

Supporting Information

(Submitted to the RSC Advances)

Role of Exchange and Correlation in the Real External Prediction of Mutagenicity: Performance of Hybrid and Meta-hybrid Exchange-correlation Functionals

Reenu and Vikas*

*Quantum Chemistry Group, Department of Chemistry & Centre of Advanced Studies in
Chemistry, Panjab University, Chandigarh-160014, India*

***Corresponding Author**

Email: qlabspu@pu.ac.in, qlabspu@yahoo.com

Phone: +91-172-2534408

Table of Contents

Supporting information Figures S1-S3

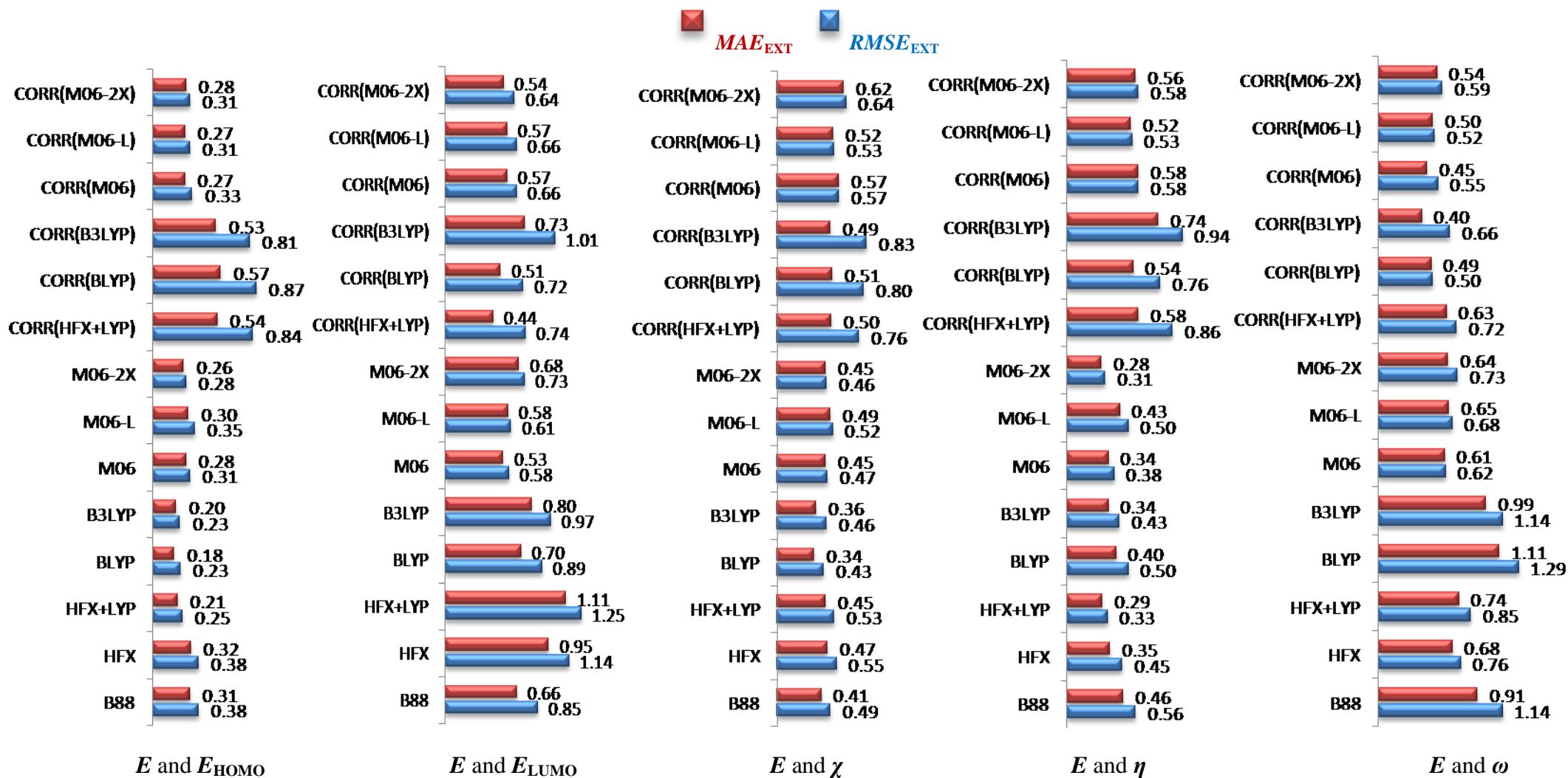
- Figure S1.** Comparison of the mean absolute error (*MAE*) and the root mean square error (*RMSE*) in the external (EXT) predictivity of various models, for the TA98 mutagenicity of nitrated-PAHs, based on the total energy (*E*), energy of the HOMO and the LUMO, absolute electronegativity (χ), chemical hardness (η) and electrophilicity index (ω) computed through exchange-only methods (HFX, B88), exchange-correlation (XC) methods (HFX+LYP, BLYP, B3LYP, M06, M06-L, M06-2X), and also based on the descriptors incorporating mainly the effect of electron correlation (CORR) from the respective XC methods.
- Figure S2(A).** Williams plots of the Standardized residuals vs Leverage (*h*) for TA100 mutagenicity for best models obtained with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively. The encircled values represent ID number of the compounds (refer to Supporting Information Table S1). The vertical (solid) line indicates warning leverage h^* , whereas the horizontal (dashed) line specifies standardized residual value of 3.0.
- Figure S2(B).** Williams plots of the Standardized residuals vs Leverage (*h*) for TA98 mutagenicity for best models developed with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively. The encircled values represent ID number of the compounds (refer to Supporting Information Table S1). The vertical (solid) line indicates warning leverage h^* , whereas the horizontal (dashed) line specifies standardized residual value of 3.0.
- Figure S3(A).** Scatter plots of the Experimental vs Predicted TA100 Mutagenicity for the best models developed with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively.
- Figure S3(B).** Scatter plots of the Experimental vs Predicted TA100 Mutagenicity for best models developed with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively.

Supporting information Tables S1 – S30

- Table S1.** List of nitrated-PAHs with their mutagenicity expressed as Log TA100 and Log TA98.
- Table S2a.** Computed value of total energy (*E*) descriptor, (all in a.u., 1a.u.=27.21165eV, 627.50956 kcal/mol) calculated with HFX, B88, HF, HFX+LYP, BLYP, B3LYP, M06, M06-L, and M06-2X level of the theory employing 6-311G(d,p) basis set for 51-nitrated PAHs.
- Table S2b.** Same as **Table S2a** but for HOMO energy descriptor (E_{HOMO}).
- Table S2c.** Same as **Table S2a** but for LUMO energy descriptor (E_{LUMO}).
- Table S2d.** Same as **Table S2a** but for absolute electronegativity descriptor (χ).
- Table S2e.** Same as **Table S2a** but for chemical hardness descriptor (η).

Table S2f.	Same as Table S2a but for electrophilicity index descriptor (ω).
Table S3.	Compounds in the training set (T), prediction set (P) and excluded set (E) in different types of splitting methods employed for TA100 mutagenicity. Excluded compounds are either response or structural outliers (determined from Williams plot).
Table S4.	Compounds in the training set (T), prediction set (P) and excluded set (E) in different types of splitting methods employed for TA98 mutagenicity. Excluded compounds are either response or structural outliers (determined from Williams plot).
Table S5.	Comparison of the key internal and external validation parameters for the models based on the total energy (E) and energy of HOMO (E_{HOMO}) computed with the Exchange (X) only, Exchange + Correlation (X+C) methods, and with the effect of electron-correlation (CORR) of the descriptors, for modeling TA98 mutagenicity of nitrated-PAHs.
Table S6.	Same as Table S5 but for the models based on the total electronic energy (E) and energy of the LUMO (E_{LUMO}).
Table S7.	Same as Table S5 but for the models based on the total electronic energy (E) and absolute electronegativity (χ).
Table S8.	Same as Table S5 but for the models based on the total energy (E) and chemical hardness (η).
Table S9.	Same as Table S5 but for the models based on the total electronic energy (E) and electrophilicity index (ω).
Table S10.	Comparison of the key internal and external validation parameters for the models developed with the total energy (E), energy of HOMO (E_{HOMO}), energy of LUMO (E_{LUMO}), absolute electronegativity (χ), chemical hardness (η) and electrophilicity index (ω) computed with the Exchange (X) only Hartree-Fock (HF) method, for modeling TA100 mutagenicity and TA98 mutagenicity of nitrated-PAHs.
Table S11.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange (X) only (B88, HFX and HF) based methods through activity sampling (ordered response) splitting method.
Table S12.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange (X) only (B88, HFX and HF) based methods through 30% (random) splitting method.
Table S13.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange (X) only (B88, HFX and HF) based methods through activity sampling (ordered response) splitting method.
Table S14.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange (X) only (B88, HFX and HF) based methods through 30% (random) splitting method.
Table S15.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange-correlation (HFX+LYP, BLYP and B3LYP) based methods through activity sampling (ordered response) splitting method.
Table S16.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange-correlation (HFX+LYP, BLYP and B3LYP) based methods through 30% (random) splitting method.
Table S17.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through activity sampling (ordered response) splitting method.
Table S18.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through 30% (random) splitting method.

Table S19.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange-correlation (HFX+LYP, BLYP and B3LYP) based methods through activity sampling (ordered response) splitting method.
Table S20.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange-correlation (HFX+LYP, BLYP and B3LYP) based methods through 30% (random) splitting method.
Table S21.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through activity sampling (ordered response) splitting method.
Table S22.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through 30% (random) splitting method.
Table S23.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through activity sampling (ordered response) splitting method.
Table S24.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through 30% (random) splitting method.
Table S25.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through activity sampling (ordered response) splitting method.
Table S26.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through 30% (random) splitting method.
Table S27.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through activity sampling (ordered response) splitting method.
Table S28.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(M06) CORR(M06-L), CORR(M06-2X)) based methods through 30% (random) splitting method.
Table S29.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through activity sampling (ordered response) splitting method.
Table S30.	Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through activity sampling 30% (random) splitting method.



QSAR models for TA98 mutagenicity

Figure S1. Comparison of the mean absolute error (MAE) and the root mean square error ($RMSE$) in the external (EXT) predictivity of various models, for the TA98 mutagenicity of nitrated-PAHs, based on the total energy (E), energy of the HOMO and the LUMO, absolute electronegativity (χ), chemical hardness (η) and electrophilicity index (ω) computed through exchange-only methods (HFX, B88), exchange-correlation (XC) methods (HFX+LYP, BLYP, B3LYP, M06, M06-L, M06-2X), and also based on the descriptors incorporating mainly the effect of electron correlation (CORR) from the respective XC methods.

Figure S2(A). Williams plots of the Standardized residuals vs Leverage (h) for TA100 mutagenicity for best models obtained with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively. The encircled values represent ID number of the compounds (refer to Supporting Information Table S1). The vertical (solid) line indicates warning leverage h^* , whereas the horizontal (dashed) line specifies standardized residual value of 3.0.

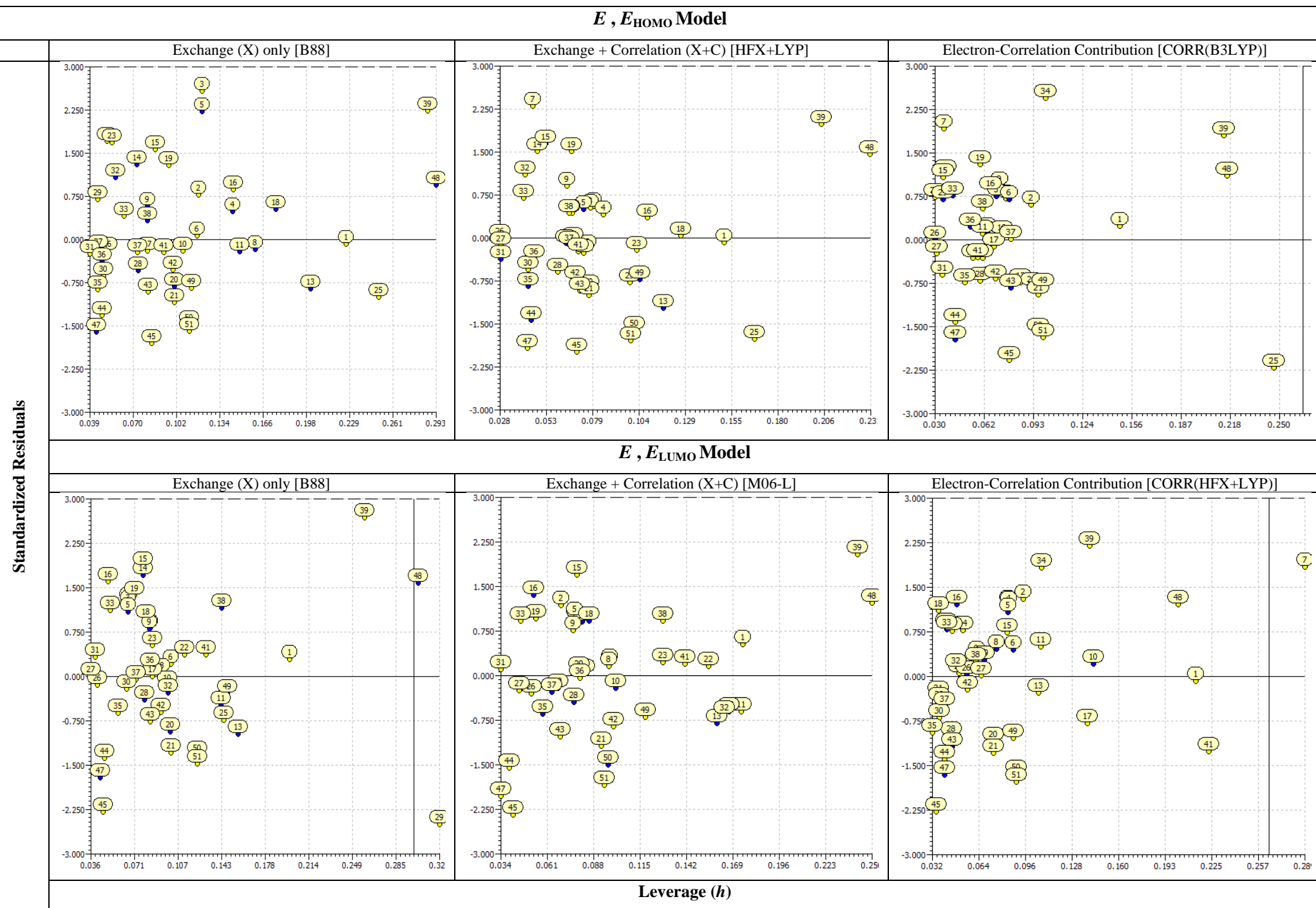


Figure S2(A) continued...

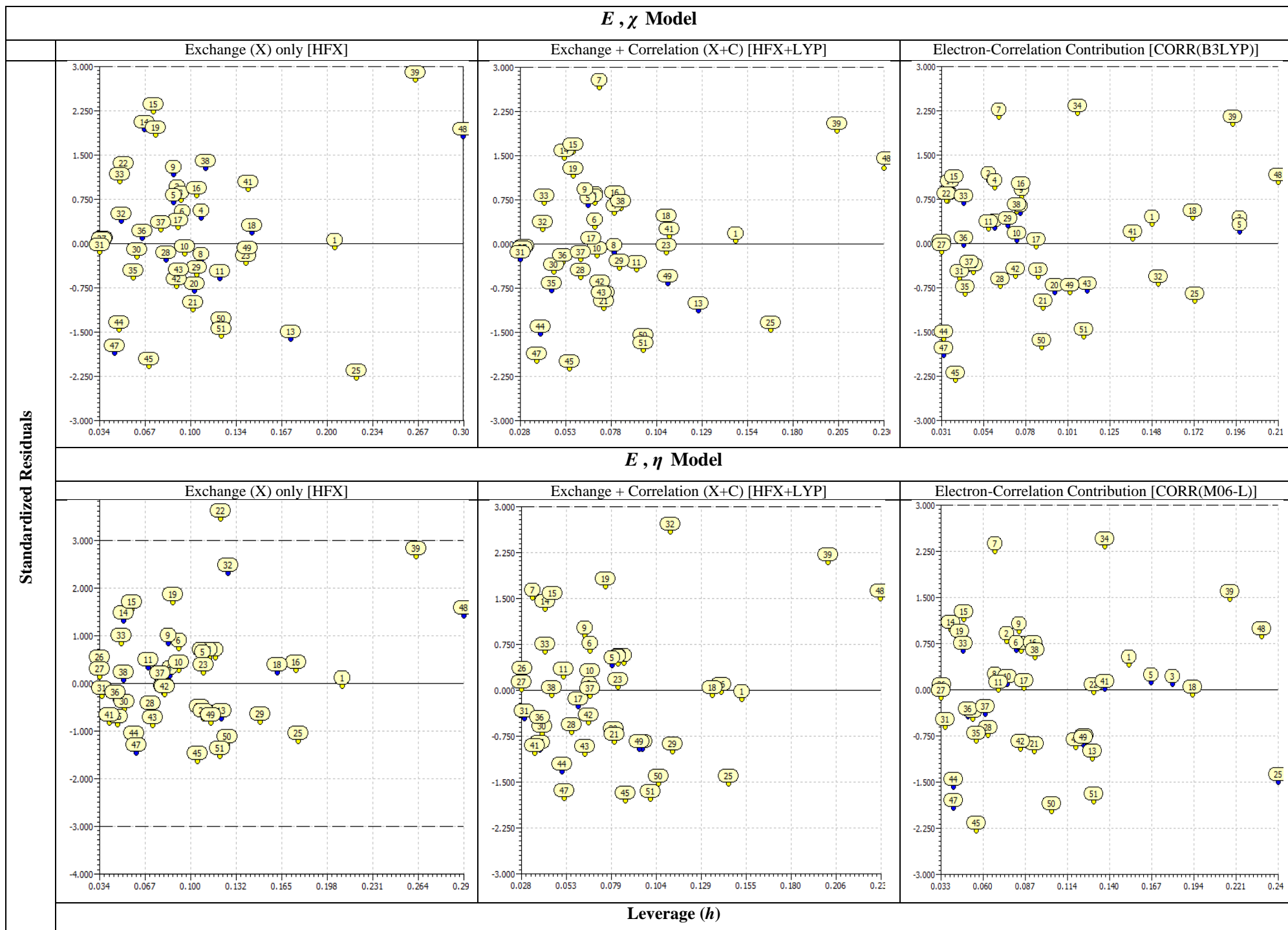


Figure S2(A) continued...

Figure S2(A) continued...

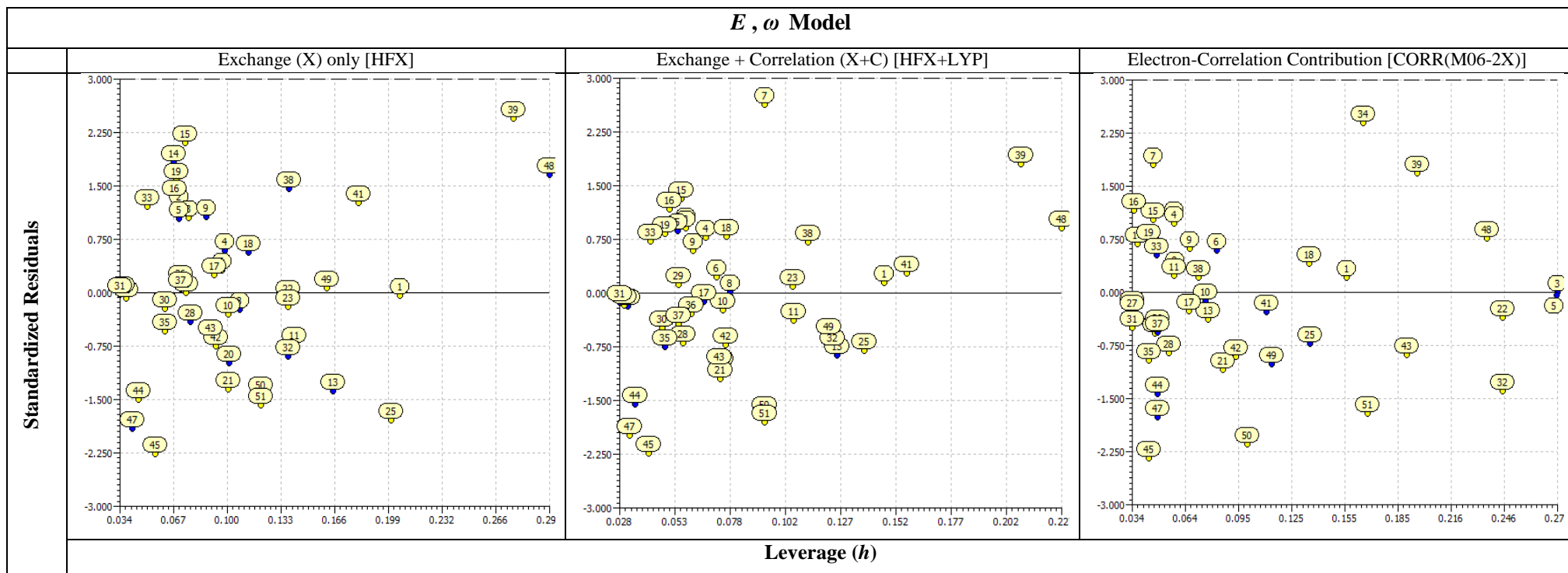


Figure S2(B). Williams plots of the Standardized residuals vs Leverage (h) for TA98 mutagenicity for best models developed with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively. The encircled values represent ID number of the compounds (refer to Supporting Information Table S1). The vertical (solid) line indicates warning leverage h^* , whereas the horizontal (dashed) line specifies standardized residual value of 3.0.

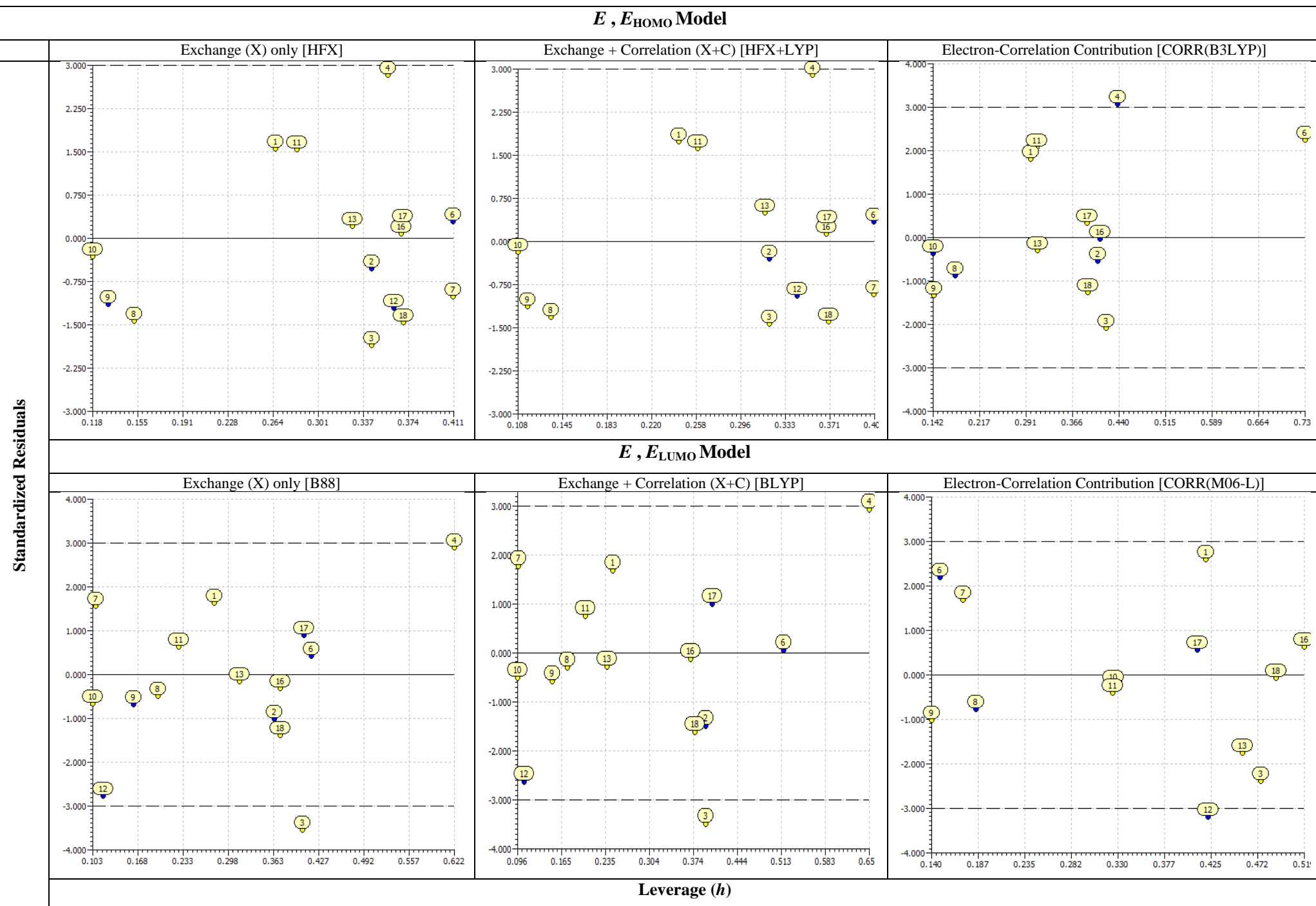


Figure S2(B) continued...

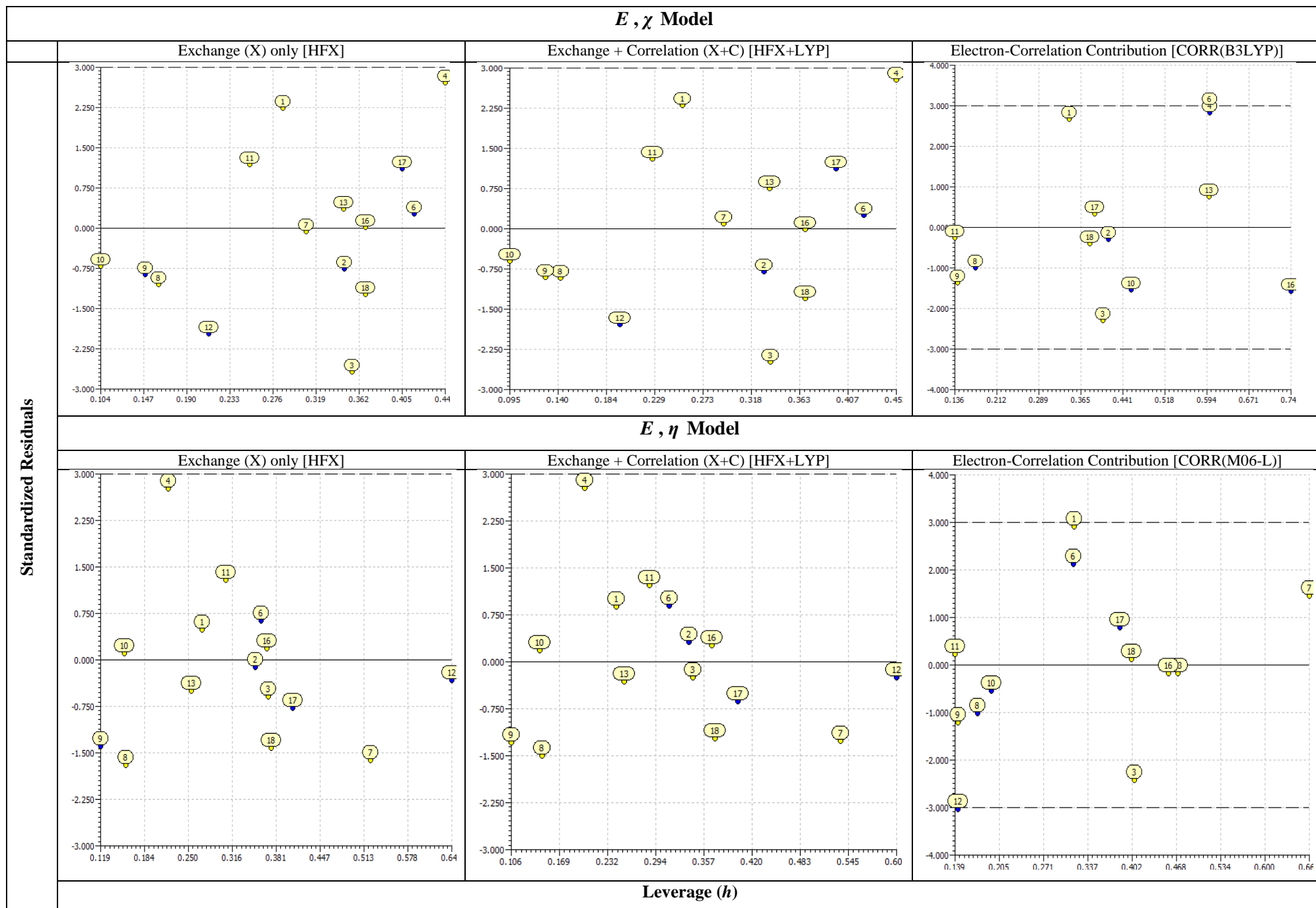


Figure S2(B) continued...

Figure S2(B) continued...

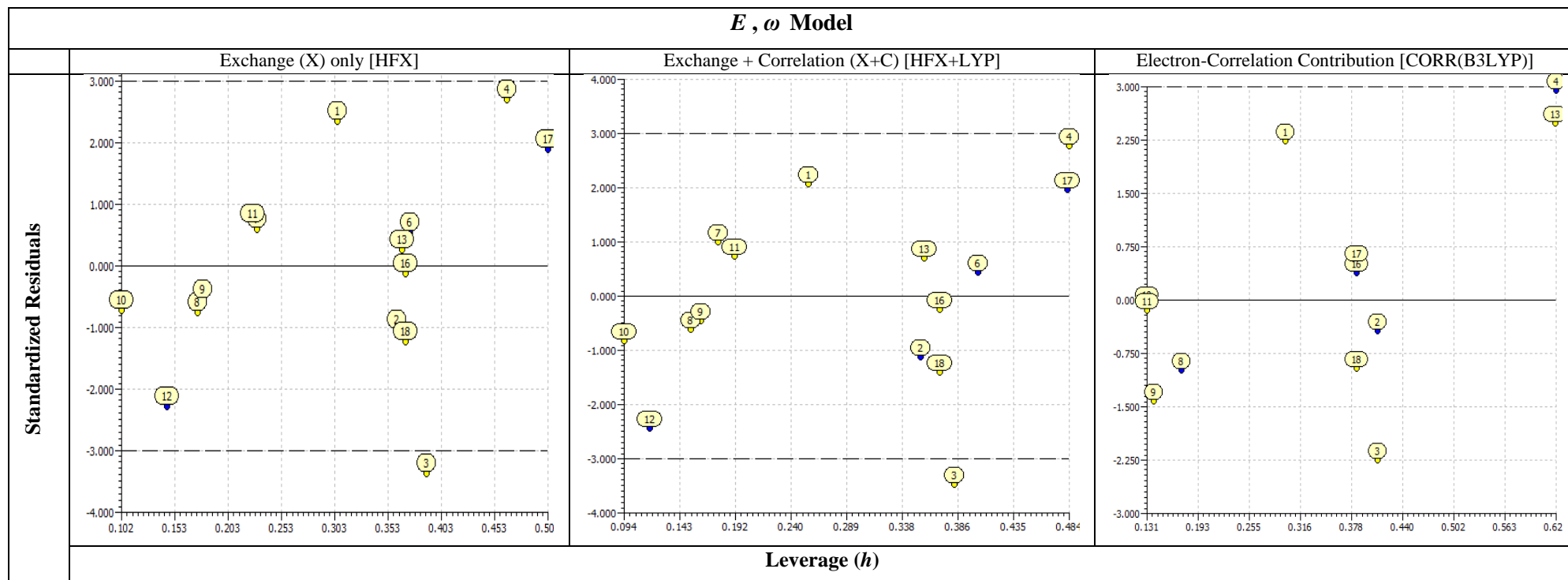


Figure S3. (A) Scatter plots of the Experimental vs Predicted TA100 Mutagenicity for the best models developed with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively.

