets for the other steps were prepared in a similar manner, by subsequently considering

Г	ON4 +b = =						D2 -4 OC	'DD dal
	QM theo	- 1	PA	١ ١	!	parameter	11	SPR model
	+ basi	s set			S	et name	7d EEM	7d QM
	HF/ST(D-3G	MP	Ά	Svob	2007_cbeg2	0.8831	0.9327
					Svob	2007_cmet2	0.8810	
					Svob	2007_chal2	0.8822	
					Svok	2007_hm2	0.8793	
				İ	В	aek1991	0.9211	
					М	ort1986	0.9176	
	B3LYP/6	6-31G*	MP	Ά	Ch	aves2006	0.9238	0.9059
					Bult2002_mul		0.9248	
			NP.	Ά	С	uy2009	0.8825	0.9169
					Ouy	2009_elem	0.8777	
				İ	Ouy2	009_elemF	0.8478	
					Bult2002_npa		0.9094	
	Legend	very god			ood	satisfactory	acceptable 0.85 – 0.9	
	R ²	0.92- 0.	95 0	0.91	- 0.92	- 0.92 0.9 - 0.91		0.8 – 0.85
ation hetwee	n calculated	d and expe	rimen	ntal r	K_{r} for	carboxylic acid	S	

Figure 3 Correlation between calculated and experimental p K_a for carboxylic acids.

one part as a test set, while the remaining parts served as a training set. For each step, the QSPR model was parameterized on the training set. Afterwards, the pK_a values of the respective test molecules were calculated via this model, and compared with experimental pK_a values. The results are summarized in the (Additional file 7: Table S4), while the cross-validation results for the best and the worst performing 5d EEM QSPR models are shown in Table 6. The cross-validation showed that the models are stable and the values of R^2 and RMSE are similar for the test set, the training set and the complete set. The robustness of EEM QSPR models and QM QSPR models is comparable.

Case study for carboxylic acids

We have shown that QSPR models based on EEM atomic charges can be used for predicting pK_a in phenols. In order to evaluate the general applicability of this approach for pK_a prediction, we tested the performance of such models for carboxylic acids. This case study is done for the charge schemes found to provide the best QM and EEM QSPR models in the case of phenols. Specifically, QM charges calculated by HF/STO-3G/MPA, B3LYP/6-31G*/MPA and B3LYP/6-31G*/NPA, and EEM charges calculated using the corresponding EEM parameters. Because 5d QSPR models provide the most accurate prediction for phenols, the case study is focused on their analogue for carboxylic acids, i.e., 7d QSPR models. Squared Pearson correlation coefficients of the analysed QSPR models are summarized in Figure 3, and all the quality and statistical criteria can be found in (Additional file 8:

Table S5). The results show that 7d EEM QSPR models are able to predict the pK_a of carboxylic acids with very good accuracy. Namely, 5 out of 12 analysed 7d EEM QSPR models were able to predict pK_a with $R^2 > 0.9$, while the best EEM QSPR model reached $R^2 = 0.925$. Therefore, we concluded that EEM QSPR models are indeed applicable also for carboxylic acids. Again QM QSPR models perform better than EEM QSPR models, but the differences are not substantial.

Conclusions

We found that the QSPR models employing EEM charges can be a suitable approach for pK_a prediction. From our 54 EEM QSPR models focused on phenols, 63% show a correlation of $R^2 > 0.9$ between the experimental and predicted p K_a . The most successful type of these EEM QSPR models employed 5 descriptors, namely the atomic charge of the hydrogen atom from the phenolic OH group, the charge on the oxygen atom from the phenolic OH group, the charge on the carbon atom binding the phenolic OH group, the charge on the oxygen from the phenoxide O⁻ from the dissociated molecule, and the charge on the carbon atom binding this oxygen. Specifically, 94% of these models have $R^2 > 0.9$, and the best one has $R^2 = 0.920$. In general, including charge descriptors from dissociated molecules, which was introduced in our work, always increases the quality of a QSPR model. The only drawback of EEM QSPR models is that the EEM parameters are currently not available for all types of atoms. Therefore the EEM parameter sets need to be expanded to larger sets of molecules and further improved.