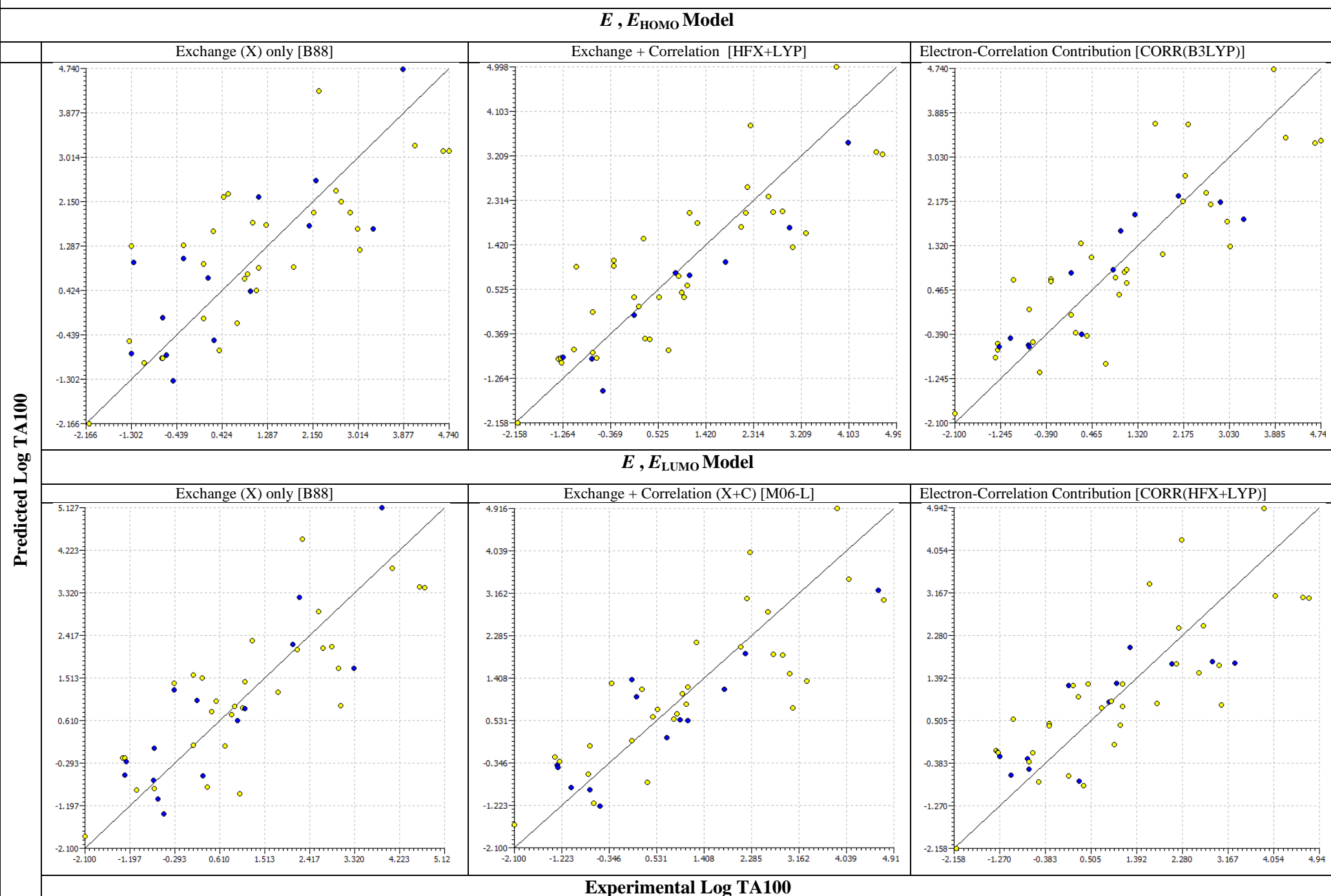


Figure S3(A) continued...

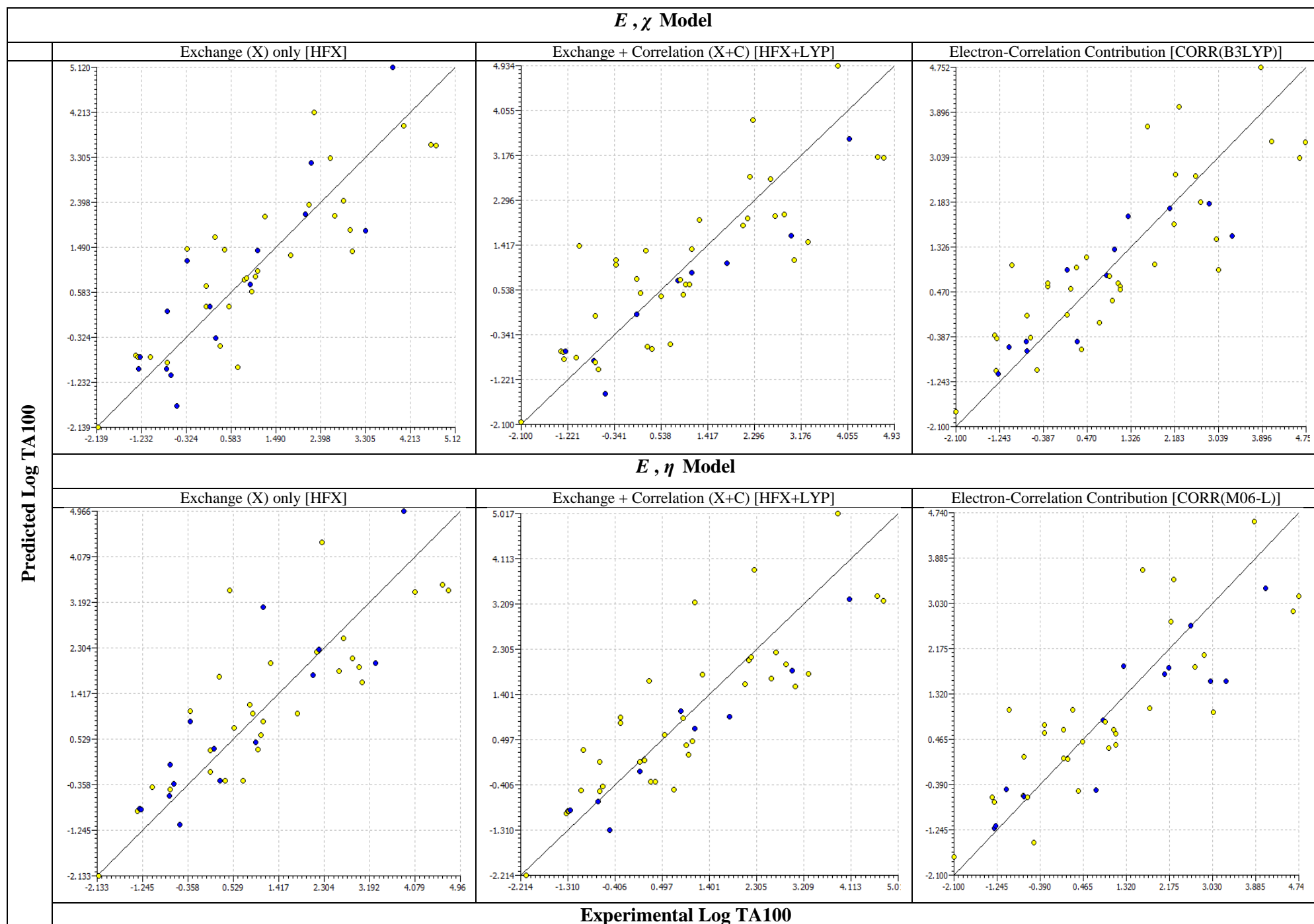


Figure S3(A) continued...

Figure S3(A) continued...

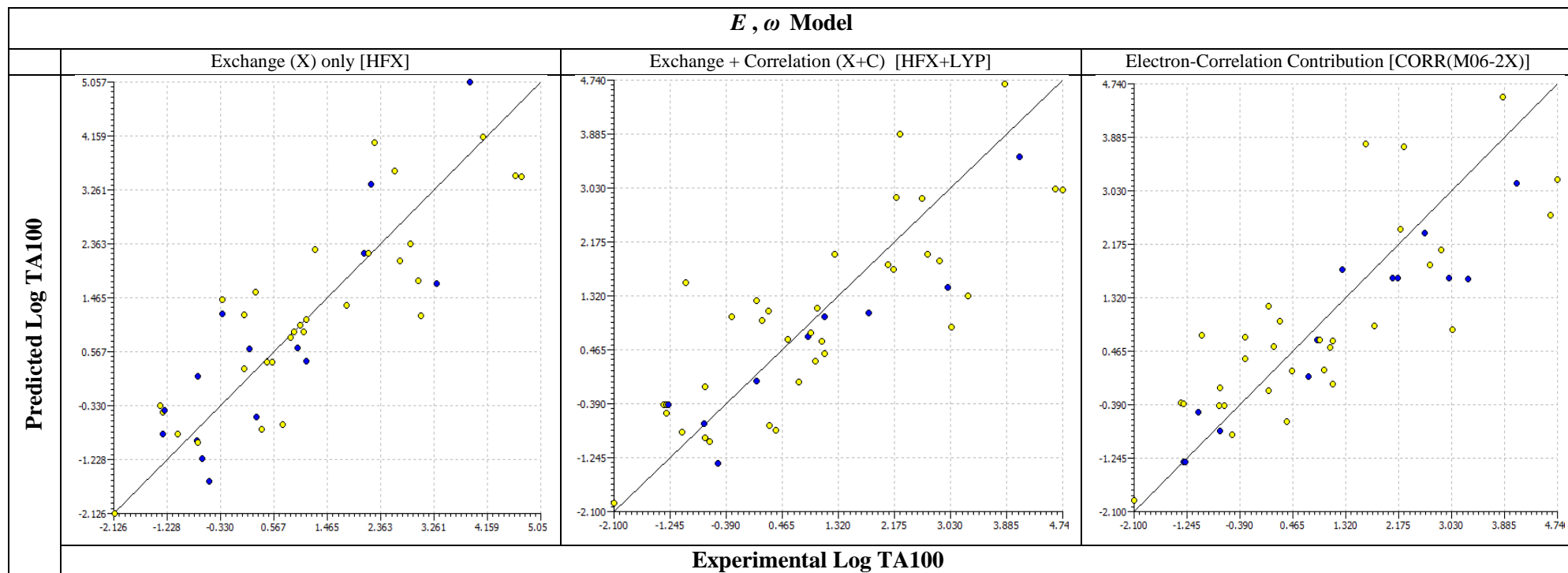


Figure S3(B). Scatter plots of the Experimental vs Predicted TA100 Mutagenicity for best models developed with Exchange (X) only, Exchange + Correlation (X+C) methods, and effect of the electron-correlation (CORR) of the energies based descriptors with 30% random splitting. Training and prediction set chemicals are represented with open (yellow) and filled (blue) circles, respectively.

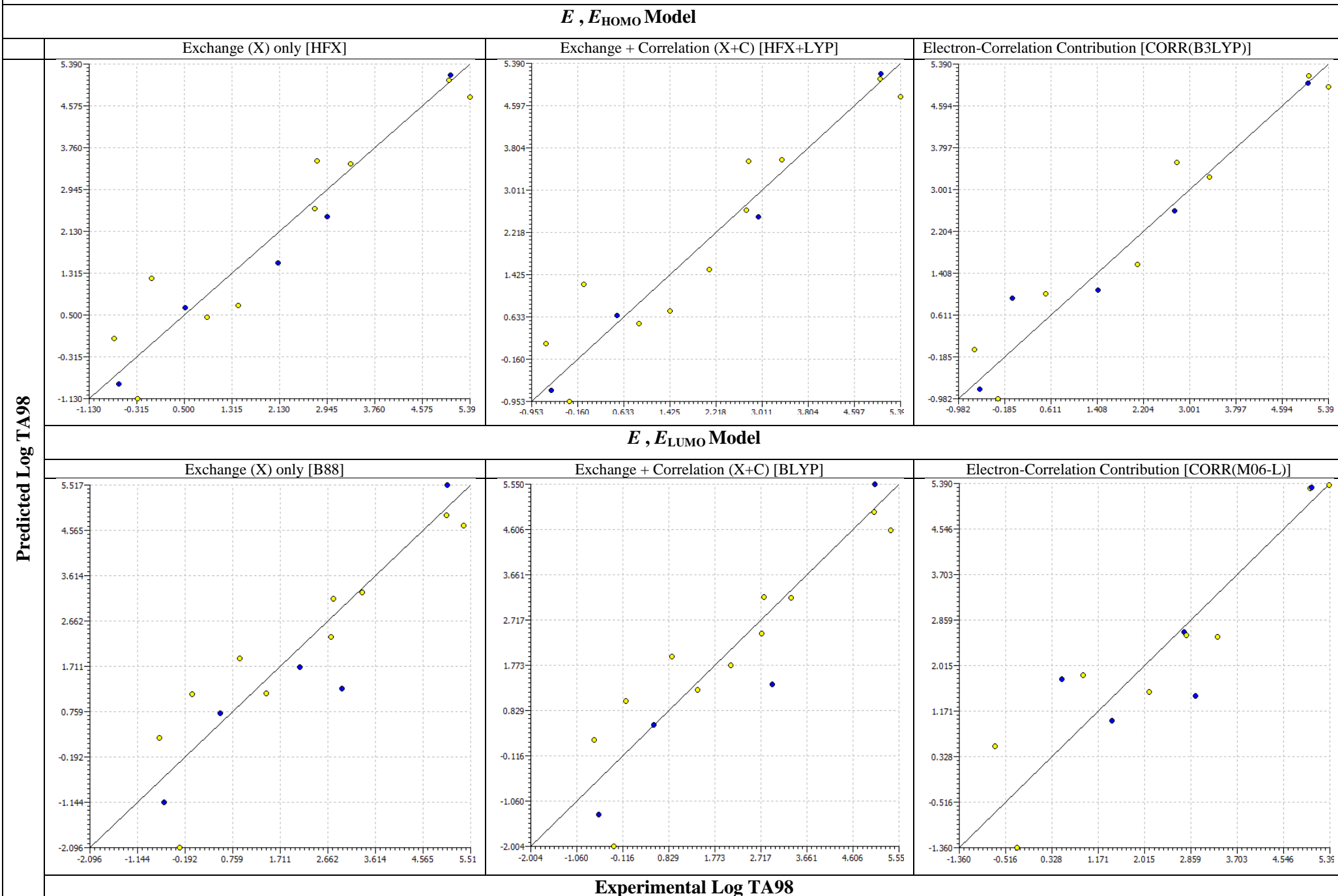


Figure S3(B) continued...

Figure S3(B) continued...

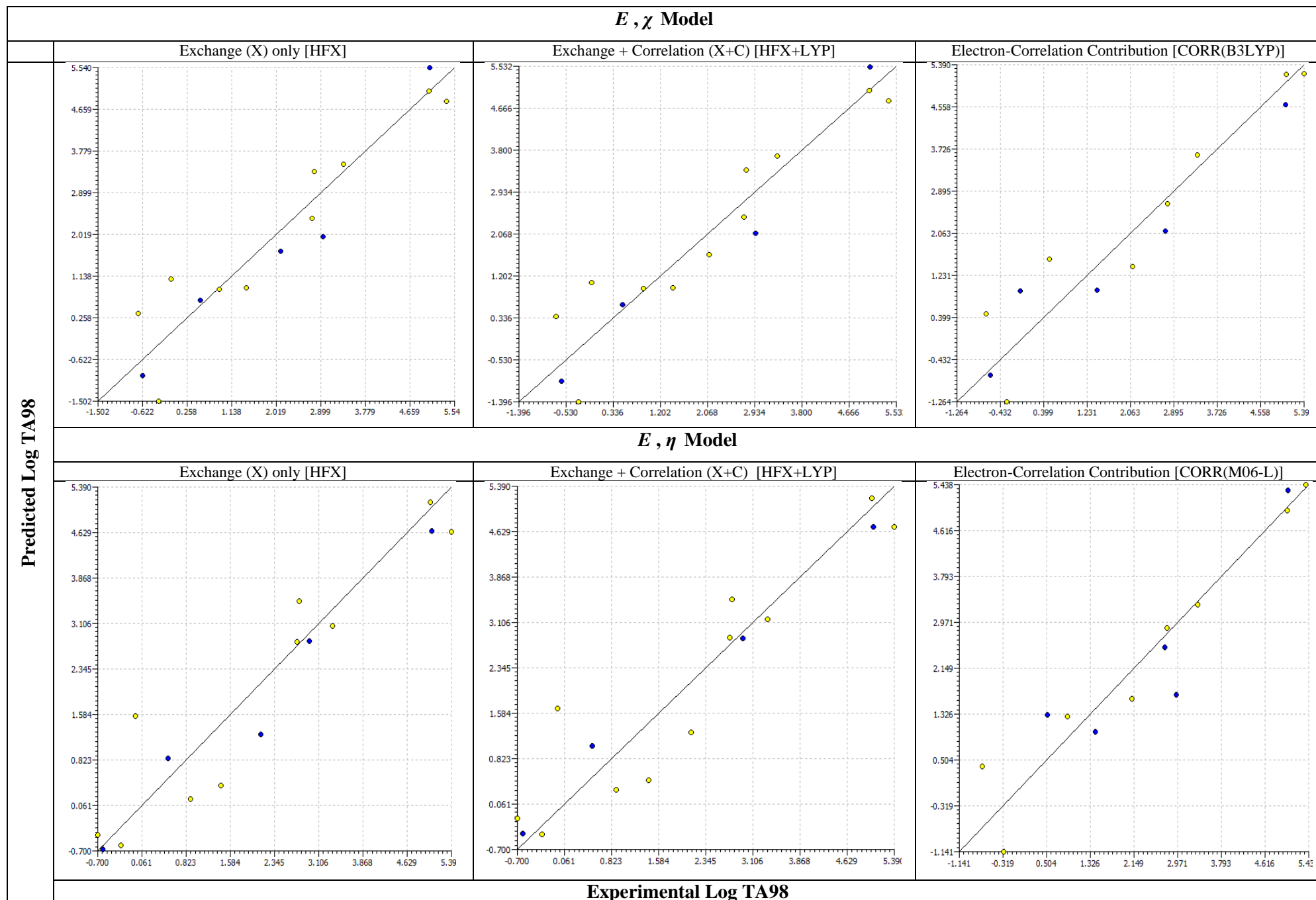


Figure S3(B) continued...

Figure S3(B) continued...

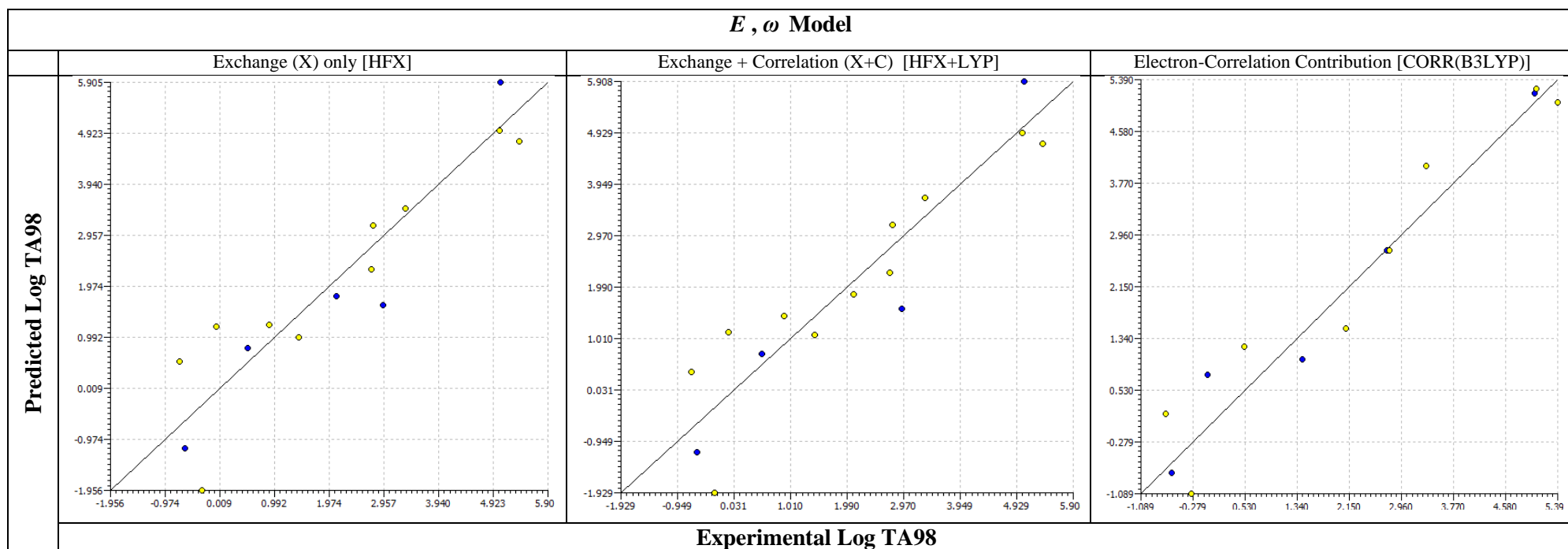


Table S1. List of nitrated-PAHs with their mutagenicity expressed as Log TA100 and Log TA98.

Name of Compound	Chemical ID number	Activity (Log TA100)	Name of Compound	Chemical ID number	Activity (Log TA98)
4-nitrotoulene	1	-2.10	1-methyl-2-nitronaphthalene	1	-0.70
2,6-dinitrotoulene	2	-1.34	1-nitronaphthlene	2	-0.61
3,4-dinitrotoulene	3	-1.30	2-nitronaphthlene	3	-0.30
2,4-dinitrotoulene	4	-1.29	1,3-dinitronaphthlene	4	-0.05
2,3-dinitrotoulene	5	-1.26	1,3-dinitrobenzene	5	0.03
6-nitroquinoline	6	-1.05	1,5-dinitronaphthlene	6	0.52
3-nitrocarbazole	7	-1.00	1,8-dinitronaphthlene	7	0.90
3,5-dinitrotoulene	8	-0.72	2-nitrofluorene	8	1.43
3-methyl-2-nitronaphthalene	9	-0.70	2-nitrophenanthrene	9	2.11
5-nitroquinoline	10	-0.70	1-nitrofluoranthene	10	2.74
2,5-dinitrotoulene	11	-0.63	1-nitropyrene	11	2.78
1,3,6,8-tetranitronaphthalene	12	-0.52	2-nitroanthracene	12	2.95
1,3-dinitrobenzene	13	-0.51	2-nitropyrene	13	3.35
2-nitrocarbazole	14	-0.30	1,3,6,8-tetranitropyrene	14	4.99
4-nitrocarbazole	15	-0.30	1,3,6-trinitropyrene	15	4.99
2,3,4-trinitrotoulene	16	0.08	1,3-dinitropyrene	16	5.04
1-methyl-2-nitronaphthalene	17	0.08	1,6-dinitropyrene	17	5.06
2,4,6-trinitrotoulene	18	0.16	1,8-dinitropyrene	18	5.39
9-nitroanthracene	19	0.26			
1-nitronaphthlene	20	0.28			
2-nitronaphthlene	21	0.37			
2,3,5-trinitrotoulene	22	0.46			
2,3,6-trinitrotoulene	23	0.55			
6-nitrobenzo[a]pyrene	24	0.71			
1,3,5-trinitrobenzene	25	0.72			
1,3-dinitronaphthlene	26	0.86			
1,5-dinitronaphthlene	27	0.91			
5-nitroacenaphthene	28	0.97			
3,4,5-trinitrotoulene	29	1.01			
2-nitrofluorene	30	1.08			
1,8-dinitronaphthlene	31	1.12			
2,4,5-trinitrotoulene	32	1.12			
2,7-dinitrofluorene	33	1.27			
1-nitrobenzo[e]pyrene	34	1.65			
2-nitrophenanthrene	35	1.79			
7-nitrofluoranthene	36	2.09			
1-nitropyrene	37	2.17			
6-nitrochrysene	38	2.21			
2,4,7-trinitro-9-fluorenone	39	2.27			
2,4,5,7-tetranitro-9-fluorenone	40	2.46			
8-nitrofluoranthene	41	2.60			
2,7-dinitro-9-fluorenone	42	2.69			
2-nitropyrene	43	2.87			
1-nitrofluoranthene	44	3.00			
2-nitroanthracene	45	3.05			
1,3,6,8-tetranitropyrene	46	3.18			
3-nitrofluoranthene	47	3.31			
1,3,6-trinitropyrene	48	3.87			
1,6-dinitropyrene	49	4.09			
1,3-dinitropyrene	50	4.63			
1,8-dinitropyrene	51	4.74			

Table S2a. Computed value of total energy (E) descriptor, (all in a.u., 1 a.u.=27.21165eV, 627.50956 kcal/mol) through exchange-only methods (HFX, B88, HF), exchange-correlation (XC) methods (HFX+LYP, BLYP, B3LYP, M06, M06-L, M06-2X), employing 6-311G(d,p) basis set, along with the descriptors incorporating mainly the effect of electron correlation (CORR) from the respective XC methods for 51-nitrated PAHs.

Name of Compound	- E_{HFX}	- E_{B88}	- E_{HF}	- $E_{\text{HFX+LYP}}$	- E_{BLYP}	- E_{B3LYP}	- E_{M06}	- $E_{\text{M06-L}}$	- $E_{\text{M06-2X}}$	- $E_{\text{CORR(HFX+LYP)}}$	- $E_{\text{CORR(BLYP)}}$	- $E_{\text{CORR(B3LYP)}}$	- $E_{\text{CORR(M06)}}$	- $E_{\text{CORR(M06-L)}}$	- $E_{\text{CORR(M06-2X)}}$
4-nitrotoluene	473.324	473.585	473.324	475.809	476.057	476.194	475.869	476.137	475.988	2.485	2.472	2.869	2.545	2.813	2.663
2,6-dinitrotoluene	676.827	677.298	676.828	680.148	680.599	680.733	680.318	680.668	680.458	3.321	3.301	3.905	3.490	3.841	3.631
3,4-dinitrotoluene	676.822	677.197	676.657	680.140	680.595	680.608	680.184	680.545	680.317	3.318	3.398	3.951	3.527	3.888	3.660
2,4-dinitrotoluene	676.837	677.306	676.837	680.157	680.606	680.741	680.325	680.675	680.464	3.320	3.300	3.904	3.488	3.838	3.627
2,3-dinitrotoluene	676.820	677.192	676.654	680.139	680.491	680.605	680.183	680.544	680.316	3.319	3.299	3.951	3.529	3.890	3.662
6-nitroquinoline	602.954	603.299	602.954	606.092	606.421	606.591	606.171	606.513	606.335	3.138	3.122	3.637	3.217	3.559	3.380
3-nitrocarbazole	717.765	718.134	717.765	721.591	721.941	722.154	721.655	722.074	721.864	3.826	3.807	4.389	3.890	4.308	4.099
3,5-dinitrotoluene	676.840	677.310	676.840	680.159	680.610	680.744	680.327	680.677	680.466	3.319	3.299	3.904	3.487	3.837	3.626
3-methyl-2-nitronaphthalene	625.999	626.306	625.999	629.376	629.666	629.865	629.422	629.793	629.600	3.378	3.360	3.866	3.424	3.794	3.602
5-nitroquinoline	602.947	603.293	602.947	606.086	606.416	606.576	606.166	606.509	606.330	3.139	3.123	3.629	3.219	3.561	3.383
2,5-dinitrotoluene	676.835	677.305	676.835	680.155	680.606	680.740	680.324	680.674	680.464	3.320	3.300	3.905	3.489	3.839	3.629
1,3,6,8-tetranitronaphthalene	1197.477	1198.417	1197.477	1203.098	1204.003	1204.169	1203.483	1204.073	1203.709	5.621	5.586	6.692	6.006	6.596	6.232
1,3-dinitrobenzene	637.793	638.266	637.793	640.853	641.307	641.415	641.031	641.354	641.158	3.059	3.041	3.622	3.238	3.560	3.365
2-nitrocarbazole	717.763	718.133	717.763	721.589	721.940	722.153	721.654	722.072	721.863	3.826	3.807	4.390	3.891	4.310	4.100
4-nitrocarbazole	717.755	718.125	717.756	721.583	721.934	722.147	721.648	722.067	721.858	3.828	3.809	4.391	3.893	4.311	4.102
2,3,4-trinitrotoluene	880.317	881.005	880.291	884.470	885.132	885.262	884.756	885.190	884.915	4.154	4.127	4.971	4.465	4.899	4.623
1-methyl-2-nitronaphthalene	625.994	626.301	625.994	629.373	629.663	629.861	629.420	629.790	629.598	3.378	3.361	3.867	3.425	3.796	3.604
2,4,6-trinitrotoluene	880.339	881.020	880.339	884.494	885.148	885.279	884.772	885.205	884.932	4.155	4.128	4.940	4.433	4.865	4.593
9-nitroanthracene	739.615	739.977	739.615	743.624	743.967	744.200	743.671	744.117	743.895	4.009	3.990	4.585	4.056	4.501	4.280
1-nitronaphthlene	586.949	587.261	586.949	590.067	590.363	590.535	590.125	590.467	590.290	3.118	3.102	3.586	3.176	3.518	3.341
2-nitronaphthlene	586.957	587.268	586.957	590.074	590.370	590.542	590.131	590.474	590.296	3.117	3.102	3.584	3.173	3.516	3.339
2,3,5-trinitrotoluene	880.167	880.918	880.167	884.329	885.144	884.876	884.641	885.085	884.794	4.162	4.226	4.709	4.474	4.918	4.627
2,3,6-trinitrotoluene	880.323	880.910	880.324	884.478	885.137	885.149	884.635	885.079	884.789	4.155	4.227	4.825	4.311	4.755	4.465
6-nitrobenzo[a]pyrene	968.071	968.502	968.071	973.400	973.806	974.128	973.426	974.024	973.740	5.329	5.305	6.057	5.355	5.953	5.670
1,3,5-trinitrobenzene	841.304	841.986	841.304	845.197	845.855	845.961	845.484	845.889	845.630	3.893	3.869	4.657	4.181	4.585	4.327
1,3-dinitronaphthlene	790.467	790.987	790.467	794.419	794.917	795.087	794.584	795.008	794.769	3.952	3.930	4.620	4.117	4.542	4.302
1,5-dinitronaphthlene	790.460	790.981	790.460	794.413	794.912	795.081	794.579	795.003	794.765	3.953	3.931	4.621	4.119	4.543	4.305
5-nitroacenaphthene	663.873	664.187	663.873	667.468	667.763	667.977	667.509	667.902	667.702	3.595	3.577	4.104	3.636	4.029	3.829
3,4,5-trinitrotoluene	880.319	881.039	880.102	884.472	885.134	885.281	884.756	885.190	884.913	4.153	4.095	5.179	4.655	5.089	4.812
2-nitrofluorene	701.758	702.091	701.758	705.558	705.873	706.098	705.598	706.017	705.808	3.801	3.782	4.340	3.840	4.260	4.050
1,8-dinitronaphthlene	790.452	790.972	790.452	794.405	794.903	795.073	794.573	794.998	794.760	3.954	3.931	4.621	4.121	4.547	4.308
2,4,5-trinitrotoluene	880.163	880.916	880.163	884.324	885.140	885.152	884.637	885.082	884.790	4.161	4.225	4.990	4.474	4.919	4.627
2,7-dinitrofluorene	905.179	905.820	905.279	909.914	910.430	910.653	910.060	910.562	910.290	4.735	4.610	5.374	4.782	5.283	5.011
1-nitrobenzo[e]pyrene	968.070	968.499	968.070	973.400	973.804	974.127	973.426	974.023	973.742	5.330	5.305	6.057	5.356	5.953	5.672
2-nitrophenanthrene	739.640	739.995	739.640	743.650	743.985	744.220	743.691	744.136	743.915	4.010	3.990	4.580	4.051	4.496	4.275
7-nitrofluoranthene	815.381	815.767	815.381	819.815	820.180	820.695	819.855	820.350	820.110	4.435	4.413	5.315	4.475	4.970	4.730
1-nitropyrene	815.402	815.786	815.402	819.838	820.202	820.461	819.877	820.372	820.132	4.437	4.416	5.060	4.475	4.971	4.730
6-nitrochrysene	892.311	892.711	892.311	897.214	897.592	897.889	897.241	897.789	897.525	4.904	4.880	5.578	4.930	5.478	5.214
2,4,7-trinitro-9-fluorenone	1182.494	1183.329	1182.494	1188.212	1189.014	1189.221	1188.512	1189.116	1188.766	5.718	5.685	6.727	6.018	6.622	6.272
2,4,5,7-tetranitro-9-fluorenone	1385.991	1387.036	1385.991	1392.545	1393.551	1393.755	1392.957	1393.644	1393.233	6.554	6.514	7.764	6.966	7.653	7.242
8-nitrofluoranthene	854.433	854.812	854.433	859.126	859.483	859.770	859.152	859.673	859.420	4.694	4.671	5.337	4.719	5.241	4.988
2,7-dinitro-9-fluorenone	978.989	979.613	978.989	983.871	984.470	984.680	984.062	984.584	984.296	4.882	4.857	5.690	5.073	5.595	5.307
2-nitropyrene	815.401	815.785	815.409	819.837	820.201	820.460	819.876	820.371	820.131	4.437	4.416	5.051	4.467	4.962	4.722
1-nitrofluoranthene	815.381	815.767	815.381	819.815	820.180	820.439	819.855	820.350	820.110	4.435	4.413	5.058	4.474	4.969	4.729
2-nitroanthracene	739.630	739.990	739.630	743.638	743.979	744.212	743.682	744.127	743.905	4.008	3.989	4.583	4.052	4.498	4.275
1,3,6,8-tetranitropyrene	1425.925	1426.940	1425.925	1432.867	1433.842	1434.094	1433.233	1433.975	1433.548	6.942	6.901	8.169	7.308	8.050	7.623
3-nitrofluoranthene	815.382	815.768	815.382	819.816	820.181	820.441	819.856	820.351	820.110	4.434	4.413	5.058	4.473	4.969	4.728
1,3,6-trinitropyrene	1222.421	1223.225	1222.421	1228.528	1229.298	1229.553	1228.784	1229.443	1229.079	6.107	6.073	7.132	6.363	7.023	6.658
1,6-dinitropyrene	1018.913	1019.506	1018.913	1024.185	1024.750	1025.008	1024.331	1024.908	1024.607	5.272	5.245	6.095	5.418	5.995	5.694
1,3-dinitropyrene	1018.912	1019.506	1018.912	1024.184	1024.751	1025.008	1024.331	1024.908	1024.606	5.272	5.245	6.096	5.419	5.997	5.694
1,8-dinitropyrene	1018.914	1019.507	1018.913	1024.185	1024.752	1025.009	1024.331	1024.910	1024.608	5.272	5.245	6.096	5.418	5.997	5.695

Table S2b. Same as **Table S2a** but for HOMO energy descriptor (E_{HOMO}).

Name of Compound	$-E_{\text{HOMO}}^{\text{HFex}}$	$-E_{\text{HOMO}}^{\text{B3LYP}}$	$-E_{\text{HOMO}}^{\text{M06}}$	$-E_{\text{HOMO}}^{\text{HFex+LYP}}$	$-E_{\text{HOMO}}^{\text{BLYP}}$	$-E_{\text{HOMO}}^{\text{B3LYP}}$	$-E_{\text{HOMO}}^{\text{M06}}$	$-E_{\text{HOMO}}^{\text{M06-L}}$	$-E_{\text{HOMO}}^{\text{M06-2X}}$	$-E_{\text{CORR}}^{\text{HFex+LYP}}$	$-E_{\text{CORR}}^{\text{BLYP}}$	$E_{\text{CORR}}^{\text{B3LYP}}$	$E_{\text{CORR}}^{\text{M06}}$	$E_{\text{CORR}}^{\text{M06-L}}$	$E_{\text{CORR}}^{\text{M06-2X}}$
4-nitrotoulene	0.363	0.199	0.363	0.397	0.233	0.278	0.289	0.252	0.329	0.034	0.034	0.085	0.074	0.112	0.035
2,6-dinitrotoulene	0.389	0.210	0.389	0.424	0.244	0.297	0.310	0.262	0.353	0.035	0.034	0.092	0.079	0.127	0.036
3,4-dinitrotoulene	0.389	0.155	0.303	0.423	0.240	0.215	0.226	0.191	0.259	0.034	0.085	0.088	0.077	0.112	0.044
2,4-dinitrotoulene	0.393	0.215	0.393	0.428	0.250	0.304	0.315	0.268	0.356	0.035	0.035	0.089	0.078	0.125	0.037
2,3-dinitrotoulene	0.386	0.155	0.303	0.420	0.184	0.215	0.226	0.191	0.259	0.034	0.029	0.088	0.077	0.112	0.044
6-nitroquinoline	0.343	0.199	0.343	0.377	0.229	0.267	0.278	0.246	0.315	0.034	0.030	0.076	0.065	0.097	0.028
3-nitrocarbazole	0.308	0.168	0.308	0.341	0.200	0.233	0.246	0.213	0.279	0.033	0.032	0.075	0.062	0.095	0.029
3,5-dinitrotoulene	0.387	0.218	0.387	0.422	0.253	0.301	0.312	0.272	0.353	0.035	0.035	0.086	0.075	0.115	0.034
3-methyl-2-nitronaphthalene	0.313	0.176	0.313	0.346	0.208	0.241	0.252	0.221	0.288	0.033	0.032	0.072	0.061	0.092	0.025
5-nitroquinoline	0.345	0.195	0.345	0.379	0.226	0.266	0.278	0.243	0.315	0.034	0.031	0.079	0.067	0.102	0.030
2,5-dinitrotoulene	0.385	0.216	0.386	0.420	0.251	0.301	0.311	0.269	0.353	0.035	0.035	0.085	0.075	0.117	0.033
1,3,6,8-tetranitronaphthalene	0.397	0.226	0.397	0.431	0.262	0.306	0.318	0.279	0.359	0.034	0.036	0.091	0.079	0.118	0.038
1,3-dinitrobenzene	0.404	0.221	0.404	0.440	0.256	0.316	0.328	0.275	0.370	0.036	0.035	0.088	0.076	0.129	0.034
2-nitrocarbazole	0.306	0.165	0.305	0.338	0.197	0.231	0.243	0.211	0.277	0.032	0.032	0.074	0.062	0.094	0.028
4-nitrocarbazole	0.300	0.161	0.301	0.333	0.192	0.226	0.238	0.206	0.272	0.033	0.031	0.075	0.063	0.095	0.029
2,3,4-trinitrotoulene	0.416	0.222	0.396	0.451	0.255	0.309	0.322	0.274	0.371	0.035	0.033	0.087	0.074	0.122	0.025
1-methyl-2-nitronaphthalene	0.314	0.176	0.314	0.347	0.208	0.241	0.252	0.221	0.289	0.033	0.032	0.073	0.062	0.093	0.025
2,4,6-trinitrotoulene	0.424	0.227	0.424	0.460	0.262	0.317	0.331	0.280	0.381	0.036	0.035	0.107	0.093	0.144	0.043
9-nitroanthracene	0.287	0.158	0.287	0.319	0.190	0.221	0.232	0.202	0.266	0.032	0.032	0.066	0.055	0.085	0.021
1-nitronaphthlene	0.319	0.179	0.319	0.352	0.212	0.246	0.257	0.226	0.293	0.033	0.033	0.073	0.062	0.093	0.021
2-nitronaphthlene	0.319	0.182	0.319	0.352	0.214	0.247	0.258	0.228	0.295	0.033	0.032	0.072	0.061	0.091	0.024
2,3,5-trinitrotoulene	0.324	0.177	0.324	0.356	0.263	0.250	0.249	0.215	0.281	0.032	0.086	0.074	0.075	0.109	0.043
2,3,6-trinitrotoulene	0.413	0.175	0.414	0.448	0.259	0.237	0.248	0.214	0.282	0.035	0.084	0.177	0.166	0.200	0.132
6-nitrobenzo[a]pyrene	0.275	0.152	0.275	0.307	0.185	0.214	0.224	0.196	0.257	0.032	0.033	0.061	0.051	0.079	0.018
1,3,5-trinitrobenzene	0.442	0.237	0.442	0.478	0.272	0.336	0.352	0.292	0.399	0.036	0.035	0.106	0.090	0.150	0.043
1,3-dinitronaphthlene	0.344	0.200	0.344	0.378	0.233	0.268	0.279	0.247	0.316	0.034	0.033	0.076	0.065	0.097	0.028
1,5-dinitronaphthlene	0.347	0.197	0.347	0.381	0.232	0.268	0.280	0.246	0.317	0.034	0.035	0.079	0.067	0.101	0.030
5-nitroacenaphthene	0.305	0.170	0.305	0.337	0.202	0.234	0.244	0.213	0.279	0.032	0.032	0.071	0.061	0.092	0.026
3,4,5-trinitrotoulene	0.412	0.192	0.346	0.446	0.254	0.270	0.317	0.272	0.365	0.034	0.062	0.076	0.029	0.074	-0.019
2-nitrofluorene	0.314	0.180	0.314	0.348	0.212	0.245	0.255	0.225	0.291	0.034	0.032	0.069	0.059	0.089	0.023
1,8-dinitronaphthlene	0.347	0.193	0.347	0.381	0.227	0.266	0.277	0.243	0.316	0.034	0.034	0.081	0.070	0.104	0.031
2,4,5-trinitrotoulene	0.322	0.174	0.322	0.354	0.259	0.236	0.246	0.213	0.279	0.032	0.085	0.086	0.076	0.109	0.043
2,7-dinitrofluorene	0.343	0.201	0.343	0.377	0.234	0.268	0.279	0.247	0.315	0.034	0.033	0.075	0.064	0.096	0.028
1-nitrobenzo[e]pyrene	0.286	0.161	0.286	0.319	0.193	0.224	0.234	0.206	0.268	0.033	0.032	0.062	0.052	0.080	0.018
2-nitrophenanthrene	0.308	0.176	0.308	0.341	0.208	0.240	0.251	0.221	0.286	0.033	0.032	0.068	0.057	0.087	0.022
7-nitrofluoranthene	0.303	0.173	0.303	0.336	0.205	0.237	0.247	0.218	0.282	0.033	0.032	0.066	0.056	0.085	0.021
1-nitropyrene	0.287	0.164	0.287	0.320	0.196	0.226	0.236	0.208	0.268	0.033	0.032	0.061	0.051	0.079	0.019
6-nitrochrysene	0.297	0.167	0.297	0.330	0.200	0.231	0.242	0.212	0.276	0.033	0.033	0.066	0.055	0.085	0.021
2,4,7-trinitro-9-fluorenone	0.382	0.216	0.382	0.416	0.250	0.299	0.311	0.266	0.349	0.034	0.034	0.083	0.071	0.116	0.033
2,4,5,7-tetranitro-9-fluorenone	0.399	0.227	0.399	0.434	0.261	0.314	0.325	0.278	0.364	0.035	0.034	0.085	0.074	0.121	0.035
8-nitrofluoranthene	0.288	0.161	0.288	0.320	0.193	0.223	0.234	0.204	0.268	0.032	0.032	0.065	0.054	0.084	0.020
2,7-dinitro-9-fluorenone	0.362	0.205	0.363	0.397	0.238	0.284	0.295	0.254	0.332	0.035	0.033	0.079	0.068	0.109	0.031
2-nitropyrene	0.284	0.162	0.284	0.316	0.194	0.223	0.234	0.206	0.267	0.032	0.032	0.061	0.050	0.078	0.017
1-nitrofluoranthene	0.305	0.173	0.305	0.338	0.205	0.237	0.248	0.218	0.284	0.033	0.032	0.068	0.057	0.087	0.021
2-nitroanthracene	0.285	0.161	0.285	0.317	0.193	0.222	0.233	0.205	0.266	0.032	0.032	0.063	0.052	0.080	0.019
1,3,6,8-tetranitropyrene	0.352	0.206	0.352	0.386	0.242	0.277	0.287	0.256	0.322	0.034	0.036	0.075	0.065	0.096	0.030
3-nitrofluoranthene	0.307	0.175	0.307	0.340	0.207	0.239	0.250	0.220	0.286	0.033	0.032	0.068	0.057	0.087	0.021
1,3,6-trinitropyrene	0.331	0.194	0.331	0.365	0.229	0.261	0.272	0.242	0.305	0.034	0.035	0.070	0.059	0.089	0.026
1,6-dinitropyrene	0.305	0.179	0.305	0.338	0.212	0.242	0.252	0.224	0.286	0.033	0.033	0.063	0.053	0.081	0.019
1,3-dinitropyrene	0.308	0.180	0.308	0.341	0.213	0.244	0.254	0.226	0.286	0.033	0.033	0.064	0.054	0.082	0.022
1,8-dinitropyrene	0.310	0.180	0.307	0.343	0.214	0.244	0.255	0.226	0.288	0.033	0.034	0.063	0.052	0.081	0.019

Table S2c. Same as **Table S2a** but for LUMO energy descriptor (E_{LUMO}).

Name of Compound	E_{HFex}^{LUMO}	$-E_{BBB}^{LUMO}$	E_{HF}^{LUMO}	$E_{HFex+LYP}^{LUMO}$	$-E_{BLYP}^{LUMO}$	$-E_{B3LYP}^{LUMO}$	$-E_{M06}^{LUMO}$	$-E_{M06-L}^{LUMO}$	$-E_{M06-2X}^{LUMO}$	$-E_{CORR(HFex+LYP)}^{LUMO}$	$-E_{CORR(BLYP)}^{LUMO}$	$-E_{CORR(B3LYP)}^{LUMO}$	$-E_{CORR(M06)}^{LUMO}$	$-E_{CORR(M06-L)}^{LUMO}$	$-E_{CORR(M06-2X)}^{LUMO}$
4-nitrotoulene	0.060	0.085	0.060	0.030	0.115	0.092	0.084	0.113	0.048	0.030	0.030	0.152	0.144	0.173	0.108
2,6-dinitrotoulene	0.040	0.100	0.040	0.006	0.133	0.111	0.103	0.132	0.068	0.034	0.033	0.151	0.143	0.172	0.108
3,4-dinitrotoulene	0.038	0.100	0.001	0.005	0.134	0.122	0.117	0.134	0.091	0.033	0.034	0.123	0.118	0.135	0.092
2,4-dinitrotoulene	0.031	0.105	0.031	0.000	0.136	0.115	0.107	0.135	0.074	0.031	0.031	0.146	0.138	0.166	0.105
2,3-dinitrotoulene	0.039	0.100	0.002	0.006	0.131	0.122	0.117	0.134	0.091	0.033	0.031	0.124	0.119	0.136	0.093
6-nitroquinoline	0.039	0.095	0.039	0.008	0.126	0.105	0.099	0.126	0.066	0.031	0.031	0.144	0.138	0.165	0.105
3-nitrocarbazole	0.069	0.074	0.069	0.040	0.104	0.082	0.075	0.102	0.040	0.029	0.030	0.151	0.144	0.171	0.109
3,5-dinitrotoulene	0.025	0.107	0.025	-0.005	0.139	0.118	0.111	0.138	0.078	0.030	0.032	0.143	0.136	0.163	0.103
3-methyl-2-nitronaphthalene	0.052	0.084	0.052	0.019	0.115	0.094	0.087	0.115	0.054	0.033	0.031	0.146	0.139	0.167	0.106
5-nitroquinoline	0.038	0.094	0.038	0.006	0.127	0.106	0.100	0.127	0.067	0.032	0.033	0.144	0.138	0.165	0.105
2,5-dinitrotoulene	0.015	0.114	0.016	-0.017	0.148	0.128	0.121	0.148	0.087	0.032	0.034	0.144	0.137	0.164	0.103
1,3,6,8-tetranitronaphthalene	-0.021	0.135	-0.021	-0.053	0.168	0.153	0.147	0.171	0.120	0.032	0.033	0.132	0.126	0.150	0.099
1,3-dinitrobenzene	0.023	0.111	0.023	-0.007	0.142	0.121	0.114	0.142	0.081	0.030	0.031	0.144	0.137	0.165	0.104
2-nitrocarbazole	0.047	0.082	0.047	0.017	0.112	0.092	0.086	0.112	0.054	0.030	0.030	0.139	0.133	0.159	0.101
4-nitrocarbazole	0.047	0.082	0.052	0.017	0.112	0.091	0.087	0.112	0.054	0.030	0.030	0.143	0.139	0.164	0.106
2,3,4-trinitrotoulene	0.020	0.113	0.010	-0.011	0.146	0.125	0.118	0.146	0.085	0.031	0.033	0.135	0.128	0.156	0.095
1-methyl-2-nitronaphthalene	0.057	0.083	0.057	0.025	0.113	0.091	0.084	0.112	0.051	0.032	0.030	0.148	0.141	0.169	0.108
2,4,6-trinitrotoulene	0.008	0.121	0.008	-0.022	0.153	0.134	0.126	0.153	0.095	0.030	0.032	0.142	0.134	0.161	0.103
9-nitroanthracene	0.036	0.084	0.036	0.004	0.117	0.098	0.093	0.118	0.066	0.032	0.033	0.134	0.129	0.154	0.102
1-nitronaphthlene	0.049	0.086	0.049	0.016	0.118	0.097	0.090	0.118	0.057	0.033	0.032	0.146	0.139	0.167	0.106
2-nitronaphthlene	0.045	0.088	0.045	0.015	0.118	0.098	0.091	0.118	0.059	0.030	0.030	0.143	0.136	0.163	0.104
2,3,5-trinitrotoulene	-0.027	0.126	-0.027	-0.058	0.154	0.169	0.145	0.162	0.118	0.031	0.028	0.142	0.118	0.135	0.091
2,3,6-trinitrotoulene	0.000	0.122	0.001	-0.034	0.160	0.146	0.141	0.159	0.115	0.034	0.038	0.147	0.142	0.160	0.116
6-nitrobenzo[a]pyrene	0.032	0.083	0.032	0.000	0.116	0.098	0.094	0.117	0.068	0.032	0.033	0.130	0.126	0.149	0.100
1,3,5-trinitrobenzene	0.000	0.128	0.000	-0.031	0.160	0.141	0.134	0.161	0.102	0.031	0.032	0.141	0.134	0.161	0.102
1,3-dinitronaphthlene	0.017	0.107	0.017	-0.015	0.140	0.121	0.115	0.141	0.085	0.032	0.033	0.138	0.132	0.158	0.102
1,5-dinitronaphthlene	0.022	0.104	0.022	-0.011	0.138	0.119	0.112	0.139	0.081	0.033	0.034	0.141	0.134	0.161	0.103
5-nitroacenaphthene	0.051	0.082	0.051	0.019	0.113	0.093	0.086	0.112	0.054	0.032	0.031	0.144	0.137	0.163	0.105
3,4,5-trinitrotoulene	0.016	0.145	-0.005	-0.017	0.149	0.131	0.126	0.152	0.094	0.033	0.004	0.126	0.121	0.147	0.089
2-nitrofluorene	0.046	0.085	0.046	0.016	0.115	0.095	0.088	0.114	0.056	0.030	0.030	0.141	0.134	0.160	0.102
1,8-dinitronaphthlene	0.029	0.095	0.029	-0.001	0.126	0.108	0.102	0.128	0.073	0.030	0.031	0.137	0.131	0.157	0.102
2,4,5-trinitrotoulene	-0.027	0.124	-0.027	-0.059	0.159	0.150	0.145	0.163	0.119	0.032	0.035	0.123	0.118	0.136	0.092
2,7-dinitrofluorene	0.016	0.105	0.016	-0.014	0.136	0.118	0.112	0.137	0.082	0.030	0.031	0.134	0.128	0.153	0.098
1-nitrobenzo[e]pyrene	0.042	0.083	0.042	0.011	0.114	0.094	0.088	0.115	0.060	0.031	0.031	0.136	0.130	0.157	0.102
2-nitrophenanthrene	0.043	0.087	0.043	0.012	0.117	0.097	0.091	0.117	0.059	0.031	0.030	0.140	0.134	0.160	0.102
7-nitrofluoranthene	0.037	0.086	0.037	0.007	0.116	0.097	0.091	0.117	0.063	0.030	0.030	0.134	0.128	0.154	0.100
1-nitropyrene	0.031	0.088	0.031	0.000	0.119	0.102	0.097	0.120	0.068	0.031	0.031	0.133	0.128	0.151	0.099
6-nitrochrysene	0.043	0.085	0.043	0.011	0.117	0.097	0.091	0.117	0.061	0.032	0.032	0.140	0.134	0.160	0.104
2,4,7-trinitro-9-fluorenone	-0.024	0.136	-0.024	-0.056	0.169	0.154	0.148	0.172	0.120	0.032	0.033	0.130	0.124	0.148	0.096
2,4,5,7-tetranitro-9-fluorenone	-0.040	0.148	-0.040	-0.073	0.182	0.167	0.161	0.185	0.135	0.033	0.034	0.127	0.121	0.145	0.095
8-nitrofluoranthene	0.059	0.083	0.059	0.029	0.113	0.091	0.084	0.112	0.048	0.030	0.030	0.150	0.143	0.171	0.107
2,7-dinitro-9-fluorenone	-0.005	0.122	-0.006	-0.036	0.154	0.138	0.132	0.157	0.103	0.031	0.032	0.132	0.126	0.151	0.097
2-nitropyrene	0.037	0.086	0.045	0.006	0.118	0.099	0.093	0.119	0.064	0.031	0.032	0.144	0.138	0.164	0.109
1-nitrofluoranthene	0.024	0.094	0.024	-0.007	0.125	0.108	0.103	0.127	0.076	0.031	0.031	0.132	0.127	0.151	0.100
2-nitroanthracene	0.030	0.091	0.030	0.000	0.121	0.103	0.098	0.123	0.070	0.030	0.030	0.133	0.128	0.153	0.100
1,3,6,8-tetranitropyrene	-0.038	0.137	-0.038	-0.073	0.173	0.161	0.155	0.176	0.131	0.035	0.036	0.123	0.117	0.138	0.093
3-nitrofluoranthene	0.021	0.095	0.021	-0.010	0.127	0.111	0.106	0.130	0.078	0.031	0.032	0.132	0.127	0.151	0.099
1,3,6-trinitropyrene	-0.017	0.123	-0.017	-0.051	0.157	0.143	0.138	0.160	0.112	0.034	0.034	0.126	0.121	0.143	0.095
1,6-dinitropyrene	0.017	0.101	0.017	-0.015	0.134	0.117	0.111	0.135	0.084	0.032	0.033	0.134	0.128	0.152	0.101
1,3-dinitropyrene	0.004	0.106	0.004	-0.028	0.139	0.124	0.118	0.141	0.092	0.032	0.033	0.128	0.122	0.145	0.096
1,8-dinitropyrene	0.006	0.106	0.012	-0.026	0.140	0.124	0.122	0.142	0.091	0.032	0.034	0.136	0.134	0.154	0.103

Table S2d. Same as **Table S2a** but for absolute electronegativity descriptor (χ).

Name of Compound	χ_{HFx}	χ_{B88}	χ_{HF}	$\chi_{\text{HFx+LYP}}$	χ_{BLYP}	χ_{B3LYP}	χ_{M06}	$\chi_{\text{M06-L}}$	$\chi_{\text{M06-2X}}$	$\chi_{\text{CORR(HFx+LYP)}}$	$\chi_{\text{CORR(BLYP)}}$	$\chi_{\text{CORR(B3LYP)}}$	$\chi_{\text{CORR(M06)}}$	$\chi_{\text{CORR(M06-L)}}$	$\chi_{\text{CORR(M06-2X)}}$
4-nitrotoluene	0.152	0.142	0.152	0.184	0.174	0.185	0.187	0.182	0.188	0.032	0.032	0.033	0.035	0.030	0.037
2,6-dinitrotoluene	0.175	0.155	0.175	0.209	0.189	0.204	0.207	0.197	0.211	0.035	0.034	0.030	0.032	0.023	0.036
3,4-dinitrotoluene	0.176	0.128	0.151	0.209	0.187	0.169	0.171	0.163	0.175	0.034	0.060	0.018	0.020	0.012	0.024
2,4-dinitrotoluene	0.181	0.160	0.181	0.214	0.193	0.210	0.211	0.202	0.215	0.033	0.033	0.029	0.030	0.021	0.034
2,3-dinitrotoluene	0.174	0.128	0.151	0.207	0.158	0.169	0.172	0.163	0.175	0.034	0.030	0.018	0.021	0.012	0.025
6-nitroquinoline	0.152	0.147	0.152	0.185	0.178	0.186	0.188	0.186	0.191	0.033	0.031	0.034	0.036	0.034	0.039
3-nitrocarbazole	0.120	0.121	0.120	0.151	0.152	0.158	0.160	0.158	0.159	0.031	0.031	0.038	0.041	0.038	0.040
3,5-dinitrotoluene	0.181	0.163	0.181	0.214	0.196	0.210	0.211	0.205	0.215	0.033	0.034	0.029	0.030	0.024	0.034
3-methyl-2-nitronaphthalene	0.131	0.130	0.131	0.164	0.162	0.168	0.170	0.168	0.171	0.033	0.032	0.037	0.039	0.037	0.041
5-nitroquinoline	0.154	0.145	0.154	0.187	0.177	0.186	0.189	0.185	0.191	0.033	0.032	0.033	0.035	0.031	0.038
2,5-dinitrotoluene	0.185	0.165	0.185	0.219	0.200	0.215	0.216	0.209	0.220	0.034	0.035	0.030	0.031	0.024	0.035
1,3,6,8-tetranitronaphthalene	0.209	0.181	0.209	0.242	0.215	0.230	0.233	0.225	0.240	0.033	0.035	0.021	0.024	0.016	0.031
1,3-dinitrobenzene	0.191	0.166	0.191	0.224	0.199	0.219	0.221	0.209	0.225	0.033	0.033	0.028	0.031	0.018	0.035
2-nitrocarbazole	0.130	0.124	0.129	0.161	0.155	0.162	0.165	0.161	0.166	0.031	0.031	0.033	0.036	0.032	0.037
4-nitrocarbazole	0.127	0.122	0.125	0.158	0.152	0.159	0.162	0.159	0.163	0.032	0.031	0.034	0.038	0.034	0.039
2,3,4-trinitrotoluene	0.198	0.168	0.193	0.231	0.201	0.217	0.220	0.210	0.228	0.033	0.033	0.024	0.027	0.017	0.035
1-methyl-2-nitronaphthalene	0.129	0.130	0.129	0.161	0.161	0.166	0.168	0.166	0.170	0.033	0.031	0.038	0.040	0.038	0.041
2,4,6-trinitrotoluene	0.208	0.174	0.208	0.241	0.208	0.226	0.229	0.217	0.238	0.033	0.034	0.018	0.021	0.009	0.030
9-nitroanthracene	0.126	0.121	0.126	0.158	0.154	0.160	0.162	0.160	0.166	0.032	0.033	0.034	0.037	0.034	0.040
1-nitronaphthlene	0.135	0.133	0.135	0.168	0.165	0.172	0.174	0.172	-0.118	0.033	0.033	0.037	0.039	0.037	-0.253
2-nitronaphthlene	0.137	0.135	0.137	0.169	0.166	0.173	0.175	0.173	0.177	0.032	0.031	0.036	0.038	0.036	0.040
2,3,5-trinitrotoluene	0.176	0.152	0.176	0.207	0.209	0.210	0.197	0.189	0.200	0.032	0.057	0.034	0.021	0.013	0.024
2,3,6-trinitrotoluene	0.207	0.149	0.207	0.241	0.210	0.192	0.195	0.186	0.198	0.035	0.061	-0.015	-0.012	-0.020	-0.008
6-nitrobenzo[a]pyrene	0.122	0.118	0.122	0.154	0.151	0.156	0.159	0.157	0.162	0.032	0.033	0.035	0.037	0.035	0.041
1,3,5-trinitrobenzene	0.221	0.183	0.221	0.255	0.216	0.239	0.243	0.226	0.251	0.034	0.034	0.018	0.022	0.005	0.030
1,3-dinitronaphthlene	0.164	0.154	0.164	0.197	0.187	0.195	0.197	0.194	0.201	0.033	0.033	0.031	0.034	0.031	0.037
1,5-dinitronaphthlene	0.163	0.151	0.163	0.196	0.185	0.194	0.196	0.192	0.199	0.034	0.035	0.031	0.033	0.030	0.037
5-nitroacenaphthene	0.127	0.126	0.127	0.159	0.158	0.164	0.165	0.163	0.167	0.032	0.032	0.037	0.038	0.036	0.040
3,4,5-trinitrotoluene	0.198	0.169	0.176	0.232	0.202	0.201	0.221	0.212	0.230	0.034	0.033	0.025	0.046	0.036	0.054
2-nitrofluorene	0.134	0.133	0.134	0.166	0.164	0.170	0.172	0.169	0.174	0.032	0.031	0.036	0.038	0.035	0.040
1,8-dinitronaphthlene	0.159	0.144	0.159	0.191	0.177	0.187	0.189	0.185	0.195	0.032	0.033	0.028	0.030	0.026	0.036
2,4,5-trinitrotoluene	0.175	0.149	0.175	0.207	0.209	0.193	0.196	0.188	0.199	0.032	0.060	0.019	0.021	0.013	0.025
2,7-dinitrofluorene	0.164	0.153	0.164	0.196	0.185	0.193	0.196	0.192	0.198	0.032	0.032	0.030	0.032	0.029	0.035
1-nitrobenzo[e]pyrene	0.122	0.122	0.122	0.154	0.154	0.159	0.161	0.160	0.164	0.032	0.032	0.037	0.039	0.038	0.042
2-nitrophenanthrene	0.133	0.132	0.133	0.165	0.163	0.169	0.171	0.169	0.173	0.032	0.031	0.036	0.038	0.036	0.040
7-nitrofluoranthene	0.133	0.130	0.133	0.165	0.161	0.167	0.169	0.167	0.173	0.032	0.031	0.034	0.036	0.034	0.040
1-nitropyrene	0.128	0.126	0.128	0.160	0.158	0.164	0.166	0.164	0.168	0.032	0.032	0.036	0.038	0.036	0.040
6-nitrochrysene	0.127	0.126	0.127	0.160	0.159	0.164	0.167	0.165	0.168	0.033	0.033	0.037	0.040	0.038	0.041
2,4,7-trinitro-9-fluorenone	0.203	0.176	0.203	0.236	0.210	0.227	0.229	0.219	0.235	0.033	0.034	0.024	0.026	0.016	0.032
2,4,5,7-tetranitro-9-fluorenone	0.220	0.188	0.220	0.254	0.222	0.241	0.243	0.232	0.249	0.034	0.034	0.021	0.024	0.012	0.030
8-nitrofluoranthene	0.115	0.122	0.115	0.146	0.153	0.157	0.159	0.158	0.158	0.031	0.031	0.043	0.044	0.043	0.044
2,7-dinitro-9-fluorenone	0.184	0.164	0.185	0.217	0.196	0.211	0.213	0.206	0.218	0.033	0.033	0.027	0.029	0.021	0.033
2-nitropyrene	0.124	0.124	0.120	0.155	0.156	0.161	0.164	0.162	0.166	0.032	0.032	0.042	0.044	0.043	0.046
1-nitrofluoranthene	0.141	0.134	0.141	0.173	0.165	0.173	0.175	0.173	0.180	0.032	0.032	0.032	0.035	0.032	0.039
2-nitroanthracene	0.128	0.126	0.128	0.159	0.157	0.163	0.166	0.164	0.168	0.031	0.031	0.035	0.038	0.036	0.041
1,3,6,8-tetranitropyrene	0.195	0.172	0.195	0.230	0.208	0.219	0.221	0.216	0.227	0.035	0.036	0.024	0.026	0.021	0.032
3-nitrofluoranthene	0.143	0.135	0.143	0.175	0.167	0.175	0.178	0.175	0.182	0.032	0.032	0.032	0.035	0.032	0.039
1,3,6-trinitropyrene	0.174	0.159	0.174	0.208	0.193	0.202	0.205	0.201	0.209	0.034	0.035	0.028	0.031	0.027	0.035
1,6-dinitropyrene	0.144	0.140	0.144	0.177	0.173	0.180	0.181	0.179	0.185	0.033	0.033	0.036	0.037	0.035	0.041
1,3-dinitropyrene	0.152	0.143	0.152	0.185	0.176	0.184	0.186	0.183	0.189	0.033	0.033	0.032	0.034	0.031	0.037
1,8-dinitropyrene	0.152	0.143	0.148	0.185	0.177	0.184	0.189	0.184	0.190	0.033	0.034	0.037	0.041	0.037	0.042

Table S2e. Same as **Table S2a** but for chemical hardness descriptor (η).