As expected, the QM QSPR models provided more accurate pK_a predictions than the EEM QSPR models. Nevertheless, these differences are not substantial. Furthermore, a big advantage of EEM QSPR models is that one can calculate the EEM charges markedly faster than the QM charges. Moreover, the EEM QSPR models are not so strongly influenced by the charge calculation approach as the QM QSPR models are. Specifically, the QM QSPR models which use atomic charges obtained from calculations with higher basis set perform better, while the EEM QSPR models do not show such marked differences. Similarly, the quality of QM OSPR models depends a lot on population analysis, but EEM QSPR models are not influenced so much. Namely, QM QSPR models which use atomic charges calculated from MPA, NPA and Hirshfeld PA performed very well, while MK provides only weak models. In the case of EEM QSPR models, MPA performs also the best, but all other PAs (including MK) provide accurate results as well. The source of the EEM parameters also did not affect the quality of the EEM QSPR models significantly.

The robustness of EEM QSPR models was successfully confirmed by cross-validation. Specifically, the accuracy of pK_a prediction for the test, training and complete set were comparable. The applicability of EEM QSPR models for other chemical classes was tested in a case study focused on carboxylic acids. This case study showed that EEM QSPR models are indeed applicable for pK_a prediction also for carboxylic acids. Namely, 5 from 12 of these models were able to predict pK_a with $R^2 > 0.9$, while the best EEM QSPR model reached $R^2 = 0.925$.

Therefore, EEM QSPR models constitute a very promising approach for the prediction of pK_a . Their main advantages are that they are accurate, and can predict pK_a values very quickly, since the atomic charge descriptors used in the QSPR model can be obtained much faster by EEM than by QM. Additionally, the quality of EEM QSPR models is less dependent on the type of atomic charges used (theory level, basis set, population analysis) than in the case of QM QSPR models. Accordingly, EEM QSPR models constitute a pK_a prediction approach which is very suitable for virtual screening.

Additional files

Additional file 1: Table S1a. The list of the phenol molecules, including their names, NCS numbers, CAS numbers and experimental pK_a values.

Additional file 2: Molecules. The SDF files with the structures of the molecules and also their dissociated forms.

Additional file 3: Table S1b. The list of the carboxylic acid molecules, including their names, NCS numbers, CAS numbers and experimental pK_2 values.

Additional file 4: Table S2. The parameters of all the QSPR models for phenols.

Additional file 5: Table S6. The table containing charge descriptors for all charge calculation approaches and predicted pK_a values for all QSPR models (for phenols).

Additional file 6: Table S3. The information about outlier molecules for phenols.

Additional file 7: Table S4. The table of cross-validation results for phenols.

Additional file 8: Table S5. The quality and statistical criteria of QSPR models for carboxylic acids.

Abbreviations

3d: 3 descriptors; 4d: 4 descriptors; 5d: 5 descriptors; 7d: 7 descriptors; AlM: Atoms in Molecules; ANN: Artificial Neural Networks; B3LYP: Becke, three-parameter, Lee-Yang-Parr; DENR: Dynamic Electronegativity Relaxation; EEM: Electronegativity Equalization Method; GDAC: Geometry-Dependent Atomic Charge; HF: Hartree-Fock; KCM: Kirchhoff Charge Model; LFER: Linear Free Energy Relationships; MK: Merz-Singh-Kollman; MLR: Multiple Linear Regression; MP2: Møller-Plesset Perturbation Theory; MPA: Mulliken Population Analysis; NPA: Natural Population Analysis; PA: Population Analysis; PEOE: Partial Equalization of Orbital Electronegativity; QEq: Charge Equilibration; QM: Quantum Mechanical; QSPR: Quantitative Structure-Property Relationship; RMSE: Root Mean Square Error; SQE: Split Charge Equilibration; TSEF: Topologically Symmetric Energy Function; WO: Without Outliers.

Competing interests

The authors declare that they have no competing interests.

Author's contributions

The concept of the study originated from JK and was reviewed and extended by RA, while the design was put together by RSV and SG and reviewed by JK and RA. SG and CMI collected and prepared the input data. SG, OS, DS and TB performed the acquisition and post-processing of data. The data were analyzed and interpreted by RSV, SG, CMI and JK. The manuscript was written by RSV and SG in cooperation with JK and CMI, and reviewed by all authors.

Authors' information

Radka Svobodová Vařeková and Stanislav Geidl wish it to be known that, in their opinion, the first two authors should be regarded as joint First Authors.