Name of Compound	η_{HFx}	η_{B88}	η_{HF}	$\eta_{\text{HFx+LYP}}$	η_{BLYP}	η_{B3LYP}	η_{M06}	$\eta_{\text{M06-L}}$	$\eta_{\text{M06-2X}}$	$\eta_{\text{CORR(HFx+LYP)}}$	$\eta_{\text{CORR(BLYP)}}$	$-\eta_{\text{COR(B3LYP)}}$	$\eta_{\text{CORR(M06)}}$	$\eta_{\text{CORR(M06-L)}}$	$\eta_{\text{CORR(M06-2X)}}$
4-nitrotoulene	0.423	0.114	0.423	0.427	0.118	0.186	0.205	0.139	0.280	0.004	0.004	0.237	0.070	0.061	0.073
2,6-dinitrotoulene	0.429	0.110	0.429	0.430	0.111	0.186	0.207	0.130	0.285	0.001	0.001	0.243	0.064	0.045	0.072
3,4-dinitrotoulene	0.427	0.055	0.304	0.428	0.106	0.093	0.109	0.057	0.168	0.001	0.051	0.211	0.041	0.023	0.048
2,4-dinitrotoulene	0.424	0.110	0.424	0.428	0.114	0.189	0.208	0.133	0.282	0.004	0.004	0.235	0.061	0.042	0.069
2,3-dinitrotoulene	0.425	0.055	0.305	0.426	0.053	0.093	0.109	0.057	0.169	0.001	-0.002	0.212	0.042	0.025	0.049
6-nitroquinoline	0.382	0.104	0.382	0.385	0.103	0.162	0.180	0.120	0.249	0.003	-0.001	0.220	0.073	0.068	0.078
3-nitrocarbazole	0.377	0.094	0.377	0.381	0.096	0.151	0.171	0.111	0.240	0.004	0.002	0.226	0.082	0.076	0.080
3,5-dinitrotoulene	0.412	0.111	0.412	0.417	0.114	0.183	0.201	0.133	0.275	0.005	0.003	0.229	0.061	0.048	0.069
3-methyl-2-nitronaphthalene	0.365	0.092	0.365	0.365	0.093	0.147	0.165	0.106	0.234	0.000	0.001	0.218	0.078	0.075	0.081
5-nitroquinoline	0.383	0.101	0.383	0.385	0.099	0.160	0.178	0.116	0.248	0.002	-0.002	0.223	0.071	0.063	0.075
2,5-dinitrotoulene	0.400	0.102	0.402	0.403	0.103	0.173	0.190	0.121	0.266	0.003	0.001	0.229	0.062	0.047	0.070
1,3,6,8-tetranitronaphthalene	0.376	0.091	0.376	0.378	0.094	0.153	0.171	0.108	0.239	0.002	0.003	0.223	0.047	0.033	0.061
1,3-dinitrobenzene	0.427	0.110	0.427	0.433	0.114	0.195	0.214	0.133	0.289	0.006	0.004	0.232	0.061	0.036	0.070
2-nitrocarbazole	0.353	0.083	0.352	0.355	0.085	0.139	0.157	0.099	0.223	0.002	0.002	0.213	0.071	0.064	0.073
4-nitrocarbazole	0.347	0.079	0.353	0.350	0.080	0.135	0.151	0.093	0.218	0.003	0.001	0.218	0.076	0.069	0.077
2,3,4-trinitrotoulene	0.436	0.109	0.406	0.440	0.109	0.184	0.203	0.128	0.286	0.004	0.000	0.222	0.054	0.033	0.070
1-methyl-2-nitronaphthalene	0.371	0.093	0.371	0.372	0.095	0.150	0.168	0.109	0.238	0.001	0.002	0.221	0.079	0.076	0.082
2,4,6-trinitrotoulene	0.432	0.106	0.432	0.438	0.109	0.183	0.205	0.127	0.286	0.006	0.003	0.249	0.041	0.017	0.059
9-nitroanthracene	0.323	0.074	0.323	0.323	0.073	0.123	0.139	0.084	0.200	0.000	-0.001	0.200	0.074	0.069	0.080
1-nitronaphthlene	0.368	0.093	0.368	0.368	0.094	0.149	0.167	0.108	-0.351	0.000	0.001	0.219	0.077	0.074	-0.506
2-nitronaphthlene	0.364	0.094	0.364	0.367	0.096	0.149	0.167	0.109	0.236	0.003	0.002	0.215	0.076	0.072	0.079
2,3,5-trinitrotoulene	0.297	0.051	0.297	0.298	0.109	0.081	0.104	0.052	0.163	0.001	0.058	0.216	0.042	0.026	0.049
2,3,6-trinitrotoulene	0.413	0.053	0.415	0.414	0.099	0.091	0.107	0.056	0.167	0.001	0.046	0.324	-0.024	-0.040	-0.017
6-nitrobenzo[a]pyrene	0.307	0.069	0.307	0.307	0.069	0.116	0.131	0.079	0.188	0.000	0.000	0.191	0.075	0.070	0.082
1,3,5-trinitrobenzene	0.442	0.109	0.442	0.447	0.112	0.195	0.218	0.131	0.297	0.005	0.003	0.247	0.043	0.011	0.059
1,3-dinitronaphthlene	0.361	0.093	0.361	0.363	0.093	0.147	0.164	0.106	0.232	0.002	0.000	0.214	0.067	0.061	0.074
1,5-dinitronaphthlene	0.369	0.093	0.369	0.370	0.094	0.149	0.167	0.107	0.236	0.001	0.001	0.220	0.067	0.060	0.073
5-nitroacenaphthene	0.356	0.088	0.356	0.356	0.089	0.141	0.158	0.101	0.225	0.000	0.001	0.215	0.077	0.072	0.080
3,4,5-trinitrotoulene	0.428	0.047	0.341	0.429	0.105	0.139	0.191	0.119	0.271	0.001	0.058	0.202	0.092	0.073	0.109
2-nitrofluorene	0.360	0.095	0.360	0.364	0.097	0.150	0.167	0.110	0.235	0.004	0.002	0.210	0.075	0.071	0.079
1,8-dinitronaphthlene	0.376	0.098	0.376	0.380	0.101	0.158	0.175	0.115	0.242	0.004	0.003	0.218	0.061	0.052	0.071
2,4,5-trinitrotoulene	0.295	0.050	0.295	0.295	0.100	0.086	0.102	0.049	0.161	0.000	0.050	0.209	0.042	0.027	0.049
2,7-dinitrofluorene	0.359	0.096	0.359	0.363	0.098	0.150	0.167	0.110	0.234	0.004	0.002	0.209	0.064	0.057	0.070
1-nitrobenzo[e]pyrene	0.328	0.078	0.328	0.330	0.079	0.130	0.146	0.091	0.208	0.002	0.001	0.198	0.078	0.076	0.084
2-nitrophenanthrene	0.351	0.089	0.351	0.353	0.091	0.143	0.160	0.104	0.227	0.002	0.002	0.208	0.076	0.073	0.080
7-nitrofluoranthene	0.340	0.087	0.340	0.343	0.089	0.140	0.156	0.101	0.219	0.003	0.002	0.200	0.072	0.069	0.080
1-nitropyrene	0.318	0.076	0.318	0.320	0.077	0.124	0.139	0.088	0.200	0.002	0.001	0.194	0.076	0.073	0.081
6-nitrochrysene	0.340	0.082	0.340	0.341	0.083	0.134	0.150	0.095	0.215	0.001	0.001	0.206	0.079	0.075	0.083
2,4,7-trinitro-9-fluorenone	0.358	0.080	0.358	0.360	0.081	0.145	0.163	0.094	0.229	0.002	0.001	0.213	0.053	0.033	0.063
2,4,5,7-tetranitro-9-fluorenone	0.359	0.079	0.359	0.361	0.079	0.147	0.164	0.093	0.229	0.002	0.000	0.212	0.048	0.024	0.060
8-nitrofluoranthene	0.347	0.078	0.347	0.349	0.080	0.132	0.150	0.093	0.220	0.002	0.002	0.215	0.089	0.087	0.088
2,7-dinitro-9-fluorenone	0.357	0.083	0.357	0.361	0.084	0.146	0.163	0.098	0.228	0.004	0.001	0.211	0.058	0.042	0.066
2-nitropyrene	0.321	0.076	0.329	0.322	0.076	0.124	0.140	0.087	0.202	0.001	0.000	0.205	0.088	0.086	0.092
1-nitrofluoranthene	0.329	0.079	0.329	0.331	0.080	0.129	0.145	0.090	0.208	0.002	0.001	0.200	0.070	0.064	0.078
2-nitroanthracene	0.315	0.070	0.315	0.317	0.072	0.119	0.135	0.081	0.196	0.002	0.002	0.196	0.076	0.073	0.081
1,3,6,8-tetranitropyrene	0.314	0.069	0.314	0.313	0.069	0.116	0.132	0.080	0.192	-0.001	0.000	0.198	0.053	0.042	0.063
3-nitrofluoranthene	0.328	0.080	0.328	0.330	0.080	0.128	0.145	0.090	0.208	0.002	0.000	0.200	0.070	0.064	0.078
1,3,6-trinitropyrene	0.314	0.071	0.314	0.314	0.072	0.118	0.133	0.082	0.193	0.000	0.001	0.196	0.062	0.054	0.070
1,6-dinitropyrene	0.322	0.078	0.322	0.323	0.078	0.125	0.141	0.089	0.202	0.001	0.000	0.197	0.075	0.071	0.081
1,3-dinitropyrene	0.312	0.074	0.312	0.313	0.074	0.120	0.135	0.085	0.194	0.001	0.000	0.192	0.068	0.062	0.075
1,8-dinitropyrene	0.316	0.074	0.319	0.317	0.074	0.120	0.133	0.084	0.196	0.001	0.000	0.199	0.083	0.073	0.084

Table S2f. Same as **Table S2a** but for electrophilicity index descriptor (ω).

Name of Compound	ω_{HFx}	ω_{B88}	ω_{HF}	$\omega_{\text{HFx+LYP}}$	ω_{BLYP}	ω_{B3LYP}	ω_{M06}	$\omega_{\text{M06-L}}$	$\omega_{\text{M06-2X}}$	$\omega_{\text{CORR(HFx+LYP)}}$	$\omega_{\text{CORR(BLYP)}}$	$-\omega_{\text{CORR(B3LYP)}}$	$\omega_{\text{CORR(M06)}}$	$\omega_{\text{CORR(M06-L)}}$	$\omega_{\text{CORR(M06-2X)}}$
4-nitrotoluene	0.027	0.088	0.027	0.039	0.128	0.092	0.085	0.119	0.063	0.128	0.128	0.002	0.009	0.008	0.009
2,6-dinitrotoluene	0.035	0.109	0.035	0.051	0.160	0.112	0.103	0.150	0.078	0.613	0.578	0.002	0.008	0.006	0.009
3,4-dinitrotoluene	0.036	0.148	0.038	0.051	0.165	0.153	0.135	0.234	0.091	0.578	0.035	0.001	0.005	0.003	0.006
2,4-dinitrotoluene	0.039	0.116	0.039	0.053	0.163	0.116	0.108	0.153	0.082	0.136	0.136	0.002	0.008	0.005	0.009
2,3-dinitrotoluene	0.035	0.148	0.037	0.050	0.234	0.153	0.136	0.233	0.091	0.578	-0.225	0.001	0.005	0.003	0.006
6-nitroquinoline	0.030	0.104	0.030	0.044	0.153	0.107	0.099	0.144	0.073	0.182	-0.481	0.003	0.009	0.008	0.010
3-nitrocarbazole	0.019	0.078	0.019	0.030	0.120	0.082	0.075	0.112	0.053	0.120	0.240	0.003	0.010	0.010	0.010
3,5-dinitrotoluene	0.040	0.119	0.040	0.055	0.168	0.120	0.111	0.158	0.084	0.109	0.193	0.002	0.008	0.006	0.009
3-methyl-2-nitronaphthalene	0.023	0.092	0.023	0.037	0.140	0.095	0.087	0.133	0.062	-	0.512	0.003	0.010	0.009	0.010
5-nitroquinoline	0.031	0.103	0.031	0.045	0.157	0.108	0.100	0.147	0.074	0.272	-0.256	0.002	0.009	0.008	0.009
2,5-dinitrotoluene	0.043	0.133	0.043	0.059	0.193	0.133	0.123	0.180	0.091	0.193	0.613	0.002	0.008	0.006	0.009
1,3,6,8-tetranitronaphthalene	0.058	0.179	0.058	0.077	0.246	0.172	0.158	0.235	0.120	0.272	0.204	0.001	0.006	0.004	0.008
1,3-dinitrobenzene	0.042	0.125	0.042	0.058	0.174	0.122	0.114	0.163	0.088	0.091	0.136	0.002	0.008	0.005	0.009
2-nitrocarbazole	0.024	0.092	0.024	0.036	0.140	0.094	0.086	0.131	0.061	0.240	0.240	0.002	0.009	0.008	0.009
4-nitrocarbazole	0.023	0.093	0.022	0.036	0.144	0.093	0.087	0.135	0.061	0.171	0.481	0.003	0.009	0.009	0.010
2,3,4-trinitrotoluene	0.045	0.129	0.046	0.061	0.184	0.128	0.119	0.172	0.091	0.136	-	0.001	0.007	0.004	0.009
1-methyl-2-nitronaphthalene	0.022	0.090	0.022	0.035	0.136	0.092	0.084	0.127	0.060	0.545	0.240	0.003	0.010	0.009	0.010
2,4,6-trinitrotoluene	0.050	0.143	0.050	0.066	0.198	0.139	0.128	0.184	0.099	0.091	0.193	0.001	0.005	0.002	0.007
9-nitroanthracene	0.024	0.099	0.024	0.038	0.161	0.103	0.095	0.152	0.069	-	-0.545	0.003	0.009	0.009	0.010
1-nitronaphthlene	0.025	0.094	0.025	0.038	0.145	0.099	0.091	0.137	-0.020	-	0.545	0.003	0.010	0.009	-0.063
2-nitronaphthlene	0.026	0.097	0.026	0.039	0.144	0.100	0.091	0.137	0.066	0.171	0.240	0.003	0.009	0.009	0.010
2,3,5-trinitrotoluene	0.052	0.225	0.052	0.072	0.199	0.271	0.186	0.340	0.123	0.512	0.028	0.003	0.005	0.003	0.006
2,3,6-trinitrotoluene	0.052	0.208	0.051	0.070	0.222	0.201	0.176	0.313	0.117	0.613	0.040	0.000	-0.003	-0.005	-0.002
6-nitrobenzo[a]pyrene	0.024	0.100	0.024	0.038	0.164	0.105	0.097	0.154	0.070	-	-	0.003	0.009	0.009	0.010
1,3,5-trinitrobenzene	0.055	0.153	0.055	0.072	0.208	0.146	0.135	0.196	0.106	0.116	0.193	0.001	0.005	0.001	0.007
1,3-dinitronaphthlene	0.037	0.127	0.037	0.053	0.187	0.129	0.118	0.177	0.087	0.272	-	0.002	0.008	0.008	0.009
1,5-dinitronaphthlene	0.036	0.122	0.036	0.052	0.182	0.126	0.115	0.173	0.084	0.578	0.613	0.002	0.008	0.007	0.009
5-nitroacenaphthene	0.023	0.090	0.023	0.036	0.139	0.095	0.087	0.131	0.062	-	0.512	0.003	0.010	0.009	0.010
3,4,5-trinitrotoluene	0.046	0.302	0.045	0.062	0.193	0.145	0.128	0.188	0.097	0.578	0.009	0.002	0.011	0.009	0.014
2-nitrofluorene	0.025	0.092	0.025	0.038	0.138	0.096	0.088	0.130	0.064	0.128	0.240	0.003	0.009	0.009	0.010
1,8-dinitronaphthlene	0.034	0.106	0.034	0.048	0.154	0.111	0.102	0.150	0.078	0.128	0.182	0.002	0.008	0.007	0.009
2,4,5-trinitrotoluene	0.052	0.222	0.052	0.072	0.218	0.217	0.188	0.358	0.123	-	0.036	0.001	0.005	0.003	0.006
2,7-dinitrofluorene	0.037	0.122	0.037	0.053	0.175	0.124	0.115	0.168	0.084	0.128	0.256	0.002	0.008	0.007	0.009
1-nitrobenzo[e]pyrene	0.023	0.095	0.023	0.036	0.149	0.097	0.089	0.141	0.065	0.256	0.512	0.003	0.010	0.010	0.011
2-nitrophenanthrene	0.025	0.097	0.025	0.038	0.145	0.099	0.091	0.138	0.065	0.256	0.240	0.003	0.010	0.009	0.010
7-nitrofluoranthene	0.026	0.096	0.026	0.039	0.145	0.100	0.092	0.139	0.068	0.171	0.240	0.003	0.009	0.009	0.010
1-nitropyrene	0.026	0.104	0.026	0.040	0.161	0.108	0.099	0.154	0.071	0.256	0.512	0.003	0.010	0.009	0.010
6-nitrochrysene	0.024	0.097	0.024	0.037	0.151	0.100	0.092	0.142	0.066	0.545	0.545	0.003	0.010	0.009	0.010
2,4,7-trinitro-9-fluorenone	0.058	0.194	0.058	0.077	0.271	0.177	0.161	0.255	0.120	0.272	0.578	0.001	0.007	0.004	0.008
2,4,5,7-tetranitro-9-fluorenone	0.067	0.223	0.067	0.089	0.311	0.197	0.181	0.289	0.135	0.289	-	0.001	0.006	0.003	0.007
8-nitrofluoranthene	0.019	0.095	0.019	0.030	0.146	0.093	0.084	0.135	0.057	0.240	0.240	0.004	0.011	0.011	0.011
2,7-dinitro-9-fluorenone	0.047	0.161	0.048	0.065	0.229	0.152	0.139	0.217	0.104	0.136	0.545	0.002	0.007	0.005	0.008
2-nitropyrene	0.024	0.101	0.022	0.037	0.160	0.105	0.095	0.151	0.068	0.512	-	0.004	0.011	0.011	0.012
1-nitrofluoranthene	0.030	0.113	0.030	0.045	0.170	0.115	0.106	0.165	0.078	0.256	0.512	0.003	0.009	0.008	0.010
2-nitroanthracene	0.026	0.113	0.026	0.040	0.171	0.111	0.102	0.165	0.072	0.240	0.240	0.003	0.010	0.009	0.010
1,3,6,8-tetranitropyrene	0.061	0.213	0.061	0.084	0.312	0.207	0.185	0.294	0.134	-0.613	-	0.001	0.007	0.005	0.008
3-nitrofluoranthene	0.031	0.114	0.031	0.046	0.174	0.120	0.109	0.170	0.080	0.256	-	0.003	0.009	0.008	0.010
1,3,6-trinitropyrene	0.048	0.177	0.048	0.069	0.259	0.173	0.157	0.247	0.113	-	0.613	0.002	0.008	0.007	0.009
1,6-dinitropyrene	0.032	0.126	0.032	0.048	0.192	0.129	0.117	0.182	0.084	0.545	-	0.003	0.009	0.009	0.010
1,3-dinitropyrene	0.037	0.138	0.037	0.054	0.209	0.141	0.128	0.198	0.092	0.545	-	0.003	0.009	0.008	0.009
1,8-dinitropyrene	0.037	0.138	0.034	0.054	0.212	0.141	0.134	0.202	0.092	0.545	-	0.003	0.010	0.009	0.011

Table S1. Compound is the name or CT, reduction or pH, and excluded or in different methods, excluded compounds are other names or structural codes (retrieved from William et al)

Name of the compound	Activity (log TA100)	Exchange (BPX, BSK, BP) only method		Exchange-Correlation (BPX, LYP, BLYP, BLYP) method		Exchange-Correlation (Min Functional, MM, MM6-L, MM6-2X) method		Electron-correlation (CORR:BPX-LYP, CORR:LYP, CORR:BSLYP) method		Electron-correlation (CORR:MM6, CORR:MM6-L, CORR:MM6-2X) method	
		Activity Sampling (Order splitting)	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	Activity Sampling (Order splitting)	30% random splitting
4-nitrobenzene	-2.10	T	T	T	T	T	T	T	T	T	T
2,6-dinitrotoluene	-1.34	P	T	P	T	P	T	P	T	P	T
3,4-dinitrotoluene	-1.30	P	T	P	T	P	T	P	T	P	T
2,4-dinitrotoluene	-1.29	P	T	P	T	P	T	P	T	P	T
2,3-dinitrotoluene	-1.26	T	P	T	P	T	P	T	P	T	P
6-nitroquinoline	-1.05	P	T	P	T	P	T	P	T	P	T
3-nitrocarbazole	-1.00	T	T	E	T	E	T	E	T	E	T
3,5-dinitrotoluene	-0.72	P	T	P	T	P	T	P	T	P	T
3-methyl-2-nitroanaphthalene	-0.70	T	P	T	P	T	P	T	P	T	P
5-nitroquinoline	-0.70	P	T	P	T	P	T	P	T	P	T
2,5-dinitrotoluene	-0.63	T	T	E	T	E	T	E	T	E	T
1,3,6,8-tetraazanonaphthalene	-0.52	E	E	E	E	E	E	E	E	E	E
1,3-dinitrobenzene	-0.51	P	T	P	T	P	T	P	T	P	T
2-nitrocarbazole	-0.30	T	T	T	T	T	T	T	T	T	T
4-nitrocarbazole	-0.30	P	T	P	T	P	T	P	T	P	T
2,3,4-trinitrotoluene	0.08	T	T	P	T	P	T	P	T	P	T
1-methyl-2-nitroanaphthalene	0.08	T	T	P	T	P	T	P	T	P	T
2,4,6-trinitrotoluene	0.16	T	T	P	T	P	T	P	T	P	T
9-nitroanthracene	0.26	P	T	P	T	P	T	P	T	P	T
1-nitroanaphthalene	0.28	T	P	T	P	T	P	T	P	T	P
2-nitroanaphthalene	0.37	P	T	P	T	P	T	P	T	P	T
2,3,5-trinitrotoluene	0.46	T	T	E	T	E	T	E	T	E	T
2,3,6-trinitrotoluene	0.55	P	T	P	T	P	T	P	T	P	T
6-nitrobenzopyrene	0.71	E	E	E	E	E	E	E	E	E	E
1,3,5-trinitrobenzene	0.72	T	T	T	T	T	T	T	T	T	T
1,5-dinitroanaphthalene	0.86	P	T	P	T	P	T	P	T	P	T
1,5-dinitroanthracene	0.91	T	T	T	T	T	T	T	T	T	T
5-nitroacenaphthene	0.97	P	T	P	T	P	T	P	T	P	T
3,4,5-trinitrotoluene	1.01	T	T	T	T	T	T	T	T	T	T
2-nitrofluorene	1.08	P	T	P	T	P	T	P	T	P	T
1,8-dinitroanaphthalene	1.12	T	T	T	T	T	T	T	T	T	T
2,4,5-trinitrotoluene	1.12	P	P	P	P	P	P	P	P	P	P
2,7-dinitrofluorene	1.27	T	T	T	T	T	T	T	T	T	T
1-nitrobenzopyrene	1.65	E	E	E	E	E	E	E	E	E	E
2-nitrophenanthrene	1.79	P	T	P	T	P	T	P	T	P	T
7-nitrofluoranthene	2.09	T	T	T	T	T	T	T	T	T	T
1-nitropyrene	2.17	P	T	P	T	P	T	P	T	P	T
6-nitrochrysenes	2.21	T	T	T	T	T	T	T	T	T	T
2,4,7-trinitro-9-fluorenone	2.27	E	E	E	E	E	E	E	E	E	E
2,4,5,7-tetraazano-9-fluorenone	2.46	E	E	E	E	E	E	E	E	E	E
8-nitrofluoranthene	2.60	P	T	P	T	P	T	P	T	P	T
2,7-dinitro-9-fluorenone	2.69	T	T	T	T	T	T	T	T	T	T
2-nitropyrene	2.87	P	T	P	T	P	T	P	T	P	T
1-nitrofluoranthene	3.00	T	T	T	T	T	T	T	T	T	T
2-nitroanthracene	3.05	P	T	P	T	P	T	P	T	P	T
1,3,6,8-tetraazapyrene	3.18	E	E	E	E	E	E	E	E	E	E
3-nitrofluoranthene	3.31	T	T	T	T	T	T	T	T	T	T
1,6-dinitropyrene	3.87	P	T	P	T	P	T	P	T	P	T
1,6-dinitropyrene	4.09	T	T	T	T	T	T	T	T	T	T
1,3-dinitropyrene	4.63	P	T	P	T	P	T	P	T	P	T
1,8-dinitropyrene	4.74	T	T	T	T	T	T	T	T	T	T
Number of Compounds in Training Set	23	30	23	30	23	30	23	30	23	30	23
Number of Compounds in Prediction Set	21	35	22	35	22	35	22	35	22	35	22
Number of Compounds in External Set	7	6	6	6	6	6	6	6	6	6	6

Name of the compound	Activity (log TA100)	Electron-correlation (CORR:BPX-LYP) method		Name of the compound	Activity (log TA100)	Electron-correlation (CORR:LYP) method	
		Activity Sampling (Order splitting)	30% random splitting			Activity Sampling (Order splitting)	30% random splitting
4-nitrobenzene	-2.10	T	T	4-nitrobenzene	-2.10	T	T
2,6-dinitrotoluene	-1.34	P	T	2,6-dinitrotoluene	-1.34	P	T
3,4-dinitrotoluene	-1.30	T	T	3,4-dinitrotoluene	-1.30	T	T
2,4-dinitrotoluene	-1.29	P	T	2,4-dinitrotoluene	-1.29	P	T
2,3-dinitrotoluene	-1.26	T	P	2,3-dinitrotoluene	-1.26	T	P
6-nitroquinoline	-1.05	P	T	6-nitroquinoline	-1.05	P	T
3-nitrocarbazole	-1.00	T	T	3-nitrocarbazole	-1.00	T	T
3,5-dinitrotoluene	-0.72	P	T	3,5-dinitrotoluene	-0.72	P	T
5-nitroquinoline	-0.70	T	P	3-methyl-2-nitroanaphthalene	-0.70	P	T
2,5-dinitrotoluene	-0.63	T	T	5-nitroquinoline	-0.70	P	T
1,3,6,8-tetraazanonaphthalene	-0.52	E	E	2,5-dinitrotoluene	-0.63	T	E
1,3-dinitrobenzene	-0.51	T	T	1,3,6,8-tetraazanonaphthalene	-0.52	E	E
2-nitrocarbazole	-0.30	P	T	1,3-dinitrobenzene	-0.51	P	P
4-nitrocarbazole	-0.30	T	T	2-nitrocarbazole	-0.30	T	T
2,3,4-trinitrotoluene	0.08	T	T	4-nitrocarbazole	-0.30	T	T
1-methyl-2-nitroanaphthalene	0.08	T	T	1-methyl-2-nitroanaphthalene	0.08	T	T
2,4,6-trinitrotoluene	0.16	P	P	2,4,6-trinitrotoluene	0.16	P	P
2-nitroanaphthalene	0.37	T	T	9-nitroanthracene	0.26	E	E
2,3,5-trinitrotoluene	0.46	P	T	1-nitroanaphthalene	0.28	T	T
2,3,6-trinitrotoluene	0.55	T	T	2-nitroanaphthalene	0.37	P	T
1,3,5-trinitrobenzene	0.72	P	P	2,3,5-trinitrotoluene	0.46	P	T
1,5-dinitroanaphthalene	0.86	T	T	2,3,6-trinitrotoluene	0.55	T	T
1,5-dinitroanthracene	0.91	P	T	1,3,5-trinitrobenzene	0.72	T	T
2-nitrofluorene	1.08	P	T	1,5-dinitroanaphthalene	0.86	P	T
1,8-dinitroanaphthalene	1.12	T	T	1,5-dinitroanthracene	0.91	P	T
2-nitrofluorene	1.12	P	P	5-nitroacenaphthene	0.97	P	P
2,7-dinitrofluorene	1.27	T	T	1,8-dinitroanaphthalene	1.12	T	T
1-nitrobenzopyrene	1.65	E	E	2,7-dinitrofluorene	1.27	T	T
2-nitrophenanthrene	1.79	P	T	1-nitrobenzopyrene	1.65	E	E
7-nitrofluoranthene	2.09	T	T	2-nitrophenanthrene	1.79	P	T
1-nitropyrene	2.17	P	T	7-nitrofluoranthene	2.09	T	T
6-nitrochrysenes	2.21	T	T	1-nitropyrene	2.17	P	T
2,4,7-trinitro-9-fluorenone	2.27	P	P	6-nitrochrysenes	2.21	T	T
2,4,5,7-tetraazano-9-fluorenone	2.46	E	E	2,4,7-trinitro-9-fluorenone	2.27	P	P
8-nitrofluoranthene	2.60	T	T	2,4,5,7-tetraazano-9-fluorenone	2.46	E	E
2,7-dinitro-9-fluorenone	2.69	P	T	8-nitrofluoranthene	2.60	T	T
2-nitropyrene	2.87	T	T	2,7-dinitro-9-fluorenone	2.69	P	T
1-nitrofluoranthene	3.00	T	T	2-nitropyrene	2.87	T	T
2-nitroanthracene	3.05	P	T	1-nitrofluoranthene	3.00	T	T
1,3,6,8-tetraazapyrene	3.18	E	E	2-nitroanthracene	3.05	P	T
3-nitrofluoranthene	3.31	T	T	1,3,6,8-tetraazapyrene	3.18	E	E
1,6-dinitropyrene	3.87	P	P	3-nitrofluoranthene	3.31	T	T
1,6-dinitropyrene	4.09	T	T	1,6-dinitropyrene	3.87	P	P
1,3-dinitropyrene	4.63	P	P	1,3-dinitropyrene	4.09	T	T
1,8-dinitropyrene	4.74	T	T	1,8-dinitropyrene	4.63	P	P
Number of Compounds in Training Set	23	30	23	Number of Compounds in Training Set	20	26	20
Number of Compounds in Prediction Set	21	35	22	Number of Compounds in Prediction Set	18	11	11
Number of Compounds in External Set	7	6	6	Number of Compounds in External Set	3	4	4

*Splitting is performed through QSARINS software. Ref:[54]. N. Chiaro, E. Papa, S. Kovach, S. Casati, P. Gramatica, QSAR Res. in Environ Chem and Ecosyst University of Insubria, Varese, Italy.

SDAS set excluding the compounds with undefined electrophilicity value (v) for CORR:BPX-LYP method (refer to Supporting Information Table S2).

SDAS set excluding the compounds with undefined electrophilicity value (v) for CORR:LYP (refer to Supporting Information Table S2).

Table S4. Compounds in the training set (T), prediction set (P) and excluded set (E) in different types of splitting methods employed^a for TA98 mutagenicity. Excluded compounds are either response or structural outliers (determined from Williams plot).

Name of the compound	Activity (log TA98)	Exchange-only (BPX, BM, HF) methods			Exchange-Correlation (BPX-LYP, BLXP, BLXP) methods			Exchange-Correlation (Mm, functional, M6, M6-L, M6-2X) methods			Electron-correlation (C0R0/EPX-LYP, C0R0/BLXP) methods			Electron-correlation (C0R0/M6, C0R0/M6-L, C0R0/M6-2X) methods		
		Activity Sampling (Order splitting)	30% random splitting	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	30% random splitting	Activity Sampling (Order splitting)	30% random splitting	30% random splitting
1-methyl-2-nitroazaphthalene	-0.70	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
1-nitroazaphthalene	-0.61	T	P	P	P	P	E	E	E	P	E	E	E	E	E	E
2-nitroazaphthalene	-0.30	P	T	T	T	T	P	P	T	P	T	P	T	P	T	P
1,3-dinitroazaphthalene	-0.05	T	T	T	T	T	P	P	T	P	E	T	E	E	E	E
1,3-dinitrobenzene	0.03	E	E	E	E	E	P	P	T	P	T	P	T	P	T	P
1,5-dinitroazaphthalene	0.52	T	P	P	P	P	P	P	T	T	T	T	T	T	T	T
1,8-dinitroazaphthalene	0.90	P	T	T	T	T	T	T	T	E	T	T	T	T	T	T
2-nitrofluorene	1.43	P	P	T	T	T	P	P	T	P	T	P	T	P	T	P
2-nitrophenanthrene	2.11	T	P	P	P	T	T	T	T	T	T	T	T	T	T	T
1-nitrofluoranthene	2.74	T	T	T	T	T	P	P	P	P	T	P	P	P	P	P
1-nitropyrene	2.78	P	T	T	T	T	T	T	T	T	P	T	T	T	T	T
2-nitroanthracene	2.95	P	P	P	P	P	E	E	P	T	P	T	P	T	P	T
2-nitropyrene	3.35	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
1,3,6,8-tetraazapyrene	4.99	E	E	E	E	E	E	E	E	E	E	E	E	E	E	E
1,3,6-dinitropyrene	4.99	T	E	E	E	E	E	E	E	E	E	E	E	E	E	E
1,5-dinitropyrene	5.04	P	T	T	T	T	P	P	P	P	P	P	T	P	T	P
1,6-dinitropyrene	5.06	T	P	P	P	P	P	T	T	T	T	P	T	P	T	T
1,8-dinitropyrene	5.39	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
Number of Compounds in Training Set		10	10	10	11	8	11	8	10	8	10	8	10	8	10	8
Number of Compounds in Prediction Set		5	5	4	4	6	4	5	4	5	4	5	4	5	4	4
Number of Compounds in External Set		2	3	3	3	4	3	3	2	5	5	4	5	4	4	4

* Name of the compound	Activity (log TA98)	Electron-correlation (C0R0/EPX-LYP) methods		* Name of the compound	Activity (log TA98)	Electron-correlation (C0R0/BLXP) methods	
		Activity Sampling (Order splitting)	30% random splitting			Activity Sampling (Order splitting)	30% random splitting
1-methyl-2-nitroazaphthalene	-0.70	T	T	1-methyl-2-nitroazaphthalene	-0.70	T	T
1-nitroazaphthalene	-0.61	T	T	1-nitroazaphthalene	-0.61	T	T
2-nitroazaphthalene	-0.30	P	T	2-nitroazaphthalene	-0.30	P	T
1,3-dinitroazaphthalene	0.03	T	P	1,3-dinitroazaphthalene	0.03	T	P
1,5-dinitroazaphthalene	0.52	P	T	1,5-dinitroazaphthalene	0.52	P	T
1,8-dinitroazaphthalene	0.90	T	T	1,8-dinitroazaphthalene	0.90	T	T
2-nitrofluorene	1.43	P	T	2-nitrofluorene	1.43	P	T
2-nitrophenanthrene	2.11	T	P	2-nitrophenanthrene	2.11	T	P
1-nitrofluoranthene	2.74	P	T	1-nitrofluoranthene	2.74	P	T
1-nitropyrene	2.78	T	T	1-nitropyrene	2.78	T	T
2-nitroanthracene	2.95	P	T	2-nitroanthracene	2.95	P	T
2-nitropyrene	3.35	T	P	1,3,6-dinitropyrene	4.99	T	T
1,3,6,8-tetraazapyrene	4.99	E	E	Number of Compounds in Training Set		7	8
1,3-dinitropyrene	5.04	T	T	Number of Compounds in Prediction Set		5	4
1,5-dinitropyrene	5.06	P	P	Number of Compounds in External Set		0	0
1,8-dinitropyrene	5.39	T	T				
Number of Compounds in Training Set		9	11				
Number of Compounds in Prediction Set		6	4				
Number of Compounds in External Set		1	1				

^a Splitting performed through QM500 software (Ref [24]). N. Dethlefs, E. Paik, S. Kowalski, & G. S. G. F. Dethlefs, QM500, User's Manual, University of Wrocław, Wrocław, Poland, 2012 (<http://www.qm500.pl>). Data set including the compounds with undefined descriptibility value (-) for C0R0/EPX-LYP method (refer to Supporting Information Table S2). Data set excluding the compounds with undefined descriptibility value (-) for C0R0/BLXP method (refer to Supporting Information Table S2).

Table S5. Comparison of the key internal and external validation parameters for the models based on the total energy (E) and energy of HOMO (E_{HOMO}) computed with the Exchange (X) only, Exchange + Correlation (X+C) methods, and with the effect of electron-correlation (CORR) of the descriptors, for modeling TA98 mutagenicity of nitrated-PAHs.

Model S.No.	Method	Descriptor Employed	Splitting Employed	R^2	Q^2_{LOO}	$R^2 - Q^2_{\text{LOO}}$	Q^2_{LMO}	Q^2_{F3}	CCC_{EXT}
Exchange (X) Only									
1.	DFT/B88	$E_{\text{B88}}, E_{\text{B88}}^{\text{HOMO}}$	activity sampling	0.948	0.862	0.085	0.681	0.943	0.955
2.	DFT/HFX	$E_{\text{HFX}}, E_{\text{HFX}}^{\text{HOMO}}$	activity sampling	0.953	0.876	0.077	0.570	0.937	0.951
Exchange + Correlation (X+C)									
3.	HFX + LYP	$E_{\text{HFX+LYP}}, E_{\text{HFX+LYP}}^{\text{HOMO}}$	activity sampling	0.941	0.885	0.056	0.699	0.984	0.994
4.	BLYP	$E_{\text{BLYP}}, E_{\text{BLYP}}^{\text{HOMO}}$	activity sampling	0.938	0.874	0.064	0.721	0.986	0.994
5.	B3LYP	$E_{\text{B3LYP}}, E_{\text{B3LYP}}^{\text{HOMO}}$	activity sampling	0.940	0.882	0.058	0.743	0.987	0.995
6.	M06	$E_{\text{M06}}, E_{\text{M06}}^{\text{HOMO}}$	activity sampling	0.963	0.912	0.051	0.244	0.910	0.922
7.	M06-L	$E_{\text{M06-L}}, E_{\text{M06-L}}^{\text{HOMO}}$	activity sampling	0.960	0.901	0.059	0.333	0.907	0.919
8.	M06-2X	$E_{\text{M06-2X}}, E_{\text{M06-2X}}^{\text{HOMO}}$	activity sampling	0.964	0.915	0.049	0.295	0.904	0.915
Electron-Correlation (CORR) only									
9.	CORR(HFX+LYP)	$E_{\text{CORR(HFX+LYP)}}, E_{\text{CORR(HFX+LYP)}}^{\text{HOMO}}$	activity sampling	0.973	0.926	0.047	0.327	0.919	0.957
10.	CORR(BLYP)	$E_{\text{CORR(BLYP)}}, E_{\text{CORR(BLYP)}}^{\text{HOMO}}$	activity sampling	0.968	0.899	0.069	0.244	0.892	0.943
11.	CORR(B3LYP)	$E_{\text{CORR(B3LYP)}}, E_{\text{CORR(B3LYP)}}^{\text{HOMO}}$	activity sampling	0.975	0.942	0.033	0.486	0.950	0.971
12.	CORR(M06)	$E_{\text{CORR(M06)}}, E_{\text{CORR(M06)}}^{\text{HOMO}}$	activity sampling	0.978	0.956	0.023	0.604	0.947	0.948
13.	CORR(M06-L)	$E_{\text{CORR(M06-L)}}, E_{\text{CORR(M06-L)}}^{\text{HOMO}}$	activity sampling	0.978	0.954	0.024	0.560	0.950	0.952
14.	CORR(M06-2X)	$E_{\text{CORR(M06-2X)}}, E_{\text{CORR(M06-2X)}}^{\text{HOMO}}$	activity sampling	0.981	0.961	0.020	0.670	0.946	0.951

Table S6. Same as **Table S5** but for the models based on the total electronic energy (E) and energy of the LUMO (E_{LUMO}).

Model S.No.	Method	Descriptor Employed	Splitting Employed	R^2	Q^2_{LOO}	$R^2-Q^2_{\text{LOO}}$	Q^2_{LMO}	Q^2_{F3}	CCC_{EXT}
Exchange (X) Only									
1.	DFT/B88	$E_{\text{B88}}, E_{\text{B88}}^{\text{LUMO}}$	activity sampling	0.941	0.861	0.080	0.694	0.858	0.878
2.	DFT/HFX	$E_{\text{HFX}}, E_{\text{HFX}}^{\text{LUMO}}$	activity sampling	0.860	0.660	0.200	0.120	0.733	0.762
Exchange + Correlation (X+C)									
3.	HFX + LYP	$E_{\text{HFX+LYP}}, E_{\text{HFX+LYP}}^{\text{LUMO}}$	activity sampling	0.886	0.711	0.175	0.542	0.588	0.872
4.	BLYP	$E_{\text{BLYP}}, E_{\text{BLYP}}^{\text{LUMO}}$	activity sampling	0.927	0.823	0.104	-	0.836	0.925
5.	B3LYP	$E_{\text{B3LYP}}, E_{\text{B3LYP}}^{\text{LUMO}}$	activity sampling	0.918	0.786	0.132	0.563	0.813	0.915
6.	M06	$E_{\text{M06}}, E_{\text{M06}}^{\text{LUMO}}$	activity sampling	0.793	0.556	0.237	0.083	0.912	0.959
7.	M06-L	$E_{\text{M06-L}}, E_{\text{M06-L}}^{\text{LUMO}}$	activity sampling	0.836	0.679	0.157	0.339	0.905	0.958
8.	M06-2X	$E_{\text{M06-2X}}, E_{\text{M06-2X}}^{\text{LUMO}}$	activity sampling	0.926	0.607	0.319	-0.951	-	-
Electron-Correlation (CORR) only									
9.	CORR(HFX+LYP)	$E_{\text{CORR(HFX+LYP)}}, E_{\text{CORR(HFX+LYP)}}^{\text{LUMO}}$	activity sampling	0.992	0.986	0.006	0.667	0.839	0.924
10.	CORR(BLYP)	$E_{\text{CORR(BLYP)}}, E_{\text{CORR(BLYP)}}^{\text{LUMO}}$	activity sampling	0.910	0.837	0.073	0.606	0.864	0.945
11.	CORR(B3LYP)	$E_{\text{CORR(B3LYP)}}, E_{\text{CORR(B3LYP)}}^{\text{LUMO}}$	activity sampling	0.929	0.851	0.078	-0.237	-	-
12.	CORR(M06)	$E_{\text{CORR(M06)}}, E_{\text{CORR(M06)}}^{\text{LUMO}}$	activity sampling	0.945	0.881	0.065	-0.133	-	-
13.	CORR(M06-L)	$E_{\text{CORR(M06-L)}}, E_{\text{CORR(M06-L)}}^{\text{LUMO}}$	activity sampling	0.948	0.873	0.074	0.136	0.822	0.831
14.	CORR(M06-2X)	$E_{\text{CORR(M06-2X)}}, E_{\text{CORR(M06-2X)}}^{\text{LUMO}}$	activity sampling	0.951	0.882	0.068	-0.032	-	-

* (-) indicates unstable models, with negative value of corresponding validation parameter.

Table S7. Same as **Table S5** but for the models based on the total electronic energy (E) and absolute electronegativity (χ).

Model S.No.	Method	Descriptor Employed	Splitting Employed	R^2	Q^2_{LOO}	$R^2 - Q^2_{\text{LOO}}$	Q^2_{LMO}	Q^2_{F3}	CCC_{EXT}
Exchange (X) Only									
1.	DFT/B88	$E_{\text{B88}}, \chi_{\text{B88}}$	activity sampling	0.951	0.873	0.078	0.557	0.945	0.954
2.	DFT/HFX	$E_{\text{HFX}}, \chi_{\text{HFX}}$	activity sampling	0.938	0.849	0.089	0.607	0.940	0.950
Exchange + Correlation (X+C)									
3.	HFX + LYP	$E_{\text{HFX+LYP}}, \chi_{\text{HFX+LYP}}$	activity sampling	0.944	0.882	0.062	0.778	0.928	0.974
4.	BLYP	$E_{\text{BLYP}}, \chi_{\text{BLYP}}$	activity sampling	0.945	0.883	0.062	0.715	0.952	0.982
5.	B3LYP	$E_{\text{B3LYP}}, \chi_{\text{B3LYP}}$	activity sampling	0.918	0.786	0.132	0.563	0.813	0.915
6.	M06	$E_{\text{M06}}, \chi_{\text{M06}}$	activity sampling	0.951	0.865	0.086	-	-	-
7.	M06-L	$E_{\text{M06-L}}, \chi_{\text{M06-L}}$	activity sampling	0.913	0.803	0.110	0.425	0.932	0.971
8.	M06-2X	$E_{\text{M06-2x}}, \chi_{\text{M06-2X}}$	activity sampling	0.909	0.791	0.119	0.454	0.945	0.976
Electron-Correlation (CORR) only									
9.	CORR(HFX+LYP)	$E_{\text{CORR(HFX+LYP)}}, \chi_{\text{CORR(HFX+LYP)}}$	activity sampling	0.992	0.986	0.006	0.667	0.839	0.924
10.	CORR(BLYP)	$E_{\text{CORR(BLYP)}}, \chi_{\text{CORR(BLYP)}}$	activity sampling	0.970	0.902	0.068	0.257	0.897	0.948
11.	CORR(B3LYP)	$E_{\text{CORR(B3LYP)}}, \chi_{\text{CORR(B3LYP)}}$	activity sampling	0.960	0.892	0.068	0.079	0.923	0.951
12.	CORR(M06)	$E_{\text{CORR(M06)}}, \chi_{\text{CORR(M06)}}$	activity sampling	0.967	0.923	0.045	0.274	0.878	0.883
13.	CORR(M06-L)	$E_{\text{CORR(M06-L)}}, \chi_{\text{CORR(M06-L)}}$	activity sampling	0.971	0.934	0.037	0.344	0.884	0.891
14.	CORR(M06-2X)	$E_{\text{CORR(M06-2X)}}, \chi_{\text{CORR(M06-2X)}}$	activity sampling	0.969	0.931	0.038	0.442	0.889	0.897

* (-) indicates unstable models, with negative value of corresponding validation parameter.

Table S8. Same as **Table S5** but for the models based on the total energy (E) and chemical hardness (η).

Model S.No.	Method	Descriptor Employed	Splitting Employed	R^2	Q^2_{LOO}	$R^2-Q^2_{\text{LOO}}$	Q^2_{LMO}	Q^2_{F3}	CCC_{EXT}
Exchange (X) Only									
1.	DFT/B88	$E_{\text{B88}}, \eta_{\text{B88}}$	activity sampling	0.909	0.786	0.124	0.491	0.746	0.847
2.	DFT/HFX	$E_{\text{HFX}}, \eta_{\text{HFX}}$	activity sampling	0.941	0.865	0.076	0.655	0.867	0.893
Exchange + Correlation (X+C)									
3.	HFX + LYP	$E_{\text{HFX+LYP}}, \eta_{\text{HFX+LYP}}$	activity sampling	0.908	0.850	0.058	0.669	0.972	0.988
4.	BLYP	$E_{\text{BLYP}}, \eta_{\text{BLYP}}$	activity sampling	0.884	0.825	0.059	0.451	0.933	0.969
5.	B3LYP	$E_{\text{B3LYP}}, \eta_{\text{B3LYP}}$	activity sampling	0.895	0.838	0.057	0.476	0.952	0.979
6.	M06	$E_{\text{M06}}, \eta_{\text{M06}}$	activity sampling	0.856	0.698	0.158	0.464	0.962	0.980
7.	M06-L	$E_{\text{M06-L}}, \eta_{\text{M06-L}}$	activity sampling	0.836	0.655	0.182	0.374	0.935	0.965
8.	M06-2X	$E_{\text{M06-2X}}, \eta_{\text{M06-2X}}$	activity sampling	0.860	0.705	0.155	0.355	0.975	0.988
Electron-Correlation (CORR) only									
9.	CORR(HFX+LYP)	$E_{\text{CORR(HFX+LYP)}}, \eta_{\text{CORR(HFX+LYP)}}$	activity sampling	0.904	0.842	0.062	0.542	0.807	0.922
10.	CORR(BLYP)	$E_{\text{CORR(BLYP)}}, \eta_{\text{CORR(BLYP)}}$	activity sampling	0.945	0.842	0.103	-0.177	-	-
11.	CORR(B3LYP)	$E_{\text{CORR(B3LYP)}}, \eta_{\text{CORR(B3LYP)}}$	activity sampling	0.928	0.842	0.086	0.690	0.766	0.908
12.	CORR(M06)	$E_{\text{CORR(M06)}}, \eta_{\text{CORR(M06)}}$	activity sampling	0.968	0.926	0.043	0.408	0.874	0.880
13.	CORR(M06-L)	$E_{\text{CORR(M06-L)}}, \eta_{\text{CORR(M06-L)}}$	activity sampling	0.972	0.936	0.036	0.374	0.888	0.896
14.	CORR(M06-2X)	$E_{\text{CORR(M06-2X)}}, \eta_{\text{CORR(M06-2X)}}$	activity sampling	0.969	0.929	0.040	0.367	0.887	0.894

* (-) indicates unstable models, with negative value of corresponding validation parameter.

Table S9. Same as **Table S5** but for the models based on the total electronic energy (E) and electrophilicity index (ω).

Model S.No.	Method	Descriptor Employed	Splitting Employed	R^2	Q^2_{LOO}	$R^2-Q^2_{\text{LOO}}$	Q^2_{LMO}	Q^2_{F3}	CCC_{EXT}
Exchange (X) Only									
1.	DFT/B88	$E_{\text{B88}}, \omega_{\text{B88}}$	activity sampling	0.947	0.894	0.053	0.649	0.673	0.744
2.	DFT/HFX	$E_{\text{HFX}}, \omega_{\text{HFX}}$	activity sampling	0.931	0.847	0.084	0.460	0.899	0.914
Exchange + Correlation (X+C)									
3.	HFX + LYP	$E_{\text{HFX+LYP}}, \omega_{\text{HFX+LYP}}$	activity sampling	0.932	0.842	0.090	0.682	0.810	0.937
4.	BLYP	$E_{\text{BLYP}}, \omega_{\text{BLYP}}$	activity sampling	0.854	0.633	0.221	0.359	0.565	0.857
5.	B3LYP	$E_{\text{B3LYP}}, \omega_{\text{B3LYP}}$	activity sampling	0.892	0.699	0.193	0.453	0.748	0.888
6.	M06	$E_{\text{M06}}, \omega_{\text{M06}}$	activity sampling	0.915	0.715	0.200	-	-	-
7.	M06-L	$E_{\text{M06-L}}, \omega_{\text{M06-L}}$	activity sampling	0.908	0.696	0.212	-	-	-
8.	M06-2X	$E_{\text{M06-2X}}, \omega_{\text{M06-2X}}$	activity sampling	0.935	0.723	0.211	-	-	-
Electron-Correlation (CORR) only									
9.	CORR(HFX+LYP)	$E_{\text{CORR(HFX+LYP)}}, \omega_{\text{CORR(HFX+LYP)}}$	activity sampling	0.948	0.761	0.187	-	-	-
10.	CORR(BLYP)	$E_{\text{CORR(BLYP)}}, \omega_{\text{CORR(BLYP)}}$	activity sampling	0.916	0.735	0.181	-	-	-
11.	CORR(B3LYP)	$E_{\text{CORR(B3LYP)}}, \omega_{\text{CORR(B3LYP)}}$	activity sampling	0.972	0.924	0.049	0.158	0.962	0.978
12.	CORR(M06)	$E_{\text{CORR(M06)}}, \omega_{\text{CORR(M06)}}$	activity sampling	0.964	0.903	0.061	0.363	0.898	0.902
13.	CORR(M06-L)	$E_{\text{CORR(M06-L)}}, \omega_{\text{CORR(M06-L)}}$	activity sampling	0.976	0.944	0.032	0.479	0.900	0.909
14.	CORR(M06-2X)	$E_{\text{CORR(M06-2X)}}, \omega_{\text{CORR(M06-2X)}}$	activity sampling	0.970	0.932	0.038	0.380	0.881	0.886

* (-) indicates unstable models, with negative value of corresponding validation parameter.

Table S10. Comparison of the key internal and external validation parameters for the models developed with the total energy (E), energy of HOMO (E_{HOMO}), energy of LUMO (E_{LUMO}), absolute electronegativity (χ), chemical hardness (η) and electrophilicity index (ω) computed with the Hartree-Fock (HF) method, for modeling TA100 mutagenicity and TA98 mutagenicity of nitrated-PAHs.

Model S.No.	Descriptor Employed	Splitting Employed	R^2	Q^2_{LOO}	$R^2-Q^2_{\text{LOO}}$	Q^2_{LMO}	Q^2_{F3}	CCC_{EXT}
TA100 mutagenicity								
1.	$E_{\text{HF}}, E_{\text{HF}}^{\text{HOMO}}$	30%	0.743	0.668	0.075	0.643	0.707	0.841
2.	$E_{\text{HF}}, E_{\text{HF}}^{\text{LUMO}}$	30%	0.680	0.603	0.077	0.574	0.816	0.682
3.	$E_{\text{HF}}, \chi_{\text{HF}}$	30%	0.778	0.711	0.067	0.692	0.731	0.859
4.	$E_{\text{HF}}, \eta_{\text{HF}}$	30%	0.647	0.536	0.111	0.529	0.653	0.793
5.	$E_{\text{HF}}, \omega_{\text{HF}}$	30%	0.787	0.731	0.056	0.707	0.736	0.860
TA98 mutagenicity								
6.	$E_{\text{HF}}, E_{\text{HF}}^{\text{HOMO}}$	activity sampling	0.958	0.888	0.070	0.757	0.935	0.950
7.	$E_{\text{HF}}, E_{\text{HF}}^{\text{LUMO}}$	activity sampling	0.891	0.748	0.143	0.307	0.699	0.763
8.	$E_{\text{HF}}, \chi_{\text{HF}}$	activity sampling	0.945	0.863	0.081	0.635	0.933	0.943
9.	$E_{\text{HF}}, \eta_{\text{HF}}$	activity sampling	0.942	0.860	0.082	0.615	0.750	0.852
10.	$E_{\text{HF}}, \omega_{\text{HF}}$	activity sampling	0.949	0.889	0.060	0.748	0.882	0.898

Table S11. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange (X) only (B88, HFX and HF) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	$RMSE_{TR}$	$RMSE_{CV}$	$RMSE_{EXT}$	F	$Ext.O.K$	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC_{TR}	CCC_{CV}	CCC_{EXT}	\bar{F}_{int}^2	Δr_{int}^2	MAE_{TR}	MAE_{CV}	MAE_{EXT}	RSS_{TR}	$PRESS_{CV}$	$PRESS_{EXT}$	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Xrand}	Q^2_{Xrand}	R^2_{Yrand}	Q^2_{Yrand}
1	E_{HFX}, E_{HF}^{HOMO}	0.7118	0.0639	0.7533	0.0224	0.7757	0.4230	0.8376	0.9496	0.7738	34.5906	5	0.7561	0.7546	0.8086	0.8737	0.8392	0.8666	0.7010	0.0779	0.6783	0.7793	0.6274	16.1378	20.7390	12.5730	0.7744	0.6258	8.4794	-21.5508	9.3239	-20.3343	9.7616	-19.8615
2	E_{HFX}, η_{HFX}	0.6852	0.0867	0.7491	0.0228	0.7719	0.2362	0.8448	0.9924	0.8392	33.8353	4	0.7131	0.7113	0.7749	0.8713	0.8254	0.8402	0.6812	0.0915	0.6390	0.7456	0.6984	16.4159	22.6539	14.7905	0.7767	0.5808	9.3805	-20.9621	8.8044	-21.2567	9.8392	-19.7071
3	E_{HFX}, χ_{HFX}	0.6796	0.0755	0.7306	0.0245	0.7551	0.1936	0.8754	1.0012	0.8016	30.8308	5	0.7382	0.7366	0.7946	0.8605	0.8198	0.8559	0.6663	0.0777	0.7157	0.8243	0.6457	17.6238	23.0534	13.4949	0.7568	0.6168	8.9207	-20.7981	8.5189	-21.7048	8.8984	-20.8981
4	E_{HF}, θ_{HF}	0.6634	0.0777	0.7152	0.0259	0.7411	0.0325	0.9000	1.0263	0.8582	28.6217	1	0.6999	0.6981	0.7646	0.8513	0.8088	0.8298	0.6313	0.0831	0.7356	0.8448	0.6971	18.6319	24.2235	15.4670	0.7433	0.5645	8.6804	-21.4844	9.1632	-20.9202	9.2232	-20.6306
5	E_{HFX}, ϕ_{HFX}	0.6487	0.0808	0.7024	0.0271	0.7295	0.0172	0.9200	1.0484	0.8708	26.9649	1	0.6911	0.6892	0.7577	0.8436	0.7997	0.8255	0.6163	0.0829	0.7534	0.8647	0.7072	19.4670	25.2796	15.9226	0.7300	0.5577	8.7087	-21.5974	9.4434	-20.3285	9.5160	-20.0754
6	E_{HF}, χ_{HF}	0.6469	0.0820	0.7018	0.0271	0.7289	0.1386	0.9209	1.0510	0.8490	26.8888	3	0.7063	0.7045	0.7696	0.8432	0.7995	0.8413	0.6156	0.0793	0.7912	0.9107	0.6890	19.5072	25.4064	15.1383	0.7351	0.5388	9.1733	-20.7737	8.4604	-21.6481	9.1590	-20.8695
7	E_{BSS}, E_{BSS}^{LUMO}	0.6215	0.0827	0.6746	0.0296	0.7042	0.0039	0.9620	1.0882	0.9534	23.8095	1	0.6297	0.6274	0.7095	0.8264	0.7804	0.7869	0.5649	0.0916	0.7779	0.8898	0.7843	21.2838	27.2379	19.0886	0.7031	0.5278	9.3865	-20.4639	9.6104	-20.6987	9.4819	-20.6350
8	E_{HF}, E_{HF}^{HOMO}	0.6187	0.0863	0.6754	0.0295	0.7049	0.3503	0.9608	1.0923	0.8940	23.8899	1	0.6744	0.6725	0.7446	0.8269	0.7807	0.8242	0.5814	0.0825	0.8225	0.9459	0.7532	21.2333	27.4404	16.7823	0.7094	0.5275	9.6012	-20.3977	8.8795	-21.1091	9.0659	-20.9328
9	E_{BSS}, χ_{BSS}	0.6047	0.0878	0.6617	0.0308	0.6924	0.0729	0.9809	1.1122	0.9852	22.5143	0	0.6046	0.6022	0.6898	0.8183	0.7706	0.7889	0.5328	0.0941	0.8245	0.9444	0.8612	22.1317	28.4482	20.3816	0.6990	0.4723	9.3955	-20.6414	8.5962	-21.2617	9.2430	-20.9216
10	E_{HF}, E_{HF}^{LUMO}	0.5884	0.0923	0.6488	0.0319	0.6807	-0.0106	0.9995	1.1348	1.0153	21.3182	0	0.5800	0.5775	0.6705	0.8100	0.7554	0.7406	0.5497	0.1019	0.7422	0.8465	0.8196	22.9769	29.6209	21.6474	0.6859	0.4761	9.0185	-21.3565	8.6193	-21.0723	9.2978	-21.0592
11	E_{BSS}, ϕ_{BSS}	0.5627	0.1155	0.6460	0.0322	0.6782	0.0757	1.0034	1.1697	1.0247	21.0766	0	0.5722	0.5696	0.6644	0.8083	0.7468	0.7315	0.5445	0.0983	0.7815	0.9211	0.8266	23.1555	31.4702	22.0523	0.6840	0.4408	8.2722	-26.5454	9.0316	-21.1739	9.0901	-23.9524
12	E_{BSS}, E_{BSS}^{HOMO}	0.5593	0.1014	0.6268	0.0339	0.6607	0.2080	1.0303	1.1742	1.0559	19.4724	0	0.5457	0.5430	0.6436	0.7957	0.7392	0.7526	0.4895	0.0997	0.8580	0.9850	0.9222	24.4159	31.7135	23.4145	0.6657	0.4483	9.3778	-20.4344	9.0118	-20.7525	8.9736	-21.2832
13	E_{HFX}, E_{HF}^{LUMO}	0.5347	0.1026	0.6010	0.0363	0.6373	-0.0427	1.0653	1.2066	1.0967	17.5688	0	0.5100	0.5071	0.6156	0.7785	0.7178	0.7050	0.4733	0.1009	0.8457	0.9679	0.8830	26.1018	33.4839	25.2561	0.6380	0.4174	9.1352	-21.7935	9.0360	-20.7048	9.2193	-21.4392
14	E_{HF}, η_{HF}	0.5316	0.1218	0.6187	0.0347	0.6534	0.2451	1.0414	1.2106	1.0393	18.8484	0	0.5599	0.5572	0.6547	0.7903	0.7222	0.7491	0.4994	0.0962	0.8537	0.9928	0.8703	24.9440	33.7087	22.6851	0.6598	0.4211	9.3622	-20.7423	9.3999	-20.4641	9.4572	-20.6024
15	E_{BSS}, η_{BSS}	0.5042	0.1160	0.5822	0.0380	0.6202	0.2183	1.0900	1.2454	1.1716	16.3314	0	0.4408	0.4374	0.5613	0.7656	0.6977	0.6548	0.4273	0.1078	0.9029	1.0424	0.9618	27.3284	35.6752	28.8243	0.6286	0.4003	9.4517	-20.5369	8.8553	-21.5242	8.8120	-21.4746

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

$N_{Ext.O.K}$ Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S12. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange (X) only (BSS, HFX and HF) based methods through 30% (random) splitting method.