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References

- Ishihama Y, Nakamura M, Miwa T, Kajima T, Asakawa N: A rapid method for pK_a determination of drugs using pressure-assisted capillary electrophoresis with photodiode array detection in drug discovery. J Pharm Sci 2002, 91(4):933–942.
- Babić S, Horvat A J, Pavlović D M, Kaštelan-Macan M: Determination of pK_a values of active pharmaceutical ingredients. *TrAC* 2007, 26(11):1043–1061.
- Manallack D: The pK_a distribution of drugs: application to drug discovery. Perspect Med Chem 2007, 1:25–38.
- Wan H, Ulander J: High-throughput pK_a screening and prediction amenable for ADME profiling. Expert Opin Drug Metabx Toxicol 2006, 2:139–155.
- Cruciani G, Milletti F, Storchi L, Sforna G, Goracci L: In silico pK_a
 prediction and ADME profiling. Chem Biodivers 2009, 6(11):
 1812–1821.
- Comer J, Tam K: Pharmacokinetic Optimization in Drug Research: Biological, Physicochemical, and Computational Strategies. Switzerland: Wiley-VCH. Verlag Helvetica Chimica Acta, Postfach, CH-8042 Zürich; 2001.
- Klebe G: Recent developments in structure-based drug design. J Mol Med 2000, 78:269–281.
- Lee AC, Crippen GM: Predicting pK_a. J Chem Inf Model 2009, 49:2013–2033.
- Rupp M, Körner R, Tetko IV: Predicting the pK_a of small molecules. Comb Chem High Throughput Screen 2010, 14(5):307–327.
- 10. Fraczkiewicz R: *In Silico Prediction of Ionization, Volume 5*. Oxford: Elsevier; 2006
- Ho J, Coote M: A universal approach for continuum solvent pK_a calculations: Are we there yet? *Theor Chim Acta* 2010, 125(1–2): 3–21
- 12. Clark J, Perrin DD: **Prediction of the strengths of organic bases.** *Q ReV Chem Soc* 1964, **18**:295–320.
- 13. Perrin DD, Dempsey B, Serjeant EP: pK_a Prediction for Organic Acids and Bases. New York: Chapman and Hall; 1981.
- 14. Blower PE, Cross KP: **Decision tree methods in pharmaceutical research.** *Curr Top Med Chem* 2006, **6:**31–39.
- Liptak MD, Gross KC, Seybold PG, Feldgus S, Shields G: Absolute pK_a determinations for substituted phenols. J Am Chem Soc 2002, 124:6421–6427.
- Toth AM, Liptak MD, Phillips DL, Shields GC: Accurate relative pK_α calculations for carboxylic acids using complete basis set and Gaussian-n models combined with continuum solvation methods.
 J Chem Phys 2001, 114:4595–4606.
- 17. Hagan MT, Demuth HB, Beale M: *In Neural, Network Design*. Boston: PWS, MA; 1996.
- Jelfs S, Ertl P, Selzer P: Estimation of pK_a for druglike compounds using semiempirical and information-based descriptors. J Chem Inf Model 2007, 47:450–459.
- 19. Dixon SL, Jurs PC: Estimation of p K_a for organic oxyacids using calculated atomic charges. *J Comput Chem* 1993, **14**:1460–1467.
- Zhang J, Kleinöder T, Gasteiger J: Prediction of pK_a values for aliphatic carboxylic acids and alcohols with empirical atomic charge descriptors. J Chem Inf Model 2006, 46:2256–2266.
- 21. Citra MJ: Estimating the pK_a of phenols, carboxylic acids and alcohols from semi-empirical quantum chemical methods. *Chemosphere* 1999, **1:**191–206.
- Gross KC, Seybold PG, Hadad CM: Comparison of different atomic charge schemes for predicting pK_a variations in substituted anilines and phenols. Int J Quantum Chem 2002, 90:445–458.
- Kreye WC, Seybold PG: Correlations between quantum chemical indices and the pK_as of a diverse set of organic phenols. Int J Quantum Chem 2009, 109:3679–3684.
- Svobodová Vařeková R, Geidl S, Ionescu CM, Skřehota O, Kudera M, Sehnal D, Bouchal T, Abagyan R, Huber HJ, Koča J: Predicting pK_α values of substituted phenols from atomic charges: Comparison of different quantum mechanical methods and charge distribution schemes. J Chem Inf Model 2011, 51(8):1795–1806.