Model	Variables	N																																
		Q^2_{1LOO}	$R^2-Q^2_{1LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Nrand}	Q^2_{Nrand}	R^2_{Yrand}	Q^2_{Yrand}
1	E_{HFX}, χ_{HFX}	0.7346	0.0619	0.7815	0.0151	0.7965	0.0847	0.7759	0.8861	0.8393	52.8549	5	0.7564	0.7187	0.7619	0.8868	0.8550	0.8767	0.7526	0.0429	0.6078	0.6860	0.6867	18.0606	23.5575	10.5672	0.8009	0.7180	7.2607	-15.0914	6.6898	-15.3408	7.1007	-15.0420
2	E_{HFX}, E_{HFX}^{HOMO}	0.7315	0.0606	0.7767	0.0154	0.7921	0.3632	0.7844	0.8914	0.7958	51.4299	5	0.7810	0.7471	0.7860	0.8840	0.8525	0.8869	0.7626	0.0616	0.5997	0.6760	0.6433	18.4569	23.8356	9.5006	0.7931	0.7194	6.6263	-15.6764	6.6681	-15.5010	7.0796	-15.1925
3	E_{HF}, ω_{HF}	0.7309	0.0562	0.7714	0.0158	0.7872	-0.0998	0.7936	0.8923	0.8836	49.9357	4	0.7300	0.6882	0.7362	0.8809	0.8519	0.8595	0.7126	0.0112	0.5920	0.6643	0.7703	18.8917	23.8839	11.7108	0.7905	0.7037	6.3514	-15.9934	7.0228	-15.0742	7.2439	-15.2148
4	E_{HFX}, ω_{HFX}	0.7197	0.0590	0.7622	0.0164	0.7786	-0.1096	0.8094	0.9108	0.9048	47.4775	4	0.7169	0.6730	0.7233	0.8755	0.8453	0.8540	0.7037	0.0223	0.6123	0.6870	0.7939	19.6533	24.8863	12.2811	0.7813	0.6964	7.1440	-15.3539	6.8225	-15.3359	6.3831	-16.1493
5	E_{HF}, χ_{HF}	0.7109	0.0674	0.7619	0.0164	0.7783	0.0641	0.8100	0.9249	0.8918	47.3911	4	0.7250	0.6824	0.7312	0.8753	0.8410	0.8587	0.7139	0.0346	0.6463	0.7296	0.7300	19.6812	25.6616	11.9292	0.7810	0.6861	7.1613	-15.4603	7.0300	-15.0703	6.5662	-16.2433
6	E_{HF}, E_{HF}^{HOMO}	0.6683	0.0746	0.7238	0.0190	0.7429	0.3233	0.8722	0.9907	0.9306	39.0062	3	0.7005	0.6541	0.7073	0.8525	0.8139	0.8412	0.6774	0.0772	0.6847	0.7725	0.7595	22.8241	29.4451	12.9911	0.7518	0.6238	6.7210	-15.8855	6.7267	-15.5165	7.4222	-14.6763
7	E_{HFX}, η_{HFX}	0.6673	0.0781	0.7265	0.0189	0.7454	0.2594	0.8681	0.9922	0.8312	39.5143	5	0.7611	0.7241	0.7665	0.8541	0.8128	0.8703	0.6953	0.0753	0.6742	0.7595	0.6598	22.6053	29.5356	10.3624	0.7491	0.6390	7.1346	-15.4306	7.0321	-15.2160	7.0952	-15.1003
8	E_{BSS}, E_{BSS}^{LUMO}	0.6404	0.0846	0.7046	0.0204	0.7250	-0.1122	0.9021	1.0315	0.9066	35.5904	3	0.7158	0.6718	0.7222	0.8406	0.7970	0.8490	0.6735	0.0816	0.7097	0.8027	0.7831	24.4123	31.9199	12.3288	0.7313	0.6139	6.6037	-16.0933	7.1237	-14.7420	7.1495	-15.2317
9	E_{BSS}, χ_{BSS}	0.6400	0.0755	0.6944	0.0211	0.7155	0.0224	0.9175	1.0322	0.9633	33.9544	1	0.6791	0.6294	0.6864	0.8342	0.7940	0.8303	0.6560	0.0858	0.7351	0.8240	0.8161	25.2539	31.9605	13.9193	0.7176	0.6210	7.2325	-15.2724	6.4840	-15.6382	6.3503	-16.2029
10	E_{HF}, E_{HF}^{LUMO}	0.6034	0.0766	0.6563	0.0237	0.6800	-0.1336	0.9731	1.0833	0.9696	28.6865	0	0.6749	0.6245	0.6823	0.8095	0.7686	0.8156	0.6208	0.0928	0.7348	0.8180	0.7996	28.4074	35.2056	14.1029	0.6846	0.5735	7.1382	-15.2424	6.7603	-15.3687	6.4490	-16.1687
11	E_{BSS}, E_{BSS}^{HOMO}	0.5589	0.0925	0.6256	0.0258	0.6515	0.2458	1.0155	1.1424	0.9841	25.2330	0	0.6651	0.6132	0.6727	0.7890	0.7374	0.8079	0.5932	0.0992	0.8247	0.9234	0.8107	30.9402	39.1528	14.5275	0.6568	0.5253	6.8770	-15.1902	6.4536	-15.7556	6.8688	-15.4849
12	E_{HFX}, E_{HFX}^{LUMO}	0.5445	0.0824	0.5993	0.0276	0.6269	-0.1423	1.0507	1.1609	1.0227	22.6868	0	0.6383	0.5823	0.6466	0.7707	0.7267	0.7858	0.5588	0.1055	0.8249	0.9146	0.8906	33.1172	40.4324	15.6876	0.6299	0.5080	6.5441	-16.7207	6.8613	-15.1975	7.2964	-15.6170
13	E_{HF}, η_{HF}	0.5399	0.1113	0.6212	0.0261	0.6473	0.2224	1.0216	1.1718	1.0141	24.7744	0	0.6444	0.5893	0.6525	0.7859	0.7244	0.7931	0.5566	0.1081	0.8250	0.9310	0.8050	31.3109	41.1944	15.4254	0.6495	0.5287	6.5382	-16.1358	6.6812	-15.4203	6.8262	-15.2212
14	E_{BSS}, ω_{BSS}	0.4717	0.1708	0.6161	0.0265	0.6426	-0.0423	1.0284	1.2502	0.9574	24.2703	0	0.6830	0.6340	0.6902	0.7824	0.7012	0.8044	0.5410	0.1167	0.8388	0.9773	0.8239	31.7288	46.8934	13.7485	0.6552	0.4773	6.8457	-17.7301	6.6132	-15.4665	7.2000	-16.7876
15	E_{BSS}, η_{BSS}	0.3906	0.1475	0.5040	0.0342	0.5382	0.1043	1.1690	1.3428	1.0468	15.7320	0	0.6210	0.5624	0.6297	0.6998	0.6154	0.7455	0.4558	0.1391	0.9990	1.1324	0.9311	40.9964	54.0931	16.4377	0.5480	0.3769	6.6256	-15.8523	7.0499	-15.3708	6.6508	-15.9554

Models are arranged according to Q^2_{1LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N.Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S13. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange (X) only (B88, HFX and HF) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	R^2	Q^2_{adj}	R^2	R^2_{adj}	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_m	Δr^2_m	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ver}	Q^2_{Ver}	R^2_{Yrand}	Q^2_{Yrand}	R^2_{Yrand}
1	E_{BSS}, ω_{BSS}	0.8940	0.0534	0.9324	0.0150	0.9474	-0.1087	0.5225	0.7416	1.3028	63.0304	0	0.4214	0.4171	0.6729	0.9730	0.9484	0.7442	0.4625	0.2217	0.4339	0.6341	1.0414	2.7298	5.4998	10.1835	0.9547	0.6493	22.6515	-68.8887	21.1679	-70.8873	21.8555	-70.8699
2	E_{HF}, ω_{HF}	0.8890	0.0596	0.9340	0.0147	0.9486	-0.0834	0.5162	0.7588	0.7827	64.6548	5	0.7912	0.7896	0.8819	0.9736	0.9458	0.8983	0.7128	0.0925	0.4373	0.6427	0.6161	2.6647	5.7580	3.6757	0.9560	0.7476	22.0542	-67.9474	23.6261	-63.6271	23.3410	-65.0586
3	E_{HF}, E_{HF}^{LUMO}	0.8875	0.0703	0.9457	0.0121	0.9577	0.4671	0.4682	0.7642	0.5799	79.3350	5	0.8853	0.8845	0.9352	0.9784	0.9462	0.9502	0.7636	0.0700	0.4096	0.6353	0.5081	2.1924	5.8397	2.0179	0.9649	0.7565	22.2405	-67.1080	23.5371	-63.1234	22.6527	-64.4347
4	E_{HF}, E_{HF}^{LUMO}	0.8761	0.0770	0.9397	0.0134	0.9531	0.4500	0.4932	0.8017	0.5739	71.1738	5	0.8877	0.8869	0.9365	0.9760	0.9409	0.9513	0.7662	0.0685	0.4245	0.6580	0.5013	2.4321	6.4275	1.9759	0.9613	0.5697	22.2525	-65.4665	22.5798	-64.3062	24.6753	-58.3598
5	E_{BSS}, χ_{BSS}	0.8729	0.0778	0.9365	0.0141	0.9506	-0.0244	0.5061	0.8122	0.5322	67.4013	5	0.9035	0.9027	0.9454	0.9747	0.9392	0.9541	0.8169	0.0620	0.4020	0.6113	0.5028	2.5615	6.5961	1.6992	0.9594	0.5569	22.5306	-64.6910	21.1065	-75.4198	22.1647	-67.7708
6	E_{HF}, η_{HF}	0.8654	0.0760	0.9246	0.0168	0.9413	0.0651	0.5517	0.8358	0.9202	56.1759	5	0.7113	0.7092	0.8368	0.9698	0.9352	0.8932	0.6965	0.1163	0.5008	0.7432	0.7741	3.0433	6.9850	5.0802	0.9474	0.6552	22.7485	-62.9075	21.0681	-70.4397	23.6500	-65.5799
7	E_{HF}, χ_{HF}	0.8632	0.0814	0.9288	0.0158	0.9446	-0.0197	0.5362	0.8426	0.5892	59.6689	5	0.8816	0.8808	0.9331	0.9715	0.9343	0.9427	0.7817	0.0595	0.4565	0.6885	0.5553	2.8750	7.0999	2.0831	0.9533	0.6353	22.6467	-63.7464	22.3072	-68.0599	21.4049	-67.6395
8	E_{BSS}, E_{BSS}^{LUMO}	0.8623	0.0853	0.9326	0.0150	0.9476	0.2249	0.5214	0.8453	0.5440	63.3035	5	0.8991	0.8984	0.9430	0.9731	0.9344	0.9553	0.7817	0.0631	0.4120	0.6302	0.4836	2.7186	7.1459	1.7754	0.9574	0.6811	22.2227	-62.9682	21.0369	-70.6898	23.3120	-63.7170
9	E_{BSS}, E_{BSS}^{LUMO}	0.8612	0.0815	0.9262	0.0164	0.9426	-0.0961	0.5456	0.8488	0.8722	57.5012	5	0.7407	0.7387	0.8534	0.9705	0.9333	0.8720	0.6925	0.1437	0.4027	0.6030	0.7259	2.9772	7.2047	4.5644	0.9526	0.6944	21.5767	-67.4983	21.4183	-68.9900	21.1444	-66.5936
10	E_{HF}, η_{HF}	0.8597	0.0822	0.9254	0.0166	0.9420	0.0480	0.5488	0.8531	1.1388	56.7945	3	0.5579	0.5546	0.7501	0.9701	0.9327	0.8523	0.6695	0.1342	0.4924	0.7392	0.9910	3.0121	7.2780	7.7807	0.9495	0.6150	22.7995	-63.2418	22.1783	-73.5844	21.7160	-66.0662
11	E_{HF}, χ_{HF}	0.8486	0.0897	0.9208	0.0176	0.9384	-0.0328	0.5654	0.8862	0.5563	53.3046	5	0.8945	0.8937	0.9404	0.9682	0.9274	0.9496	0.7888	0.0585	0.4469	0.6729	0.5339	3.1971	7.8541	1.8566	0.9490	0.6067	21.1746	-64.9958	22.3567	-67.8936	22.7263	-61.2029
12	E_{HF}, ω_{HF}	0.8471	0.0843	0.9118	0.0196	0.9314	-0.0892	0.5966	0.8908	0.7243	47.5212	5	0.8212	0.8199	0.8989	0.9645	0.9259	0.9141	0.7271	0.0886	0.4797	0.7092	0.5765	3.5595	7.9359	3.1472	0.9412	0.4597	22.2852	-66.8020	21.3160	-68.6635	22.5408	-63.4197
13	E_{BSS}, η_{BSS}	0.7856	0.1238	0.8835	0.0259	0.9094	0.0511	0.6858	1.0549	1.1478	35.1183	1	0.5509	0.5476	0.7461	0.9525	0.8968	0.8465	0.5636	0.1703	0.5870	0.8661	0.9898	4.7027	11.1276	7.9041	0.9225	0.4913	23.3746	-62.0947	22.4306	-68.1555	22.2886	-64.8389
14	E_{HF}, E_{HF}^{LUMO}	0.7480	0.1427	0.8594	0.0312	0.8907	-0.0977	0.7532	1.1435	1.2496	28.5148	0	0.4677	0.4638	0.6991	0.9422	0.8764	0.7363	0.5216	0.1981	0.5976	0.8857	1.0200	5.6727	13.0771	9.3684	0.9057	0.3073	23.9993	-61.6300	21.5777	-68.3425	21.8983	-65.5037
15	E_{HF}, E_{HF}^{LUMO}	0.6597	0.2003	0.8200	0.0400	0.8600	-0.0956	0.8523	1.3288	1.1782	21.5007	1	0.5267	0.5232	0.7325	0.9247	0.8328	0.7623	0.5506	0.1900	0.7038	1.0566	0.9473	7.2642	17.6565	8.3295	0.8847	0.1199	22.5905	-62.4541	21.2489	-71.6427	21.2575	-65.9351

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S14. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange (X) only (B88, HFX and HF) based methods through 30% (random) splitting method.

Model	Variables	Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	N	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{F}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Year}	Q^2_{Year}	R^2_{Xrand}	Q^2_{Xrand}	R^2_{Yrand}	Q^2_{Yrand}
1	E_{HF}, E_{HF}^{HOMO}	0.8963	0.0542	0.9363	0.0142	0.9505	0.3682	0.4562	0.6601	0.3709	67.1375	5	0.9643	0.9643	0.9672	0.9746	0.9469	0.9826	0.8863	0.0205	0.3858	0.5554	0.3024	2.0808	4.3567	0.6879	0.9541	0.7359	24.1909	-56.6951	23.1834	-66.5499	22.0206	-61.7668	
2	E_{HFX}, E_{HFX}^{HOMO}	0.8917	0.0567	0.9336	0.0147	0.9484	0.3799	0.4656	0.6744	0.3785	64.3175	5	0.9629	0.9628	0.9659	0.9735	0.9445	0.9820	0.8824	0.0220	0.3936	0.5671	0.3204	2.1674	4.5480	0.7163	0.9532	0.7187	22.9371	-58.4484	22.9182	-67.8165	22.4006	-60.0329	
3	E_{HF}, E_{HF}^{HOMO}	0.8878	0.0613	0.9346	0.0145	0.9491	0.1252	0.4623	0.6865	0.5594	65.2636	5	0.9189	0.9188	0.9255	0.9739	0.9425	0.9622	0.8140	0.0475	0.3916	0.5698	0.4661	2.1376	4.7127	1.5648	0.9543	0.7698	23.3206	-60.1189	22.7955	-65.2709	21.4112	-64.5934	
4	E_{HFX}, E_{HFX}^{HOMO}	0.8828	0.0651	0.9329	0.0149	0.9478	0.0850	0.4681	0.7017	0.5502	63.5912	5	0.9215	0.9215	0.9279	0.9732	0.9398	0.9644	0.8144	0.0461	0.3842	0.5617	0.4720	2.1908	4.9239	1.5138	0.9532	0.7186	21.9321	-61.7354	22.2159	-66.3975	21.5205	-63.3489	
5	E_{BSS}, E_{BSS}^{HOMO}	0.8826	0.0636	0.9309	0.0154	0.9462	0.4568	0.4752	0.7021	0.3764	61.5797	5	0.9633	0.9633	0.9663	0.9724	0.9393	0.9815	0.9066	0.0203	0.3923	0.5776	0.3068	2.2586	4.9299	0.7084	0.9525	0.7535	22.0475	-61.2091	21.5975	-69.0357	22.1889	-61.8658	
6	E_{BSS}, E_{BSS}^{HOMO}	0.8802	0.0670	0.9322	0.0151	0.9472	0.1505	0.4707	0.7092	0.4902	62.8406	5	0.9377	0.9377	0.9428	0.9729	0.9382	0.9702	0.8392	0.0358	0.4140	0.6134	0.4079	2.2156	5.0293	1.2016	0.9537	0.7526	20.9841	-66.8600	23.0754	-66.5819	21.6673	-65.8154	
7	E_{HFX}, E_{HFX}^{HOMO}	0.8692	0.0586	0.9073	0.0206	0.9279	0.0936	0.5504	0.7411	0.4529	45.0176	5	0.9469	0.9468	0.9512	0.9626	0.9327	0.9716	0.9356	0.0264	0.4143	0.5841	0.3548	3.0295	5.4920	1.0255	0.9299	0.6286	21.2630	-65.4999	22.0305	-68.3950	21.7036	-63.0758	
8	E_{HF}, E_{HF}^{HOMO}	0.8529	0.0661	0.8958	0.0232	0.9190	0.0880	0.5834	0.7860	0.4285	39.6891	5	0.9524	0.9524	0.9563	0.9578	0.9243	0.9748	0.9427	0.0264	0.4664	0.6553	0.3259	3.4033	6.1775	0.9182	0.9224	0.5815	22.9847	-62.0211	21.9714	-69.1585	22.7891	-61.6714	
9	E_{HF}, E_{HF}^{HOMO}	0.8515	0.0852	0.9186	0.0181	0.9367	-0.0072	0.5156	0.7898	0.7370	51.7953	5	0.8592	0.8591	0.8707	0.9673	0.9241	0.9375	0.7525	0.0765	0.4329	0.6448	0.6180	2.6582	6.2381	2.7159	0.9454	0.5902	21.8853	-64.0207	23.2411	-67.2532	21.7012	-63.4208	
10	E_{BSS}, E_{BSS}^{HOMO}	0.8478	0.0651	0.8880	0.0249	0.9129	0.0973	0.6047	0.7995	0.5561	36.6929	5	0.9199	0.9198	0.9264	0.9545	0.9206	0.9557	0.8823	0.0526	0.4451	0.6112	0.4586	3.6570	6.3916	1.5464	0.9167	0.5958	22.0327	-61.6641	21.7324	-69.8288	22.1344	-61.1897	
11	E_{HFX}, E_{HFX}^{HOMO}	0.8367	0.0969	0.9146	0.0190	0.9336	-0.0331	0.5282	0.8282	0.7644	49.1755	5	0.8486	0.8485	0.8609	0.9656	0.9166	0.9364	0.7479	0.0763	0.4338	0.6523	0.6560	2.7904	6.8594	2.9218	0.9441	0.6665	21.0764	-65.4603	21.8851	-69.6489	22.5857	-61.8280	
12	E_{BSS}, E_{BSS}^{HOMO}	0.8252	0.1027	0.9073	0.0206	0.9279	-0.0376	0.5503	0.8569	0.8481	45.0298	5	0.8136	0.8135	0.8287	0.9626	0.9103	0.9178	0.6996	0.0959	0.4535	0.6887	0.6620	3.0288	7.3430	3.5961	0.9380	0.5603	21.8862	-69.8507	23.5395	-63.3053	22.1350	-67.1439	
13	E_{HF}, E_{HF}^{HOMO}	0.7377	0.1493	0.8547	0.0323	0.8870	-0.0551	0.6889	1.0495	1.1018	27.4691	2	0.6854	0.6852	0.7110	0.9401	0.8652	0.8630	0.6254	0.1425	0.5701	0.8325	0.8985	4.7462	11.0150	6.0694	0.8992	0.4532	21.4773	-65.7349	21.1768	-66.4496	21.2866	-65.3895	
14	E_{BSS}, E_{BSS}^{HOMO}	0.7064	0.1812	0.8556	0.0321	0.8877	-0.0585	0.6869	1.1103	1.1413	27.6559	1	0.6624	0.6622	0.6898	0.9405	0.8521	0.8584	0.5974	0.1467	0.5533	0.8595	0.9116	4.7177	12.3286	6.5130	0.9000	0.3259	20.3550	-71.7720	20.7038	-71.0775	23.5845	-62.1699	
15	E_{HFX}, E_{HFX}^{HOMO}	0.6636	0.2161	0.8454	0.0344	0.8797	-0.0646	0.7107	1.1886	1.1444	25.5999	1	0.6606	0.6604	0.6882	0.9360	0.8308	0.8611	0.6205	0.1432	0.5916	0.9281	0.9512	5.0511	14.1276	6.5482	0.8998	0.2864	23.0206	-63.1027	22.8256	-63.6073	23.8703	-63.1084	

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N. Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S15. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange-correlation (HFX+LYP, BLYP and B3LYP) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																															
		Q^2_{LOO}	R^2_{LOO}	R^2_{adj}	R^2_{adj}	R^2	ΔK	$RMSE_{TR}$	$RMSE_{CV}$	$RMSE_{EXT}$	$Ext.OK$	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC_{TR}	CCC_{CV}	CCC_{EXT}	\bar{F}^2_{int}	Δr^2_{int}	MAE_{TR}	MAE_{CV}	MAE_{EXT}	RSS_{TR}	$PRESS_{CV}$	$PRESS_{EXT}$	R^2_{LMO}	Q^2_{LMO}	R^2_{Year}	Q^2_{Year}	R^2_{Xrand}	Q^2_{Xrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{HFX+LYP}^{HOMO}, E_{HFX+LYP}^{HOMO}$	0.7102	0.0643	0.7520	0.0225	0.7746	0.4233	0.8398	0.9521	0.8344	5	0.7461	0.7461	0.7775	0.8730	0.8382	0.8787	0.7054	0.0851	0.6806	0.7819	0.6753	16.2222	20.8512	15.3153	0.7775	0.6160	9.0839	-20.9894	9.2415	-20.3120	8.8312	-21.1261
2	$E_{HFX+LYP}, \eta_{HFX+LYP}$	0.6867	0.0866	0.7506	0.0227	0.7733	0.2414	0.8422	0.9900	0.8964	5	0.7069	0.7069	0.7432	0.8721	0.8266	0.8583	0.6895	0.0966	0.6402	0.7470	0.7482	16.3144	22.5441	17.6789	0.7765	0.5693	9.1633	-20.9206	8.5543	-21.5851	8.8513	-21.2185
3	$E_{HFX+LYP}, \chi_{HFX+LYP}$	0.6745	0.0769	0.7265	0.0249	0.7514	0.1945	0.8819	1.0092	0.8557	5	0.7330	0.7330	0.7660	0.8581	0.8164	0.8719	0.6747	0.0864	0.7212	0.8306	0.6924	17.8894	23.4237	16.1075	0.7563	0.5868	9.6330	-20.2007	9.3072	-20.3673	8.8481	-21.5107
4	E_{BLYP}, χ_{BLYP}	0.6442	0.0838	0.7008	0.0272	0.7280	0.0973	0.9225	1.0551	0.9294	2	0.6850	0.6849	0.7239	0.8426	0.7974	0.8519	0.6123	0.0918	0.7629	0.8786	0.7800	19.5737	25.6021	19.0037	0.7308	0.5560	8.2667	-22.2341	9.2662	-20.4744	9.0706	-20.9955
5	$E_{BLYP}, E_{BLYP}^{HOMO}$	0.6359	0.0938	0.7027	0.0270	0.7297	0.2344	0.9196	1.0674	0.9313	2	0.6837	0.6837	0.7228	0.8437	0.7934	0.8530	0.6145	0.0917	0.7517	0.8725	0.7828	19.4509	26.2039	19.0805	0.7346	0.5412	8.7711	-21.1473	8.8202	-21.2763	8.4491	-21.7246
6	$E_{HFX+LYP}, \omega_{HFX+LYP}$	0.6340	0.0842	0.6900	0.0282	0.7182	0.0040	0.9390	1.0701	0.9185	1	0.6923	0.6923	0.7304	0.8360	0.7897	0.8478	0.6205	0.0933	0.7612	0.8726	0.7580	20.2798	26.3393	18.5600	0.7192	0.5506	9.6501	-20.1649	8.7927	-21.1156	8.9105	-21.5348
7	$E_{B3LYP}, E_{B3LYP}^{LUMO}$	0.6212	0.0786	0.6698	0.0300	0.6998	0.0066	0.9691	1.0887	0.9832	0	0.6475	0.6474	0.6910	0.8234	0.7783	0.8203	0.5891	0.1029	0.7415	0.8391	0.8079	21.6027	27.2612	21.2657	0.7047	0.4927	9.4259	-20.9756	9.1074	-20.7971	9.4459	-20.8007
8	E_{B3LYP}, χ_{B3LYP}	0.6174	0.0872	0.6751	0.0295	0.7046	0.0943	0.9614	1.0941	0.9969	0	0.6376	0.6375	0.6823	0.8267	0.7800	0.8320	0.5857	0.0998	0.8262	0.9494	0.8564	21.2566	27.5318	21.8647	0.7079	0.5223	8.9350	-21.0976	8.9309	-20.9682	9.3311	-20.6773
9	$E_{BLYP}, E_{BLYP}^{LUMO}$	0.6070	0.0930	0.6700	0.0300	0.7000	-0.0065	0.9688	1.1088	0.9637	1	0.6613	0.6613	0.7032	0.8236	0.7717	0.8324	0.5897	0.0976	0.7906	0.9079	0.7992	21.5852	28.2782	20.4313	0.7061	0.5061	8.8386	-21.4346	8.9275	-21.4680	9.6972	-20.2766
10	$E_{B3LYP}, \omega_{B3LYP}$	0.5935	0.0860	0.6474	0.0321	0.6795	0.0847	1.0014	1.1277	1.0363	0	0.6083	0.6083	0.6567	0.8091	0.7625	0.7897	0.5586	0.1096	0.7666	0.8867	0.8516	23.0659	29.2514	23.6283	0.6772	0.4773	9.2834	-29.2417	9.4739	-20.4226	8.7891	-29.2942
11	$E_{B3LYP}, E_{B3LYP}^{HOMO}$	0.5756	0.0981	0.6411	0.0326	0.6738	0.2485	1.0103	1.1523	1.0680	0	0.5840	0.5840	0.6354	0.8051	0.7512	0.8055	0.5449	0.1073	0.8489	0.9764	0.9253	23.4767	30.5376	25.0934	0.6797	0.4585	9.3052	-21.0522	9.6978	-20.2786	9.3429	-21.0456
12	E_{BLYP}, ω_{BLYP}	0.5573	0.0854	0.6069	0.0357	0.6426	0.0078	1.0574	1.1769	1.0841	0	0.5714	0.5713	0.6243	0.7824	0.7298	0.7680	0.5272	0.1135	0.8260	0.9292	0.8941	25.7155	31.8579	25.8565	0.6457	0.4365	8.5693	-22.7165	9.3737	-20.0531	9.1412	-21.2872
13	$E_{HFX+LYP}, E_{HFX+LYP}^{LUMO}$	0.5289	0.1074	0.5999	0.0364	0.6363	-0.0372	1.0667	1.2140	1.1115	0	0.5494	0.5494	0.6051	0.7777	0.7141	0.7681	0.5064	0.1134	0.8464	0.9712	0.9084	26.1716	33.8989	27.1818	0.6420	0.4090	8.6417	-22.3318	9.0061	-20.9481	8.5792	-22.2710
14	E_{B3LYP}, η_{B3LYP}	0.4939	0.1350	0.5917	0.0371	0.6289	0.2795	1.0776	1.2584	1.1653	0	0.5048	0.5047	0.5660	0.7721	0.6953	0.7511	0.4819	0.1179	0.8951	1.0462	0.9849	26.7071	36.4215	29.8745	0.6352	0.3820	9.3319	-21.6752	8.9937	-20.8505	9.8314	-20.4240
15	E_{BLYP}, η_{BLYP}	0.4165	0.2857	0.6724	0.0298	0.7022	0.2848	0.9653	1.3511	0.9938	0	0.6399	0.6398	0.6844	0.8251	0.6809	0.8300	0.5849	0.1003	0.7574	0.9498	0.8538	21.4295	41.9882	21.7263	0.7278	0.4099	9.3087	-22.1177	8.6959	-21.4791	9.2233	-22.2913

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_ExtOK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S16. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using exchange-correlation (HFX+LYP, BLYP and B3LYP) based methods through 30% (random) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{F}^2_m	Δr^2_m	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Year}	Q^2_{Year}	R^2_{Nrand}	Q^2_{Nrand}	R^2_{Yrand}	Q^2_{Yrand}
1	E_{BLYP} / η_{BLYP}	0.7206	0.0530	0.7599	0.0137	0.7736	0.1968	0.8259	0.9175	1.6134	56.3837	0	0.0765	0.0764	0.1360	0.8724	0.8452	0.5029	0.1946	0.0069	0.6561	0.7241	1.0737	24.5558	30.3053	23.4282	0.7774	0.7090	5.7280	-12.5098	5.8243	-12.2167	5.7656	-12.3989
2	$E_{HFX+LYP} / \eta_{HFX+LYP}$	0.7193	0.0512	0.7566	0.0139	0.7705	0.2638	0.8316	0.9197	0.6273	55.3960	5	0.8604	0.8604	0.8694	0.8704	0.8443	0.9200	0.7929	0.0358	0.6715	0.7382	0.5216	24.8932	30.4511	3.5417	0.7758	0.6967	5.8044	-12.3388	5.5544	-12.5512	5.9020	-12.4195
3	$E_{HFX+LYP} / E_{HFX+LYP}^{HOMO}$	0.7181	0.0472	0.7511	0.0142	0.7653	0.3512	0.8409	0.9215	0.6402	53.8078	5	0.8546	0.8546	0.8640	0.8671	0.8427	0.9196	0.7642	0.0424	0.6758	0.7404	0.4998	25.4555	30.5727	3.6886	0.7684	0.6975	5.7730	-12.4151	5.7259	-12.3442	5.9728	-12.1645
4	$E_{BLYP} / E_{BLYP}^{HOMO}$	0.6855	0.0526	0.7222	0.0159	0.7381	0.2347	0.8884	0.9735	1.0451	46.4957	0	0.6125	0.6125	0.6375	0.8493	0.8217	0.7693	0.5721	0.0488	0.7156	0.7842	0.7857	28.4102	34.1162	9.8295	0.7413	0.6640	5.4992	-12.8002	5.9906	-12.1287	5.9231	-12.0892
5	$E_{HFX+LYP} / \chi_{HFX+LYP}$	0.6813	0.0523	0.7174	0.0161	0.7336	0.0715	0.8960	0.9800	0.6828	45.4273	5	0.8346	0.8346	0.8453	0.8463	0.8190	0.9066	0.7596	0.0483	0.7195	0.7876	0.5311	28.9003	34.5720	4.1958	0.7386	0.6619	5.7890	-12.4175	5.7839	-12.2668	5.8909	-12.3150
6	E_{BLYP} / χ_{BLYP}	0.6622	0.0548	0.6999	0.0171	0.7171	0.0414	0.9233	1.0088	0.8912	41.8221	4	0.7183	0.7182	0.7364	0.8352	0.8061	0.8298	0.7062	0.0859	0.7464	0.8161	0.6945	30.6868	36.6352	7.1477	0.7218	0.6396	6.2234	-11.7809	5.6540	-12.4897	5.6782	-12.3711
7	E_{B3LYP} / χ_{B3LYP}	0.6363	0.0560	0.6737	0.0186	0.6923	0.0578	0.9628	1.0468	0.9086	37.1276	4	0.7071	0.7071	0.7260	0.8182	0.7883	0.8268	0.6696	0.0890	0.7982	0.8710	0.7412	33.3731	39.4516	7.4307	0.6973	0.6153	6.0531	-12.2267	6.0337	-12.0689	5.5223	-12.7575
8	$E_{HFX+LYP} / \omega_{HFX+LYP}$	0.6310	0.0565	0.6686	0.0189	0.6875	-0.0912	0.9703	1.0544	0.7216	36.3037	5	0.8153	0.8152	0.8272	0.8148	0.7845	0.8910	0.7794	0.0547	0.7965	0.8678	0.5480	33.8938	40.0212	4.6864	0.6930	0.6056	5.7669	-12.3057	5.6544	-12.5262	5.9757	-12.3540
9	$E_{B3LYP} / E_{B3LYP}^{HOMO}$	0.6157	0.0591	0.6550	0.0197	0.6747	0.2897	0.9900	1.0761	1.0253	34.2262	0	0.6271	0.6270	0.6511	0.8058	0.7736	0.7724	0.6125	0.1103	0.8241	0.8987	0.8221	35.2820	41.6890	9.4610	0.6789	0.5937	5.4405	-13.0584	5.4296	-12.6781	5.8064	-12.3999
10	$E_{BLYP} / E_{BLYP}^{LUMO}$	0.6063	0.0617	0.6480	0.0201	0.6681	-0.0932	1.0000	1.0891	0.7427	33.2104	5	0.8043	0.8043	0.8169	0.8010	0.7676	0.8802	0.7981	0.0545	0.8177	0.8915	0.5847	36.0030	42.7007	4.9642	0.6737	0.5759	5.9042	-12.3315	5.7791	-12.2880	5.7610	-12.2556
11	$E_{B3LYP} / E_{B3LYP}^{LUMO}$	0.5835	0.0639	0.6260	0.0214	0.6474	-0.0971	1.0307	1.1203	0.7204	30.2957	5	0.8159	0.8158	0.8277	0.7860	0.7509	0.8855	0.8213	0.0432	0.8410	0.9157	0.5506	38.2454	45.1815	4.6713	0.6490	0.5710	5.6929	-12.4902	5.6884	-12.4562	5.4763	-12.8388
12	$E_{B3LYP} / \omega_{B3LYP}$	0.5412	0.0696	0.5872	0.0236	0.6108	-0.0681	1.0829	1.1757	0.7637	25.8964	5	0.7931	0.7931	0.8064	0.7584	0.7203	0.8625	0.8768	0.0323	0.8871	0.9670	0.5611	42.2140	49.7654	5.2489	0.6135	0.5165	5.6379	-13.0705	5.4357	-12.7487	5.7049	-12.7142
13	E_{BLYP} / ω_{BLYP}	0.5262	0.0686	0.5703	0.0246	0.5949	-0.1342	1.1048	1.1948	0.7517	24.2270	5	0.7995	0.7995	0.8124	0.7460	0.7070	0.8800	0.7590	0.0507	0.8894	0.9648	0.5735	43.9443	51.3906	5.0859	0.5993	0.4986	5.9825	-12.4072	5.5046	-12.5163	5.8950	-12.3335
14	E_{B3LYP} / η_{B3LYP}	0.5208	0.0926	0.5900	0.0234	0.6135	0.1973	1.0792	1.2015	1.1270	26.1864	0	0.5494	0.5494	0.5785	0.7604	0.7094	0.6925	0.5371	0.0031	0.9060	0.9981	0.8760	41.9272	51.9724	11.4303	0.6226	0.5008	5.7097	-12.6794	5.6244	-12.6333	6.1563	-12.1235
15	$E_{HFX+LYP} / E_{HFX+LYP}^{LUMO}$	0.5022	0.0754	0.5521	0.0256	0.5777	-0.1044	1.1280	1.2246	0.9171	22.5702	4	0.7016	0.7016	0.7208	0.7323	0.6891	0.7904	0.7816	0.0381	0.9238	1.0038	0.7075	45.8078	53.9915	7.5698	0.5823	0.4722	5.4644	-13.1121	5.4830	-12.8185	5.5700	-12.9921

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S17. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{T1}	Q^2_{T2}	Q^2_{T3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{F}_m^2	Δr_m^2	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Year}	Q^2_{Year}	R^2_{Yrand}	Q^2_{Yrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{M06-L}, E_{M06-L}^{LUMO}$	0.6061	0.1087	0.6863	0.0285	0.7148	-0.1507	0.9477	1.1138	0.8564	25.0673	3	0.704	0.6948	0.7671	0.8337	0.7767	0.85	0.6734	0.0428	0.7993	0.9312	0.6423	20.6559	28.5327	14.6669	0.7156	0.5170	8.9048	-21.3705	9.0194	-20.6567	8.8857	-21.1523
2	$E_{M06-2X}, \omega_{M06-2X}$	0.5977	0.1046	0.6726	0.0298	0.7024	-0.142	0.9682	1.1256	0.9098	23.5971	1	0.6659	0.6555	0.7372	0.8252	0.769	0.8243	0.6243	0.0471	0.7889	0.9146	0.7072	21.5598	29.1392	16.5556	0.7070	0.5029	8.9929	-21.1497	9.034	-20.8322	9.1942	-21.1689
3	E_{M06}, E_{M06}^{LUMO}	0.5889	0.1087	0.6673	0.0302	0.6976	-0.148	0.9759	1.1379	0.8737	23.0655	1	0.6919	0.6823	0.7576	0.8218	0.7634	0.8367	0.6478	0.0861	0.8052	0.9341	0.6765	21.9064	29.7801	15.2679	0.6998	0.5108	9.063	-20.7371	8.9168	-21.0906	9.1405	-20.9918
4	$E_{M06-2X}, \chi_{M06-2X}$	0.5591	0.118	0.6448	0.0323	0.6771	-0.0466	1.0084	1.1784	0.7806	20.9716	5	0.7541	0.7464	0.8065	0.8075	0.746	0.8679	0.7086	0.0291	0.8636	1.0032	0.6148	23.3874	31.9369	12.1875	0.6840	0.4634	8.8934	-21.4506	9.3654	-20.5769	9.0199	-21.3408
5	$E_{M06-2X}, E_{M06-2X}^{LUMO}$	0.5569	0.1123	0.6361	0.0331	0.6692	-0.1455	1.0207	1.1813	0.8955	20.2279	1	0.6763	0.6663	0.7454	0.8018	0.74	0.8157	0.6138	0.1117	0.8199	0.947	0.6872	23.9628	32.0982	16.0391	0.6733	0.4473	9.2154	-20.9118	9.3731	-20.6479	9.0196	-21.3004
6	E_{M06}, χ_{M06}	0.5565	0.1201	0.6443	0.0323	0.6766	-0.0468	1.0092	1.1818	0.7807	20.9211	5	0.754	0.7463	0.8065	0.8071	0.745	0.8693	0.7113	0.0119	0.8653	1.0077	0.6071	23.4256	32.1243	12.1909	0.6802	0.4656	8.912	-21.6633	9.0612	-20.9569	9.1491	-21.0229
7	E_{M06}, ω_{M06}	0.5391	0.1182	0.6231	0.0343	0.6573	-0.1084	1.0388	1.2047	0.9293	19.1839	1	0.6515	0.6406	0.7258	0.7933	0.7306	0.789	0.582	0.0629	0.8344	0.9684	0.7034	24.82	33.3823	17.2711	0.6630	0.4273	8.9816	-21.9561	9.6895	-19.909	9.2673	-21.1627
8	E_{M06-L}, χ_{M06-L}	0.5193	0.126	0.6099	0.0355	0.6454	-0.0467	1.0568	1.2304	0.7946	18.197	5	0.7452	0.7372	0.7995	0.7845	0.7174	0.8591	0.6963	0.0024	0.9039	1.0479	0.6405	25.6888	34.8171	12.629	0.6537	0.3933	8.6882	-21.8218	8.9025	-20.9187	9.0413	-21.0772
9	$E_{M06-L}, \omega_{M06-L}$	0.4953	0.123	0.5802	0.0382	0.6183	-0.0488	1.0963	1.2608	0.9983	16.2015	0	0.5978	0.5852	0.6835	0.7642	0.6981	0.7252	0.5374	0.0578	0.9112	1.0535	0.773	27.6452	36.5583	19.9331	0.6187	0.4057	9.3782	-22.4399	8.7937	-21.2473	8.6473	-23.5715
10	$E_{M06-2X}, E_{M06-2X}^{HOMO}$	0.4888	0.1367	0.588	0.0375	0.6255	0.153	1.086	1.2689	0.8582	16.7008	2	0.7027	0.6935	0.7661	0.7696	0.6954	0.8249	0.6459	0.0071	0.9022	1.0477	0.6951	27.1282	37.0297	14.7315	0.6371	0.3720	9.1317	-20.8141	9.5143	-20.106	9.2712	-21.0597
11	E_{M06}, E_{M06}^{HOMO}	0.484	0.1389	0.5852	0.0377	0.6229	0.1093	1.0897	1.2747	0.8562	16.5212	2	0.7042	0.6949	0.7672	0.7677	0.6924	0.8278	0.65	0.0102	0.9101	1.0582	0.6955	27.312	37.3733	14.6608	0.6301	0.4248	9.1619	-21.0887	9.082	-20.9086	9.532	-20.3311
12	$E_{M06-L}, E_{M06-L}^{HOMO}$	0.4349	0.1566	0.5508	0.0408	0.5916	0.0309	1.1341	1.334	0.8854	14.4855	1	0.6836	0.6738	0.7511	0.7434	0.6567	0.8063	0.6276	0.0024	0.9396	1.0929	0.7227	29.5827	40.9295	15.6783	0.6033	0.3438	8.7519	-21.5956	9.1232	-20.9926	8.9122	-21.3319
13	$E_{M06-2X}, \eta_{M06-2X}$	0.3856	0.1803	0.5225	0.0434	0.5659	0.301	1.1692	1.391	0.9948	13.0388	0	0.6006	0.5881	0.6858	0.7228	0.6213	0.7377	0.5452	0.0009	0.9524	1.1162	0.8255	31.4403	44.5033	19.7931	0.5809	0.2967	8.7939	-21.6215	9.3398	-20.5981	9.3088	-20.5606
14	E_{M06}, η_{M06}	0.372	0.1849	0.5126	0.0443	0.5569	0.2925	1.1813	1.4064	1.0036	12.5682	0	0.5935	0.5809	0.6802	0.7154	0.6107	0.7303	0.5405	0.0009	0.9693	1.136	0.8309	32.0958	45.4913	20.1428	0.5657	0.2913	9.217	-21.1696	8.9569	-20.9562	8.9464	-21.4671
15	E_{M06-L}, η_{M06-L}	0.3404	0.1933	0.4871	0.0466	0.5337	0.0672	1.2118	1.4413	1.0351	11.4462	0	0.5676	0.5541	0.6598	0.696	0.5836	0.6928	0.5231	0.0422	1.0221	1.2016	0.8467	33.775	47.7775	21.4295	0.5483	0.2225	8.9472	-21.8243	8.6665	-21.3824	9.2377	-21.4807

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecolox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N.Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S18. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through 30% (random) splitting method.

Model	Variables	N.																																
		Q^2_{LOO}	R^2_{LOO}	R^2_{adj}	R^2_{adj}	R^2_{adj}	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_m	Δr^2_m	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Xrand}	Q^2_{Xrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{M06-L}, E_{LUMO}^{LUMO}$	0.6457	0.059	0.6836	0.0211	0.7047	-0.1039	0.9344	1.0235	0.7741	33.4027	5	0.8045	0.7692	0.7973	0.8267	0.7963	0.8539	0.7385	0.0611	0.7329	0.8094	0.6806	27.0677	32.4746	7.7897	0.7072	0.6324	6.631	-14.9168	6.5415	-14.9054	6.8114	-14.65
2	E_{M06-L}, χ_{M06}	0.645	0.0586	0.6824	0.0212	0.7036	0.0425	0.9361	1.0245	0.8487	33.2288	4	0.765	0.7226	0.7563	0.826	0.7956	0.8356	0.6982	0.0366	0.7665	0.8447	0.6791	27.1674	32.5369	9.3648	0.7058	0.6239	7.0021	-14.4278	6.4345	-14.9124	6.6507	-14.9168
3	$E_{M06-2X}, \omega_{M06-2X}$	0.6444	0.0573	0.6804	0.0213	0.7017	-0.0991	0.939	1.0253	0.8381	32.9376	4	0.7708	0.7295	0.7624	0.8247	0.795	0.8262	0.6796	0.0736	0.7251	0.7989	0.7122	27.3359	32.5894	9.1321	0.7047	0.6189	6.4535	-15.2116	6.263	-15.2692	6.8421	-14.8103
4	$E_{M06-2X}, \chi_{M06-2X}$	0.6391	0.0591	0.6766	0.0216	0.6982	0.0362	0.9447	1.033	0.8283	32.3816	4	0.7762	0.7358	0.7679	0.8223	0.7915	0.8418	0.7144	0.0354	0.7836	0.8632	0.664	27.6636	33.0781	8.9188	0.7020	0.6082	6.5018	-15.2048	6.603	-14.8244	6.6091	-15.1715
5	$E_{M06-L}, E_{LUMO}^{LUMO}$	0.6355	0.0597	0.6734	0.0218	0.6951	-0.1043	0.9493	1.0381	0.815	31.9242	4	0.7833	0.7442	0.7753	0.8202	0.7892	0.8358	0.6921	0.0706	0.7311	0.805	0.697	27.9392	33.4086	8.6348	0.6995	0.6169	6.5893	-14.9388	6.3198	-15.155	6.5335	-15.1558
6	E_{M06-L}, ω_{M06}	0.6198	0.0631	0.6602	0.0227	0.6829	-0.0821	0.9682	1.0602	0.9577	30.149	1	0.7008	0.6468	0.6898	0.8116	0.7792	0.7711	0.5522	0.1025	0.7574	0.8361	0.7881	29.0626	34.844	11.9233	0.6894	0.5835	6.521	-15.6171	7.1178	-14.0682	6.9362	-14.9774
7	E_{M06-L}, χ_{M06-L}	0.6156	0.0637	0.6564	0.0229	0.6793	0.0207	0.9736	1.066	0.857	29.6611	4	0.7604	0.7172	0.7516	0.8091	0.7755	0.8252	0.6854	0.0722	0.797	0.8776	0.7075	29.3874	35.2282	9.5474	0.6817	0.5926	6.7324	-14.8209	6.2714	-15.1945	6.9439	-14.6665
8	$E_{M06-2X}, E_{LUMO}^{LUMO}$	0.6119	0.0623	0.6509	0.0233	0.6742	-0.1068	0.9814	1.0712	0.87	28.9697	3	0.7531	0.7085	0.744	0.8054	0.7727	0.8091	0.6293	0.0848	0.75	0.8248	0.7097	29.8602	35.5689	9.8387	0.6771	0.5876	6.2714	-15.4525	6.6731	-14.6313	6.7733	-14.9116
9	$E_{M06-L}, \omega_{M06-L}$	0.5888	0.069	0.6335	0.0244	0.6579	-0.0499	1.0057	1.1025	1.0841	26.9231	0	0.6165	0.5473	0.6024	0.7937	0.7585	0.7067	0.4399	0.1277	0.8148	0.9007	0.9221	31.3536	37.6819	15.2798	0.6668	0.5456	6.5828	-16.1135	6.7211	-14.6945	6.5068	-16.2047
10	$E_{M06-L}, E_{HOMO}^{HOMO}$	0.5881	0.0715	0.6352	0.0243	0.6595	0.3063	1.0033	1.1036	0.9461	27.12	1	0.708	0.6553	0.6972	0.7948	0.7562	0.7899	0.6135	0.0705	0.8077	0.891	0.7751	31.2034	37.7531	11.6363	0.6655	0.5566	6.4845	-15.3083	6.5776	-14.8402	6.3259	-15.5547
11	$E_{M06-2X}, E_{HOMO}^{HOMO}$	0.5831	0.0726	0.6311	0.0246	0.6557	0.3457	1.0089	1.1103	0.9304	26.6599	2	0.7176	0.6666	0.7072	0.792	0.7527	0.7946	0.6254	0.0838	0.813	0.8975	0.7556	31.5565	38.2123	11.2545	0.6607	0.5602	7.2481	-14.2227	6.6043	-14.7153	6.3517	-15.3904
12	$E_{M06-L}, E_{HOMO}^{HOMO}$	0.551	0.0827	0.6076	0.0262	0.6337	0.2823	1.0406	1.1521	0.9651	24.2221	0	0.6962	0.6413	0.685	0.7758	0.7302	0.769	0.5479	0.1032	0.8266	0.9131	0.7963	33.5692	41.1502	12.1073	0.6405	0.5199	6.7844	-14.6291	7.0666	-14.2034	6.7418	-15.101
13	$E_{M06-2X}, \eta_{M06-2X}$	0.4616	0.1262	0.5584	0.0294	0.5878	0.1338	1.1039	1.2616	1.0442	19.9668	0	0.6443	0.5801	0.6312	0.7404	0.6705	0.7137	0.4312	0.145	0.9217	1.0399	0.884	37.7746	49.3395	14.1748	0.5983	0.4594	6.7489	-14.9611	6.9357	-14.347	6.3502	-15.6883
14	E_{M06-L}, η_{M06}	0.4527	0.1304	0.5533	0.0298	0.5831	0.1358	1.1102	1.272	1.0627	19.5782	0	0.6315	0.565	0.618	0.7366	0.6644	0.7031	0.4145	0.1505	0.932	1.0533	0.9087	38.2118	50.1586	14.6824	0.5971	0.4514	6.4859	-15.2957	6.7911	-14.5454	6.8381	-14.8391
15	E_{M06-L}, η_{M06-L}	0.4363	0.1327	0.5383	0.0308	0.5691	0.1038	1.1287	1.2909	1.1237	18.4869	0	0.588	0.5137	0.5729	0.7253	0.6507	0.6631	0.3501	0.1788	0.9562	1.0814	1.0061	39.4954	51.6593	16.4154	0.5858	0.4303	6.6439	-15.3211	6.6466	-14.8362	6.4947	-15.5033

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N.Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S19. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange-correlation (HFx+LYP, BLYP and B3LYP) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	R^2_{LOO}	R^2_{adj}	R^2_{adj}	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{T1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{F}^2_{int}	ΔF^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Year}	Q^2_{Year}	R^2_{Yrand}	Q^2_{Yrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{HFx+LYP}^{HOMO}$ $E_{HFx+LYP}^{HOMO}$	0.8921	0.0561	0.9333	0.0148	0.9481	0.3791	0.4667	0.6732	0.3717	63.9963	5	0.9642	0.9642	0.9671	0.9734	0.9447	0.9826	0.8862	0.0213	0.3952	0.5680	0.3145	2.1777	4.5318	0.6907	0.9528	0.7629	21.9794	-61.7079	21.7589	-70.1808	22.4744	-61.4373
2	E_{B3LYP}^{HOMO} E_{B3LYP}^{HOMO}	0.8897	0.0581	0.9329	0.0149	0.9478	0.4409	0.4683	0.6805	0.3691	63.5213	5	0.9647	0.9647	0.9676	0.9732	0.9432	0.9827	0.8879	0.0204	0.3837	0.5550	0.3065	2.1931	4.6312	0.6811	0.9521	0.7383	22.1647	-61.2524	22.0663	-68.2821	22.2077	-61.2150
3	E_{BLYP}^{HOMO} χ_{BLYP}	0.8847	0.0653	0.9357	0.0143	0.9500	0.1003	0.4584	0.6959	0.4817	66.4643	5	0.9399	0.9398	0.9448	0.9743	0.9406	0.9720	0.8238	0.0347	0.3879	0.5799	0.3941	2.1009	4.8430	1.1601	0.9574	0.7270	21.8175	-67.1722	22.0540	-67.3895	20.7748	-66.5595
4	$E_{HFx+LYP}^{HOMO}$ $\chi_{HFx+LYP}$	0.8831	0.0651	0.9335	0.0148	0.9483	0.0812	0.4661	0.7006	0.5405	64.1557	5	0.9243	0.9242	0.9304	0.9734	0.9401	0.9658	0.8105	0.0443	0.3875	0.5681	0.4707	2.1726	4.9080	1.4609	0.9558	0.7078	21.7435	-62.2515	21.4043	-70.4791	22.1623	-60.9956
5	E_{BLYP}^{HOMO} E_{BLYP}^{HOMO}	0.8814	0.0644	0.9304	0.0155	0.9458	0.4835	0.4770	0.7057	0.3781	61.1130	5	0.9630	0.9629	0.9660	0.9722	0.9387	0.9817	0.8854	0.0203	0.3848	0.5665	0.3002	2.2749	4.9797	0.7147	0.9512	0.7480	20.8031	-64.4687	21.8562	-68.5860	21.4284	-62.1607
6	E_{B3LYP}^{HOMO} χ_{B3LYP}	0.8794	0.0687	0.9333	0.0148	0.9481	0.1015	0.4669	0.7116	0.4985	63.9202	5	0.9356	0.9356	0.9408	0.9734	0.9380	0.9703	0.8163	0.0373	0.3950	0.5881	0.4116	2.1801	5.0640	1.2423	0.9545	0.7643	23.5082	-60.1483	21.5496	-70.2395	21.5255	-63.6436
7	$E_{HFx+LYP}^{HOMO}$ $\eta_{HFx+LYP}$	0.8627	0.0605	0.9013	0.0219	0.9232	0.0925	0.5679	0.7593	0.4647	42.0728	5	0.9440	0.9440	0.9486	0.9601	0.9294	0.9694	0.9431	0.0041	0.4268	0.6012	0.3796	3.2253	5.7655	1.0796	0.9241	0.5293	22.3573	-63.8950	21.4242	-67.5565	23.5924	-58.3102
8	E_{B3LYP}^{HOMO} η_{B3LYP}	0.8513	0.0628	0.8896	0.0245	0.9141	0.0957	0.6006	0.7903	0.5406	37.2526	5	0.9243	0.9242	0.9304	0.9551	0.9233	0.9583	0.9095	0.0462	0.4353	0.6074	0.4339	3.6068	6.2455	1.4612	0.9143	0.6270	23.9290	-57.5111	22.7135	-65.9103	21.6030	-65.7885
9	E_{BLYP}^{HOMO} η_{BLYP}	0.8403	0.0633	0.8760	0.0276	0.9035	0.0892	0.6365	0.8190	0.5912	32.7816	5	0.9094	0.9094	0.9168	0.9493	0.9166	0.9490	0.8675	0.0581	0.4487	0.6115	0.4868	4.0512	6.7078	1.7475	0.9057	0.5090	22.8803	-62.7192	23.5435	-67.2161	22.7790	-62.8621
10	$E_{HFx+LYP}^{HOMO}$ $\omega_{HFx+LYP}$	0.8363	0.0968	0.9140	0.0191	0.9331	-0.0438	0.5301	0.8291	0.7803	48.8159	5	0.8422	0.8421	0.8550	0.9654	0.9166	0.9342	0.7364	0.0790	0.4562	0.6839	0.6484	2.8096	6.8737	3.0443	0.9441	0.6290	21.6940	-64.7598	20.9693	-72.3987	22.7684	-61.7637
11	E_{B3LYP}^{LUMO} E_{B3LYP}^{LUMO}	0.8230	0.1040	0.9061	0.0209	0.9270	-0.0447	0.5537	0.8622	0.8297	44.4478	5	0.8216	0.8215	0.8361	0.9621	0.9090	0.9245	0.6878	0.0904	0.4537	0.6942	0.6379	3.0655	7.4344	3.4416	0.9363	0.6446	21.9277	-74.3680	20.8247	-68.6930	22.4754	-71.1257
12	E_{B3LYP}^{LUMO} E_{B3LYP}^{LUMO}	0.7862	0.1319	0.8948	0.0234	0.9182	-0.0568	0.5862	0.9475	0.8874	39.2752	5	0.7959	0.7958	0.8125	0.9573	0.8914	0.9150	0.6778	0.1000	0.4712	0.7324	0.6990	3.4362	8.9772	3.9370	0.9330	0.4793	21.7020	-70.7957	23.9076	-63.0904	21.9027	-68.1805
13	E_{B3LYP}^{HOMO} ω_{B3LYP}	0.6994	0.1926	0.8611	0.0309	0.8920	-0.0625	0.6735	1.1236	1.0298	28.9039	4	0.7252	0.7250	0.7475	0.9429	0.8509	0.8879	0.6370	0.1242	0.5435	0.8579	0.8450	4.5361	12.6237	5.3027	0.9057	0.3130	22.7737	-64.4224	21.9101	-68.6705	22.4299	-63.9915
14	$E_{HFx+LYP}^{LUMO}$ $E_{HFx+LYP}^{LUMO}$	0.6972	0.1906	0.8557	0.0321	0.8878	-0.0668	0.6866	1.1277	1.1299	27.6832	1	0.6691	0.6689	0.6960	0.9405	0.8470	0.8696	0.6139	0.1375	0.5644	0.8790	0.9486	4.7136	12.7167	6.3836	0.9033	0.4389	22.4945	-62.6439	21.8520	-70.1848	21.9754	-65.0578
15	E_{BLYP}^{HOMO} ω_{BLYP}	0.6171	0.2400	0.8164	0.0408	0.8572	-0.0575	0.7745	1.2680	1.1610	21.0035	1	0.6507	0.6504	0.6790	0.9231	0.8080	0.8540	0.5910	0.1498	0.6257	0.9798	0.9607	5.9986	16.0789	6.7400	0.8747	0.2503	21.9825	-66.6553	21.4262	-70.2615	22.0693	-64.8659

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARENS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N. Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S20. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using exchange-correlation (HFx+LYP, BLYP and B3LYP) based methods through 30% (random) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	$R^2_{Q^2_{LOO}}$	R^2_{adj}	R^2_{adj}	R^2_{adj}	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{E1}	Q^2_{E2}	Q^2_{E3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{ext}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Yest}	Q^2_{Yest}	R^2_{Yrand}	Q^2_{Yrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{HFx+LYP} \cdot E_{HOMO}^{HOMO}$	0.8851	0.0558	0.9261	0.0148	0.9409	0.3848	0.4749	0.6624	0.2453	63.7044	5	0.9875	0.9875	0.9842	0.9696	0.9407	0.9938	0.9576	0.0080	0.4089	0.5663	0.2116	2.4813	4.8261	0.2406	0.9434	0.6994	19.8221	-53.9691	19.8985	-58.0855	21.3591	-50.4427
2	$E_{BLYP} \cdot X_{BLYP}$	0.8826	0.0621	0.9308	0.0138	0.9447	0.0855	0.4596	0.6695	0.4298	68.2891	5	0.9617	0.9616	0.9516	0.9715	0.9391	0.9819	0.8710	0.0224	0.3969	0.5697	0.3362	2.3239	4.9301	0.7391	0.9499	0.7150	20.1971	-56.0847	19.7550	-58.1195	19.6207	-58.2302
3	$E_{B3LYP} \cdot E_{HOMO}^{HOMO}$	0.8824	0.0576	0.9250	0.0150	0.9400	0.4467	0.4786	0.6701	0.2270	62.6802	5	0.9893	0.9893	0.9865	0.9691	0.9389	0.9947	0.9621	0.0067	0.3970	0.5514	0.1958	2.5194	4.9389	0.2061	0.9452	0.7427	20.1310	-53.3814	20.5526	-56.5028	19.5937	-54.0735
4	$E_{HFx+LYP} \cdot X_{HFx+LYP}$	0.8822	0.0618	0.9300	0.0140	0.9440	0.0665	0.4624	0.6707	0.5261	67.4389	5	0.9426	0.9425	0.9275	0.9712	0.9392	0.9739	0.8548	0.0318	0.3917	0.5532	0.4492	2.3516	4.9482	1.1073	0.9488	0.7782	19.9449	-54.7623	19.3021	-57.3220	18.2851	-56.6318
5	$E_{B3LYP} \cdot X_{B3LYP}$	0.8782	0.0650	0.9291	0.0142	0.9433	0.0863	0.4655	0.6818	0.4590	66.4853	5	0.9563	0.9563	0.9448	0.9708	0.9370	0.9796	0.8600	0.0253	0.4014	0.5747	0.3648	2.3834	5.1139	0.8428	0.9495	0.7200	20.3673	-54.2044	20.6016	-55.7319	21.6294	-52.6924
6	$E_{BLYP} \cdot E_{HOMO}^{HOMO}$	0.8738	0.0639	0.9221	0.0156	0.9377	0.4762	0.4877	0.6940	0.2318	60.2189	5	0.9889	0.9888	0.9859	0.9679	0.9342	0.9944	0.9596	0.0066	0.4123	0.5801	0.1841	2.6160	5.2985	0.2149	0.9435	0.7206	21.1418	-51.2546	20.9238	-54.5191	20.9746	-51.8951
7	$E_{HFx+LYP} \cdot X_{HFx+LYP}$	0.8498	0.0582	0.8850	0.0230	0.9080	0.0862	0.5926	0.7572	0.3264	39.4896	5	0.9779	0.9779	0.9721	0.9518	0.9215	0.9875	0.8822	0.0140	0.4534	0.6032	0.2858	3.8628	6.3073	0.4261	0.9078	0.6691	19.6280	-57.3645	19.1549	-57.7598	19.5265	-56.9078
8	$E_{HFx+LYP} \cdot X_{HFx+LYP}$	0.8416	0.0903	0.9148	0.0170	0.9319	-0.0530	0.5101	0.7777	0.8520	54.7039	5	0.8495	0.8493	0.8099	0.9647	0.9191	0.9372	0.7556	0.0686	0.4373	0.6369	0.7376	2.8617	6.6529	2.9036	0.9402	0.6822	19.7244	-56.0226	20.7992	-56.2872	19.3957	-56.9965
9	$E_{B3LYP} \cdot X_{B3LYP}$	0.8379	0.0574	0.8691	0.0262	0.8953	0.0874	0.6322	0.7868	0.4263	34.2066	5	0.9623	0.9623	0.9524	0.9448	0.9146	0.9785	0.8313	0.0288	0.4662	0.6068	0.3403	4.3970	6.8094	0.7271	0.8956	0.4757	21.5021	-54.5061	21.7267	-51.5034	21.0012	-54.4140
10	$E_{BLYP} \cdot X_{BLYP}$	0.8301	0.0947	0.9060	0.0188	0.9248	-0.0540	0.5359	0.8055	0.8933	49.1816	5	0.8346	0.8343	0.7910	0.9609	0.9123	0.9297	0.7032	0.0784	0.4316	0.6376	0.6994	3.1589	7.1369	3.1917	0.9305	0.6301	20.2387	-59.4732	20.4010	-57.3591	19.5696	-62.9501
11	$E_{BLYP} \cdot X_{BLYP}$	0.8245	0.0592	0.8546	0.0291	0.8837	0.0804	0.6664	0.8185	0.5047	30.3885	5	0.9472	0.9471	0.9333	0.9382	0.9065	0.9692	0.7889	0.0423	0.4787	0.6107	0.4025	4.8852	7.3697	1.0189	0.8858	0.4507	21.0491	-54.0528	20.3026	-57.9467	20.1140	-54.8191
12	$E_{B3LYP} \cdot E_{HOMO}^{HOMO}$	0.7951	0.1217	0.8960	0.0208	0.9168	-0.0654	0.5636	0.8844	0.9713	44.0727	5	0.8044	0.8041	0.7529	0.9566	0.8958	0.9184	0.6891	0.0886	0.4432	0.6673	0.7978	3.4946	8.6038	3.7738	0.9269	0.6311	18.8755	-60.0352	18.6105	-60.7592	20.0680	-60.2992
13	$E_{HFx+LYP} \cdot E_{HOMO}^{HOMO}$	0.7112	0.1747	0.8573	0.0285	0.8859	-0.0748	0.6601	1.0500	1.2549	31.0498	1	0.6735	0.6731	0.5875	0.9395	0.8541	0.8717	0.6298	0.1230	0.5245	0.7907	1.1104	4.7930	12.1280	6.2993	0.8947	0.5419	20.5400	-56.4358	20.5850	-55.5373	19.9968	-58.9030
14	$E_{B3LYP} \cdot X_{B3LYP}$	0.7089	0.1819	0.8635	0.0273	0.8908	-0.0707	0.6457	1.0542	1.1402	32.6340	3	0.7305	0.7301	0.6595	0.9423	0.8556	0.8901	0.6459	0.1117	0.5089	0.7823	0.9923	4.5857	12.2256	5.2004	0.9024	0.4876	20.3042	-57.3648	20.4722	-56.7230	18.6994	-60.2581
15	$E_{BLYP} \cdot X_{BLYP}$	0.6329	0.2212	0.8176	0.0365	0.8541	-0.0659	0.7465	1.1839	1.2882	23.4074	1	0.6560	0.6555	0.5654	0.9213	0.8150	0.8569	0.6026	0.1354	0.5849	0.8891	1.1097	6.1295	15.4177	6.6379	0.8668	0.3586	19.7015	-60.8864	21.2735	-54.7731	19.1376	-60.2717

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N. Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S21. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																
		Q^2_{100}	$R^2-Q^2_{100}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Xntd}	Q^2_{Xntd}	R^2_{Yntd}	Q^2_{Yntd}
1	E_{M06}, η_{M06}	0.9376	0.0294	0.9539	0.0132	0.9671	0.0922	0.3856	0.5308	0.8451	73.4735	4	0.7584	0.7546	0.8419	0.9833	0.9681	0.8524	0.5745	0.1954	0.306	0.4597	0.6485	1.1893	2.2536	4.2849	0.9665	0.4282	27.7949	-111.7891	28.2131	-109.2969	28.3554	-106.8579
2	E_{M06-L}, η_{M06-L}	0.9338	0.0322	0.9523	0.0136	0.966	0.0992	0.3922	0.5469	0.901	70.9336	3	0.7253	0.7211	0.8203	0.9827	0.966	0.8365	0.5489	0.2102	0.312	0.4754	0.6781	1.2304	2.393	4.8705	0.9665	0.4517	27.9991	-103.9355	29.1644	-103.9607	29.0461	-98.2654
3	$E_{M06-2X}, \eta_{M06-2X}$	0.9255	0.0391	0.9503	0.0142	0.9645	0.0812	0.4004	0.5803	0.8164	67.9586	4	0.7745	0.771	0.8525	0.9819	0.9618	0.8638	0.5971	0.1827	0.3433	0.5338	0.622	1.2824	2.6937	3.9987	0.9667	0.4467	29.4742	-94.3226	29.123	-106.5453	28.8858	-95.8147
4	$E_{M06-2X}, E_{M06-2X}^{HOMO}$	0.9154	0.0488	0.9498	0.0143	0.9642	0.3813	0.4024	0.6182	0.6584	67.2372	5	0.8533	0.8511	0.9041	0.9817	0.9569	0.9154	0.7345	0.1167	0.361	0.5729	0.5509	1.2956	3.0576	2.6008	0.9669	0.2947	28.3913	-94.96	28.2176	-107.765	29.0638	-90.382
5	E_{M06}, E_{M06}^{HOMO}	0.912	0.051	0.9482	0.0148	0.963	0.4472	0.409	0.6306	0.6381	65.03	5	0.8622	0.8601	0.9099	0.9811	0.9552	0.9221	0.7564	0.1082	0.3621	0.5745	0.5332	1.338	3.181	2.4428	0.9665	0.2437	29.3185	-89.4435	29.5409	-107.8017	28.1431	-92.6082
6	$E_{M06-L}, E_{M06-L}^{HOMO}$	0.9007	0.0589	0.9435	0.0161	0.9596	0.4873	0.4271	0.6698	0.6469	59.4245	5	0.8584	0.8562	0.9074	0.9794	0.9497	0.9194	0.747	0.1118	0.3755	0.598	0.537	1.4591	3.5886	2.5106	0.9630	0.3326	29.892	-84.3877	28.2755	-110.5739	28.6256	-90.1226
7	E_{M06-2X}, X_{M06-2X}	0.8869	0.0709	0.9409	0.0169	0.9578	0.0975	0.4367	0.7148	0.6766	56.7217	5	0.8451	0.8427	0.8987	0.9784	0.9432	0.9139	0.7736	0.1141	0.3855	0.6288	0.6009	1.5257	4.0872	2.7467	0.9620	0.0473	29.9173	-84.8367	28.1887	-108.8898	28.409	-90.3705
8	E_{M06}, X_{M06}	0.8653	0.0857	0.9314	0.0196	0.951	0.0841	0.4705	0.78	0.6957	48.5188	5	0.8362	0.8337	0.8929	0.9749	0.9327	0.9107	0.7692	0.1183	0.4241	0.6922	0.6077	1.771	4.8678	2.9044	0.9568	-0.1088	28.285	-87.7112	29.086	-105.774	28.0153	-87.1623
9	E_{M06-L}, X_{M06-L}	0.8644	0.0861	0.9306	0.0198	0.9504	0.0934	0.4732	0.7828	0.6626	47.9485	5	0.8514	0.8491	0.9028	0.9746	0.9325	0.9176	0.7791	0.1103	0.4172	0.6784	0.576	1.791	4.9021	2.6344	0.9569	0.0372	28.7003	-86.8684	29.0013	-107.4807	28.7019	-85.8066
10	$E_{M06-L}, E_{M06-L}^{HOMO}$	0.7279	0.2013	0.9009	0.0283	0.9292	-0.0166	0.5655	1.1087	0.873	32.8217	5	0.7421	0.7381	0.8313	0.9633	0.8789	0.8602	0.6618	0.0803	0.4757	0.8531	0.7203	2.558	9.8346	4.5731	0.9414	-0.7682	27.9867	-100.0206	29.8237	-104.964	27.5712	-98.7785
11	$E_{M06-2X}, \omega_{M06-2X}$	0.7232	0.2113	0.9083	0.0262	0.9345	-0.0224	0.544	1.1183	0.9597	35.6573	1	0.6884	0.6835	0.7961	0.9661	0.8781	0.834	0.6064	0.0805	0.4353	0.8082	0.8368	2.3679	10.0041	5.5256	0.9503	-0.7994	28.4668	-96.7516	29.2101	-104.6621	29.1888	-93.7139
12	E_{M06}, ω_{M06}	0.7146	0.2	0.8805	0.0342	0.9146	-0.0158	0.621	1.1355	1.1443	26.7821	1	0.5569	0.55	0.7101	0.9554	0.8719	0.7491	0.4609	0.2018	0.5243	0.937	0.9638	3.0856	10.3153	7.8571	0.9247	-0.7434	28.4578	-113.1536	29.5066	-103.8563	28.8376	-111.896
13	$E_{M06-L}, \omega_{M06-L}$	0.6956	0.2123	0.871	0.0368	0.9079	-0.0152	0.6451	1.1727	1.2723	24.642	0	0.4523	0.4437	0.6417	0.9517	0.8692	0.6739	0.3532	0.25	0.5378	0.9653	1.0343	3.3289	11.0024	9.7132	0.9141	-0.9541	29.4002	-132.8891	29.0162	-108.4973	28.6865	-134.2754
14	E_{M06}, E_{M06}^{LUMO}	0.6909	0.2308	0.8904	0.0313	0.9217	-0.0185	0.5948	1.1817	1.005	29.42	1	0.6582	0.6529	0.7764	0.9592	0.8644	0.8114	0.5658	0.1462	0.5068	0.9364	0.8591	2.8306	11.1708	6.0607	0.9357	-1.1518	28.0177	-106.0502	27.4887	-113.4558	29.3138	-104.1459
15	$E_{M06-2X}, E_{M06-2X}^{LUMO}$	0.607	0.319	0.8963	0.0296	0.926	-0.0247	0.5784	1.3325	1.0272	31.2611	1	0.643	0.6375	0.7665	0.9616	0.8405	0.8049	0.5513	0.0901	0.472	0.9372	0.8735	2.6762	14.2045	6.3304	0.9429	-0.9509	29.6514	-103.5237	30.5324	-101.0151	26.6299	-110.496

Models are arranged according to Q^2_{100} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012.