- Mulliken RS: Electronic structures of molecules XI. Electroaffinity, molecular orbitals and dipole moments. J Chem Phys 1935, 3(9):573–585.
- Mulliken RS: Criteria for construction of good self-consistent-field molecular orbital wave functions, and significance of LCAO-MO population analysis. J Chem Phys 1962, 36(12):3428–3439.
- Lowdin PO: On the non-orthogonality problem connected with the use of atomic wave functions in the theory of molecules and crystals. J Chem Phys 1950, 18(3):365–375.
- 28. Reed AE, Weinstock RB, Weinhold F: **Natural-population analysis.** *J Chem Phys* 1985, **83**(2):735–746.
- Bader RFW, Larouche A, Gatti C, Carroll MT, Macdougall PJ, Wiberg KB: Properties of atoms in molecules - dipole-moments and transferability of properties. J Chem Phys 1987, 87(2):1142–1152.
- Hirshfeld FL: Bonded-atom fragments for describing molecular charge-densities. Theor Chim Acta 1977, 44(2):129–138.
- Breneman CM, Wiberg KB: Determining atom-centered monopoles from molecular electrostatic potentials - the need for high sampling density in formamide conformational-analysis. J Comput Chem 1990, 11(3):361–373.
- 32. Besler BH, Merz KM, Kollman PA: **Atomic charges derived from** semiempirical methods. *J Comput Chem* 1990, **11**(4):431–439.
- Kelly CP, Cramer CJ, Truhlar DG: Accurate partial atomic charges for high-energy molecules using class IV charge models with the MIDI! basis set. Theor Chem Acc 2005, 113(3):133–151.
- Abraham RJ, Griffiths L, Loftus P: Approaches to charge calculations in molecular mechanics. J Comput Chem 1982, 3(3):407–416.
- 35. Gasteiger J, Marsili M: **Iterative partial equalization of orbital electronegativity a rapid access to atomic charges.** *Tetrahedron* 1980, **36**(22):3219–3228.
- Cho KH, Kang YK, No KT, Scheraga HA: A fast method for calculating geometry-dependent net atomic charges for polypeptides. J Phys Chem B 2001, 105(17):3624–3634.
- Oliferenko AA, Pisarev SA, Palyulin VA, Zefirov NS: Atomic charges via electronegativity equalization: Generalizations and perspectives. Adv Quantum Chem 2006, 51:139–156.
- Shulga DA, Oliferenko AA, Pisarev SA, Palyulin VA, Zefirov NS:
 Parameterization of empirical schemes of partial atomic charge calculation for reproducing the molecular electrostatic potential. Dokl Chem 2008, 419:57–61.
- Mortier WJ, Ghosh SK, Shankar S: Electronegativity equalization method for the calculation of atomic charges in molecules. JAm Chem Soc 1986, 108:4315–4320.
- Rappe AK, Goddard WA: Charge equilibration for molecular-dynamics simulations. J Phys Chem 1991, 95(8):3358–3363.
- 41. Nistor RA, Polihronov JG, Muser MH, Mosey NJ: A generalization of the charge equilibration method for nonmetallic materials. *J Chem Phys* 2006, **125**(9):094108–094118.
- Czodrowski P, Dramburg I, Sotriffer CA, Klebe G: Development, validation, and application of adapted PEOE charges to estimate pK_a values of functional groups in protein-ligand complexes. Proteins Struct Funct Bioinf 2006, 65:424–437.
- Gieleciak R, Polanski J: Modeling robust QSAR. 2. Iterative variable elimination schemes for CoMSA: Application for modeling benzoic acid pK_a values. J Chem Inf Model 2007, 47:547–556.
- Svobodová Vařeková R, Jiroušková Z, Vaněk J, Suchomel S, Koča J: Electronegativity equalization method: Parameterization and validation for large sets of organic, organohalogene and organometal molecule. Int J Mol Sci 2007, 8:572–582.
- Baekelandt BG, Mortier WJ, Lievens JL, Schoonheydt RA: Probing the reactivity of different sites within a molecule or solid by direct computation of molecular sensitivities via an extension of the electronegativity equalization method. J Am Chem Soc 1991, 113(18):6730–6734.
- Jiroušková Z, Svobodová Vařeková R, Vaněk J, Koča J: Electronegativity equalization method: Parameterization and validation for organic molecules using the Merz–Kollman–Singh charge distribution scheme. J Comput Chem 2009, 30:1174–1178.
- Chaves J, Barroso JM, Bultinck P, Carbo-Dorca R: Toward an alternative hardness kernel matrix structure in the Electronegativity Equalization Method (EEM). J Chem Inf Model 2006, 46(4):1657–1665.

- Bultinck P, Langenaeker W, Lahorte P, De Proft, F, Geerlings P, Waroquier M, Tollenaere J: The electronegativity equalization method I:
 Parametrization and validation for atomic charge calculations.
 J Phys Chem A 2002, 106(34):7887–7894.
- Ouyang Y, Ye F, Liang Y: A modified electronegativity equalization method for fast and accurate calculation of atomic charges in large biological molecules. *Phys Chem* 2009, 11:6082–6089.
- Bultinck P, Vanholme R, Popelier PLA, De Proft, F, Geerlings P: High-speed calculation of AIM charges through the electronegativity equalization method. J Phys Chem A 2004, 108(46):10359–10366.
- Yang ZZ, Wang CS: Atom-bond electronegativity equalization method. 1. Calculation of the charge distribution in large molecules. J Phys Chem A 1997, 101:6315–6321.
- Menegon G, Loos M, Chaimovich H: Parameterization of the electronegativity equalization method based on the charge model 1. J Phys Chem A 2002, 106:9078–9084.
- Svobodová Vařeková R, Koča J: Optimized and parallelized implementation of the electronegativity equalization method and the atom-bond electronegativity equalization method. J Comput Chem 2006, 3:396–405.
- 54. Frisch MJ, Trucks GW, Schlegel HB, Scuseria GE, Robb MA, Cheeseman JR, Montgomery JAJr, Vreven T, Kudin KN, Burant JC, Millam JM, Iyengar SS, Tomasi J, Barone V, Mennucci B, Cossi M, Scalmani G, Rega N, Petersson GA, Nakatsuji H, Hada M, Ehara M, Toyota K, Fukuda R, Hasegawa J, Ishida M, Nakajima T, Honda Y, Kitao O, Nakai H, et al.: Gaussian 09, Revision E.01. Wallingford: Gaussian, Inc.; 2004.
- Keith TA: AIMAII, Version 11.12.19. USA: TK Gristmill Software, Overland Park KS; 2011. [aim.tkgristmill.com].
- Habibi-Yangjeh A, Danandeh-Jenagharad M, Nooshyar M: Application of artificial neural networks for predicting the aqueous acidity of various phenols using QSAR. J Mol Model 2006, 12:338–347.
- Hanai T, Koizumi K, Kinoshita T, Arora R, Ahmed F: Prediction of pK_a values of phenolic and nitrogen-containing compounds by computational chemical analysis compared to those measured by liquid chromatography. *J Chromatogr A* 1997, 762:55–61.
- 58. Tehan BG, Lloyd EJ, Wong MG, Pitt WR, Montana JG, Manallack DT, Gancia E: Estimation of p K_{α} Using semiempirical molecular orbital methods. Part 1: Application to phenols and carboxylic acids. Quant Struct-Act Relat 2002, **21:**457–472.
- NCI Open Database Compounds. Retrieved from [http://cactus.nci.nih. gov/] on August 10, 2010.
- Sadowski J, Gasteiger J: From atoms and bonds to three-dimensional atomic coordinates: Automatic model builders. Chem ReV 1993, 93:2567–2581
- Howard P, Meylan W: Physical/Chemical Property Database (PHYSPROP).
 North Syracuse NY: Syracuse Research Corporation, Environmental Science Center: 1999.
- Skřehota O, Svobodová Vařeková R, Geidl S, Kudera M, Sehnal D, Ionescu CM, Koča J: QSPR designer – a program to design and evaluate QSPR models. Case study on pK_a prediction. J Cheminf 2011, 3(Suppl 1):P16.
- Bultinck P, Langenaeker W, Lahorte P, De Proft, F, Geerlings P, Van Alsenoy, C, Tollenaere JP: The electronegativity equalization method II: Applicability of different atomic charge schemes. J Phys Chem A 2002, 106(34):7895–7901.
- Lemm S, Blankertz B, Dickhaus T, Müller KR: Introduction to machine learning for brain imaging. NeuroImage 2011, 56(2):387–399.
- Organisation for Economic Co-operation and Development: Guidance Document on the Validation of (Quantitative)Structure-Activity Relationships [(Q)SAR] Models. Paris: OECD; 2007. [http://search.oecd.org/ officialdocuments/displaydocumentpdf/?cote=env/jm/ mono(2007)2&doclanguage=en] (accessed April 6,2013).

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