(<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S22. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using meta exchange-correlation functionals (M06, M06-L and M06-2X) based methods through 30% (random) splitting method.

Model	Variables	N.																																
		Q^2_{100}	$R^2-Q^2_{100}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	$RMSE_{TR}$	$RMSE_{CV}$	$RMSE_{EXT}$	F	$ExtLOK$	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC_{TR}	CCC_{CV}	CCC_{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE_{TR}	MAE_{CV}	MAE_{EXT}	RSS_{TR}	$PRESS_{CV}$	$PRESS_{EXT}$	R^2_{LMO}	Q^2_{LMO}	R^2_{Ytest}	Q^2_{Ytest}	R^2_{Xtest}	Q^2_{Xtest}	R^2_{Ytest}	Q^2_{Ytest}
1	E_{M06-L}, X_{M06-L}	0.8029	0.1097	0.8908	0.0218	0.9126	-0.0994	0.5821	0.8743	0.515	41.7915	5	0.9428	0.9386	0.9316	0.9543	0.9051	0.971	0.7986	0.0278	0.478	0.6885	0.4896	3.7267	8.4085	1.061	0.9225	0.4249	20.7231	-57.8864	19.7506	-57.2044	19.9462	-60.7468
2	$E_{M06-L}, E_{M06-L}^{HOMO}$	0.7966	0.117	0.8919	0.0216	0.9135	0.2004	0.5791	0.8883	0.31	42.2546	5	0.9793	0.9777	0.9752	0.9548	0.9015	0.9888	0.9073	0.0078	0.5292	0.7718	0.281	3.6894	8.679	0.3845	0.9236	0.6483	20.2452	-57.6463	19.5393	-57.3815	19.6521	-57.8741
3	$E_{M06-L}, E_{M06-L}^{HOMO}, E_{M06-L}^{LUMO}$	0.7952	0.1174	0.8907	0.0219	0.9126	0.1423	0.5824	0.8913	0.3531	41.744	5	0.9731	0.9711	0.9678	0.9543	0.9007	0.9855	0.8954	0.0119	0.5286	0.7682	0.295	3.7306	8.7384	0.4988	0.9238	0.6750	19.8096	-57.7607	20.4369	-56.3584	19.124	-60.238
4	E_{M06-2X}, X_{M06-2X}	0.7907	0.1185	0.8865	0.0227	0.9092	-0.0934	0.5934	0.9009	0.9009	40.06	5	0.9542	0.9508	0.9453	0.9524	0.899	0.9758	0.8378	0.0179	0.499	0.7225	0.452	3.8732	8.9288	0.8489	0.9196	0.4544	20.0443	-58.6461	20.2271	-57.4871	19.5866	-59.1815
5	$E_{M06-2X}, E_{M06-2X}^{HOMO}$	0.7854	0.1248	0.8876	0.0225	0.9101	0.2822	0.5904	0.9124	0.2758	40.5031	5	0.9836	0.9824	0.9804	0.9529	0.8965	0.991	0.9216	0.0048	0.5441	0.7967	0.2611	3.8346	9.1571	0.3043	0.9199	0.5766	20.2852	-56.762	20.2144	-56.5654	19.9734	-57.6441
6	$E_{M06-2X}, E_{M06-2X}^{HOMO}, E_{M06-2X}^{LUMO}$	0.7813	0.1209	0.8777	0.0245	0.9022	-0.0981	0.6159	0.921	0.4729	36.8954	5	0.9518	0.9482	0.9423	0.9486	0.894	0.9753	0.8178	0.0233	0.4927	0.7082	0.4476	4.1729	9.3311	0.8947	0.9132	0.4133	20.2397	-59.3655	20.3822	-57.6207	20.1287	-59.5271
7	$E_{M06-2X}, \omega_{M06-2X}$	0.7121	0.1341	0.8078	0.0384	0.8462	-0.1409	0.7722	1.0566	0.7292	22.0146	5	0.8853	0.8768	0.8629	0.9167	0.8547	0.9391	0.7623	0.0483	0.5371	0.7703	0.6395	6.5598	12.2811	2.127	0.8577	0.0407	20.0714	-61.6787	20.5407	-54.9415	20.3017	-61.3724
8	$E_{M06-2X}, \eta_{M06-2X}$	0.705	0.1545	0.8244	0.0351	0.8595	0.1217	0.7382	1.0697	0.3096	24.4702	5	0.9793	0.9778	0.9753	0.9244	0.8565	0.9875	0.8573	0.0171	0.6652	0.9419	0.2801	5.994	12.5874	0.3834	0.8729	0.3548	20.2079	-56.909	20.4296	-55.1677	19.4959	-58.4826
9	E_{M06}, η_{M06}	0.6983	0.1576	0.8199	0.036	0.8559	0.1282	0.7475	1.0817	0.3832	23.7615	5	0.9683	0.966	0.9621	0.9224	0.8536	0.9802	0.8243	0.0261	0.6705	0.95	0.3381	6.1471	12.8713	0.5873	0.8679	0.4637	20.5301	-55.4222	18.9807	-59.4511	20.5426	-56.2351
10	$E_{M06-L}, E_{M06-L}^{LUMO}$	0.6791	0.157	0.7952	0.041	0.8361	-0.1355	0.7972	1.1156	0.6062	20.4115	5	0.9207	0.9149	0.9053	0.9108	0.841	0.9583	0.7895	0.0309	0.5808	0.8088	0.5794	6.9906	13.6894	1.4697	0.8481	0.3388	19.0538	-70.3668	20.0604	-58.4087	21.0157	-65.4341
11	E_{M06-L}, η_{M06-L}	0.6546	0.1818	0.7955	0.0409	0.8364	0.1379	0.7965	1.1574	0.5017	20.4533	5	0.9457	0.9417	0.9351	0.9109	0.833	0.9653	0.7744	0.0482	0.7001	0.994	0.4346	6.9787	14.7354	1.007	0.8519	0.3739	19.4985	-61.362	19.7692	-56.9152	20.278	-58.6698
12	E_{M06}, E_{M06}^{LUMO}	0.5557	0.2372	0.7411	0.0518	0.7929	-0.1318	0.8963	1.3127	0.5836	15.3127	5	0.9265	0.9211	0.9122	0.8845	0.7745	0.9591	0.8453	0.0274	0.6606	0.9816	0.533	8.8362	18.9564	1.3625	0.8152	0.0829	20.5617	-63.9972	19.6951	-58.4899	20.5158	-65.3048
13	$E_{M06-2X}, E_{M06-2X}^{LUMO}$	0.5184	0.2645	0.7286	0.0543	0.7829	-0.1317	0.9176	1.3668	0.732	14.4237	5	0.8844	0.8759	0.8619	0.8782	0.7548	0.9321	0.8429	0.0383	0.6927	1.0432	0.683	9.2626	20.5485	2.1431	0.8016	-0.0577	20.0308	-63.4254	19.6596	-57.2943	20.2048	-64.3493
14	E_{M06}, ω_{M06}	0.5123	0.2562	0.7107	0.0579	0.7686	-0.1303	0.9474	1.3753	0.6217	13.2826	5	0.9166	0.9105	0.9003	0.8691	0.7503	0.953	0.8555	0.0396	0.7197	1.0622	0.6105	9.8742	20.806	1.5461	0.7869	-0.2663	20.0267	-62.0108	19.7598	-57.9131	19.9072	-61.7964
15	$E_{M06-L}, \omega_{M06-L}$	0.3121	0.4052	0.6466	0.0707	0.7173	-0.1208	1.0471	1.6333	0.6763	10.1494	5	0.9013	0.8941	0.8821	0.8354	0.6628	0.9394	0.8856	0.0487	0.8545	1.2895	0.6522	12.0607	29.3461	1.8297	0.7490	-0.5231	19.9278	-59.3158	20.769	-55.7784	20.516	-58.0956

Models are arranged according to Q^2_{100} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecolox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N. ExtLOK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S23. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_m	Δr^2_m	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Yvr}	Q^2_{Yvr}	R^2_{Xmdl}	Q^2_{Xmdl}	R^2_{Ymdl}	Q^2_{Ymdl}
1	$E_{CORR(B3LYP)}^{HOMO}$	0.7316	0.0638	0.7759	0.0195	0.7954	0.2918	0.8486	0.9720	0.6796	40.8131	5	0.8194	0.8164	0.8687	0.8860	0.8537	0.8979	0.7899	0.0741	0.6628	0.7601	0.5374	17.2837	22.6726	10.1623	0.7983	0.7184	8.5818	-19.8806	8.8958	-19.6206	8.5980	-20.3638
2	$E_{CORR(B3LYP)}^{HOMO}$	0.7140	0.0662	0.7593	0.0209	0.7802	0.0759	0.8795	1.0033	0.8843	37.2770	1	0.6942	0.6892	0.7778	0.8765	0.8433	0.8465	0.6090	0.0998	0.6903	0.7933	0.7837	18.5629	24.1581	17.2038	0.7875	0.6954	8.7341	-20.7888	9.2896	-19.1411	8.7284	-20.5103
3	$E_{CORR(B3LYP)}^{HOMO}$	0.7074	0.0691	0.7552	0.0213	0.7764	0.2506	0.8870	1.0148	0.7811	36.4691	4	0.7614	0.7575	0.8266	0.8742	0.8391	0.8456	0.6957	0.1080	0.6886	0.7902	0.6194	18.8822	24.7146	13.4224	0.7838	0.6829	8.6179	-20.1660	8.7765	-19.7794	8.9349	-19.7431
4	$E_{CORR(B3LYP)}^{HOMO}$	0.6148	0.0941	0.6812	0.0277	0.7090	0.2006	1.0121	1.1643	0.8699	25.5776	3	0.7041	0.6992	0.7850	0.8297	0.7825	0.7957	0.6789	0.1158	0.8110	0.9302	0.6906	24.5826	32.5320	16.6488	0.7217	0.5871	8.7722	-20.3955	8.8063	-19.4989	8.3950	-20.6244
5	$E_{CORR(HFX+LYP)}^{HOMO}$	0.5811	0.1030	0.6540	0.0301	0.6841	0.2542	1.0545	1.2142	0.8869	22.7332	2	0.6924	0.6874	0.7765	0.8124	0.7608	0.8100	0.6760	0.0995	0.8953	1.0283	0.7326	26.6866	35.3827	17.3041	0.6978	0.5577	8.5996	-20.1505	8.6091	-19.9511	8.9524	-19.3428
6	$E_{CORR(BLYP)}^{HOMO}$	0.5613	0.1018	0.6310	0.0321	0.6631	0.2905	1.0889	1.2426	1.0263	20.6632	1	0.5881	0.5813	0.7007	0.7974	0.7454	0.7027	0.5875	0.0836	0.9167	1.0454	0.8590	28.4592	37.0588	23.1742	0.6749	0.3861	8.7603	-20.7541	8.7057	-19.9187	8.5276	-21.3903
7	$E_{CORR(BLYP)}^{HOMO}$	0.5598	0.1032	0.6310	0.0321	0.6631	0.3283	1.0889	1.2446	1.0114	20.6643	1	0.6000	0.5934	0.7093	0.7974	0.7445	0.7104	0.5989	0.1029	0.9138	1.0445	0.8385	28.4582	37.1781	22.5061	0.6737	0.5409	9.0654	-20.4579	8.5227	-20.1933	8.8310	-21.1491
8	$E_{CORR(BLYP)}^{HOMO}$	0.5570	0.1053	0.6302	0.0322	0.6624	0.3680	1.0900	1.2486	1.0002	20.6001	1	0.6088	0.6024	0.7158	0.7969	0.7425	0.7170	0.6057	0.1143	0.9169	1.0527	0.8168	28.5170	37.4144	22.0082	0.6747	0.5176	8.7120	-21.2602	9.1549	-19.2641	8.4688	-21.1907
9	$E_{CORR(HFX+LYP)}^{HOMO}$	0.5504	0.1192	0.6381	0.0315	0.6696	0.0980	1.0783	1.2579	0.9148	21.2803	2	0.6728	0.6674	0.7622	0.8021	0.7413	0.7855	0.6690	0.0379	0.8921	1.0368	0.7653	27.9066	37.9781	18.4107	0.6846	0.5373	8.6560	-20.1316	8.6377	-19.7756	8.6007	-20.4133
10	$E_{CORR(BLYP)}^{LUMO}$	0.5354	0.1162	0.6184	0.0332	0.6516	0.0777	1.1073	1.2787	1.0257	19.6386	0	0.5886	0.5818	0.7011	0.7891	0.7306	0.7090	0.5923	0.0329	0.9385	1.0804	0.8319	29.4267	39.2400	23.1462	0.6662	0.5254	8.4540	-20.8180	9.1036	-19.1342	8.8774	-19.8428
11	$E_{CORR(HFX+LYP)}^{LUMO}$	0.5315	0.1260	0.6249	0.0326	0.6575	0.2027	1.0978	1.2840	1.0298	20.1608	0	0.5854	0.5785	0.6987	0.7934	0.7325	0.7101	0.5982	0.0067	0.9229	1.0718	0.8315	28.9255	39.5683	23.3294	0.6733	0.5461	8.6228	-19.8037	8.7390	-19.6208	8.7043	-20.0552
12	$E_{CORR(HFX+LYP)}^{LUMO}$	0.5299	0.1298	0.6272	0.0324	0.6596	0.1022	1.0945	1.2863	0.9869	20.3487	0	0.6192	0.6129	0.7232	0.7949	0.7300	0.7332	0.6128	0.0553	0.9256	1.0808	0.8134	28.7494	39.7093	21.4278	0.6779	0.5120	8.6127	-20.4149	8.2008	-20.5684	8.8164	-19.9156
13	$E_{CORR(B3LYP)}^{LUMO}$	0.4928	0.1407	0.5987	0.0349	0.6336	0.0198	1.1356	1.3360	1.0909	18.1550	0	0.5347	0.5270	0.6618	0.7757	0.6984	0.6598	0.5239	0.0790	0.9535	1.1156	0.8731	30.9503	42.8382	26.1821	0.6513	0.4949	8.2016	-21.3063	8.7201	-19.9390	8.9866	-20.4215
14*	$E_{CORR(BLYP)}^{HOMO}$	0.6948	0.0736	0.7412	0.0272	0.7684	0.2315	0.7479	0.8587	0.9434	28.2024	0	0.4828	0.4820	0.6316	0.8690	0.8315	0.7161	0.4592	0.0476	0.6525	0.7582	0.7615	11.1881	14.7457	16.0193	0.7702	0.6389	10.5104	-25.7058	11.1067	-23.6962	10.6530	-25.2916
15*	$E_{CORR(HFX+LYP)}^{HOMO}$	0.6213	0.1061	0.6970	0.0303	0.7273	0.1896	0.9478	1.1170	0.9486	24.0054	1	0.6616	0.6616	0.7269	0.8421	0.7833	0.7993	0.5223	0.1645	0.7354	0.8645	0.7866	18.8643	26.2012	17.9951	0.7345	0.5720	10.0636	-23.0298	10.1334	-23.2785	9.5411	-23.6776

*Computed using data set represented with (c) and (b) respectively in Supporting Information Table S3.

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N. Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S24. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through 30% (random) splitting method.

Model	Variables	Q^2_{LOO}	R^2_{LOO}	R^2_{adj}	R^2_{adj}	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	N	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{F}^2_{int}	ΔF^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Yext}	Q^2_{Yext}	R^2_{Yext}	Q^2_{Yext}	R^2_{Yext}	Q^2_{Yext}
1	$E_{CORR(B3LYP)} \cdot \omega_{CORR(B3LYP)}$	0.7300	0.0423	0.7576	0.0147	0.7723	0.4161	0.8490	0.9245	0.5696	52.5632	5	0.8574	0.8463	0.8975	0.8715	0.8495	0.9117	0.8230	0.0489	0.7074	0.7740	0.4538	24.5056	29.0577	3.8934	0.7767	0.7037	5.8120	-13.7357	6.1898	-13.0184	6.2578	-13.0349	
2	$E_{CORR(B3LYP)} \cdot \eta_{CORR(B3LYP)}$	0.7285	0.0522	0.7666	0.0141	0.7807	0.1350	0.8331	0.9270	0.8462	55.1825	1	0.6852	0.6608	0.7737	0.8769	0.8500	0.7921	0.5166	0.1456	0.7054	0.7806	0.7110	23.5975	29.2155	8.5931	0.7865	0.7093	6.0928	-13.6483	6.1851	-13.0131	6.0133	-13.4949	
3	$E_{CORR(B3LYP)} \cdot E^{HOMO}_{CORR(B3LYP)}$	0.7269	0.0524	0.7650	0.0142	0.7793	0.2428	0.8358	0.9298	0.6599	54.7232	5	0.8085	0.7937	0.8624	0.8759	0.8493	0.8739	0.6864	0.0880	0.6987	0.7739	0.5388	23.7518	29.3930	5.2259	0.7819	0.7144	6.0064	-13.7885	6.2754	-12.9220	5.7498	-13.8372	
4	$E_{CORR(B3LYP)} \cdot \chi_{CORR(B3LYP)}$	0.6892	0.0502	0.7226	0.0168	0.7394	0.3932	0.9082	0.9918	0.6894	43.9734	5	0.7911	0.7748	0.8498	0.8502	0.8237	0.8616	0.7337	0.0815	0.7519	0.8230	0.5025	28.0449	33.4434	5.7031	0.7411	0.6684	6.3193	-13.0769	6.1666	-12.9331	6.5960	-12.6326	
5	$E_{CORR(BLYP)} \cdot \chi_{CORR(BLYP)}$	0.6755	0.0506	0.7085	0.0177	0.7261	0.4132	0.9310	1.0134	0.8213	41.1001	2	0.7034	0.6804	0.7868	0.8414	0.8140	0.7893	0.6028	0.1161	0.7655	0.8373	0.6626	29.4686	34.9151	8.0954	0.7276	0.6541	6.0782	-13.7373	6.2074	-13.0593	5.7923	-14.8742	
6	$E_{CORR(BLYP)} \cdot \eta_{CORR(BLYP)}$	0.6755	0.0480	0.7056	0.0178	0.7235	0.3482	0.9355	1.0135	0.8506	40.5513	1	0.6819	0.6572	0.7714	0.8395	0.8135	0.7705	0.6133	0.1132	0.7713	0.8390	0.7241	29.7572	34.9238	8.6831	0.7262	0.5960	5.8767	-14.3557	6.2288	-12.8752	6.1137	-14.3599	
7	$E_{CORR(BLYP)} \cdot E^{HOMO}_{CORR(BLYP)}$	0.6748	0.0493	0.7063	0.0178	0.7241	0.3775	0.9344	1.0145	0.8271	40.6856	1	0.6993	0.6760	0.7839	0.8400	0.8133	0.7833	0.6153	0.1128	0.7701	0.8403	0.6860	29.6860	34.9897	8.2082	0.7241	0.6025	5.9723	-14.5012	6.0097	-13.2461	6.0453	-13.9386	
8	$E_{CORR(HFX+LYP)} \cdot \eta_{CORR(HFX+LYP)}$	0.6746	0.0551	0.7123	0.0174	0.7297	0.2939	0.9249	1.0148	0.7243	41.8418	4	0.7693	0.7514	0.8342	0.8437	0.8145	0.8428	0.7305	0.0614	0.7938	0.8716	0.5633	29.0875	35.0146	6.2960	0.7310	0.6591	5.9475	-13.3280	6.3815	-12.7102	6.1515	-13.2789	
9	$E_{CORR(HFX+LYP)} \cdot \chi_{CORR(HFX+LYP)}$	0.6604	0.0629	0.7055	0.0179	0.7233	0.3913	0.9358	1.0367	0.7951	40.5177	4	0.7220	0.7005	0.8002	0.8394	0.8046	0.8175	0.6626	0.0137	0.7752	0.8547	0.6367	29.7750	36.5435	7.5869	0.7271	0.6611	6.0901	-13.2748	6.1081	-13.2016	5.8029	-13.6337	
10	$E_{CORR(HFX+LYP)} \cdot \eta_{CORR(HFX+LYP)}$	0.6595	0.0555	0.6966	0.0184	0.7150	0.2005	0.9497	1.0381	0.8081	38.8916	2	0.7129	0.6907	0.7937	0.8338	0.8042	0.7959	0.6426	0.1059	0.8086	0.8866	0.6448	30.6651	36.6404	7.8356	0.7217	0.6253	6.2968	-13.0442	6.1381	-13.0603	5.8093	-13.5624	
11	$E_{CORR(HFX+LYP)} \cdot E^{LUMO}_{CORR(HFX+LYP)}$	0.6545	0.0624	0.6986	0.0183	0.7169	0.2128	0.9466	1.0457	0.8553	39.2459	1	0.6784	0.6534	0.7689	0.8351	0.8007	0.7868	0.6044	0.0431	0.7734	0.8536	0.6889	30.4667	37.1805	8.7786	0.7218	0.6418	6.0580	-13.5194	5.6578	-13.6882	5.9416	-13.4927	
12	$E_{CORR(BLYP)} \cdot E^{LUMO}_{CORR(BLYP)}$	0.6469	0.0627	0.6909	0.0187	0.7097	0.1624	0.9586	1.0571	0.9293	37.8858	1	0.6204	0.5909	0.7272	0.8302	0.7956	0.7566	0.5318	0.0841	0.7960	0.8753	0.7793	31.2429	37.9922	10.3624	0.7145	0.6371	6.3975	-12.7487	6.2690	-13.1105	6.1649	-13.1672	
13	$E_{CORR(B3LYP)} \cdot E^{LUMO}_{CORR(B3LYP)}$	0.6030	0.0936	0.6770	0.0196	0.6966	0.0659	0.9800	1.1209	1.0021	35.5805	0	0.5585	0.5243	0.6827	0.8211	0.7683	0.6961	0.4339	0.1528	0.8235	0.9261	0.8117	32.6529	42.7216	12.0501	0.7078	0.6061	5.9775	-13.6795	6.0562	-13.3007	5.6018	-14.7448	
14*	$E_{CORR(BLYP)} \cdot \omega_{CORR(BLYP)}$	0.6766	0.0585	0.7120	0.0230	0.7351	0.0959	0.7445	0.8226	1.0635	31.9059	0	0.4638	0.4588	0.4594	0.8473	0.8162	0.5607	0.4721	0.1355	0.6354	0.7097	0.8428	14.4105	17.5914	12.4405	0.7351	0.6273	7.9407	-18.9697	8.4925	-17.3485	7.5899	-18.8297	
15*	$E_{CORR(HFX+LYP)} \cdot \omega_{CORR(HFX+LYP)}$	0.6627	0.0640	0.7057	0.0210	0.7267	0.2099	0.8787	0.9762	1.0645	34.5727	0	0.6648	0.6607	0.5989	0.8417	0.8055	0.7905	0.5912	0.1370	0.6979	0.7778	0.9469	22.3911	27.6362	13.5990	0.7287	0.6295	7.2501	-15.6239	7.1459	-15.8096	7.5236	-15.5867	

*Computed using data set represented with (c) and (b) respectively in Supporting Information Table S3.

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S25. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N.																																
		Q^2_{LOO}	R^2	Q^2_{LOO}	R^2_{adj}	R^2	R^2_{adj}	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{E1}	Q^2_{E2}	Q^2_{E3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{CORR(HFX+LYP)}$, E_{LMO}	0.9857	0.006	0.9884	0.0033	0.9917	0.1925	0.1967	0.2588	0.8696	300.0248	5	0.8305	0.8174	0.8385	0.9959	0.9928	0.9236	0.834	0.0702	0.149	0.2045	0.7153	0.3095	0.5357	3.7814	0.9917	0.6673	30.568	-90.9253	28.6292	-114.2375	26.0326	-109.1877
2	$E_{CORR(HFX+LYP)}$, $E_{CORR(HFX+LYP)}$	0.969	0.0178	0.9815	0.0053	0.9868	0.3371	0.2485	0.3808	0.6743	187.0138	5	0.8981	0.8902	0.9029	0.9934	0.9845	0.9481	0.8655	0.0556	0.1921	0.3014	0.5028	0.4941	1.16	2.2737	0.9881	0.6725	31.3447	-86.6643	29.8416	-102.8297	29.9057	-93.7522
3	$E_{CORR(HFX+LYP)}$, $H_{CORR(HFX+LYP)}$	0.9433	0.0255	0.9564	0.0125	0.9688	0.1122	0.382	0.5152	1.1211	77.7319	4	0.7184	0.6965	0.7316	0.9842	0.9717	0.8754	0.757	0.024	0.3003	0.4295	0.9603	1.1672	2.1235	6.2839	0.9682	0.4754	29.0884	-104.0019	28.1578	-112.7071	27.9369	-108.8094
4	$E_{CORR(B3LYP)}$, $E_{CORR(B3LYP)}$	0.9418	0.0333	0.9652	0.01	0.9751	0.1537	0.3413	0.522	0.4817	97.9877	5	0.948	0.944	0.9504	0.9874	0.9704	0.9708	0.8963	0.0363	0.2944	0.4691	0.3333	0.9319	2.1801	1.1602	0.9768	0.4862	28.8528	-102.3972	30.4638	-96.361	28.4378	-104.8297
5	$E_{CORR(HFX+LYP)}$, $E_{CORR(HFX+LYP)}$, E_{HOMO}	0.9263	0.0465	0.9619	0.0109	0.9728	0.3776	0.3567	0.5875	0.6179	89.4806	5	0.9144	0.9078	0.9185	0.9862	0.9631	0.9556	0.8738	0.0512	0.3036	0.5149	0.4609	1.0181	2.7617	1.9091	0.9764	0.3271	29.4168	-98.4941	29.0534	-105.5568	29.2124	-100.887
6	$E_{CORR(B3LYP)}$, $\omega_{CORR(B3LYP)}$	0.9235	0.0485	0.9608	0.0112	0.972	0.4051	0.3621	0.5986	0.4235	86.7637	5	0.9598	0.9567	0.9617	0.9858	0.9617	0.9782	0.9267	0.027	0.3116	0.5309	0.3087	1.0491	2.8669	0.8967	0.9754	0.1582	28.5195	-100.3837	29.2347	-107.8823	27.8915	-103.8511
7	$E_{CORR(B3LYP)}$, $X_{CORR(B3LYP)}$	0.9017	0.0679	0.9573	0.0122	0.9695	0.1141	0.3777	0.6786	0.6948	79.5624	5	0.8918	0.8834	0.8969	0.9845	0.9495	0.9475	0.8864	0.0064	0.3323	0.6047	0.5562	1.1412	3.684	2.4139	0.9743	0.2572	28.0261	-178.0529	29.0357	-106.6961	30.4532	-173.2299
8	$E_{CORR(B3LYP)}$, $E_{CORR(B3LYP)}$, E_{HOMO}	0.8987	0.0693	0.9551	0.0128	0.9679	0.2028	0.3874	0.6887	0.7105	75.4813	5	0.8869	0.8781	0.8922	0.9837	0.9479	0.9434	0.8627	0.0584	0.3559	0.6309	0.5702	1.2009	3.7949	2.5241	0.9727	0.2441	29.9875	-122.7008	28.5371	-105.595	29.9511	-128.4131
9	$E_{CORR(B3LYP)}$, $X_{CORR(B3LYP)}$	0.8922	0.0677	0.9439	0.016	0.9599	0.4292	0.4332	0.7105	0.5993	59.8744	5	0.9195	0.9133	0.9233	0.9795	0.9451	0.9507	0.8919	0.0462	0.3467	0.5743	0.5298	1.5014	4.0384	1.7958	0.9659	0.0786	29.5733	-92.1222	27.6814	-106.7811	29.5312	-96.0842
10	$E_{CORR(B3LYP)}$, E_{LMO} , $E_{CORR(B3LYP)}$	0.8744	0.0887	0.9485	0.0147	0.9632	0.0589	0.4151	0.7667	0.6419	65.4247	5	0.9077	0.9005	0.912	0.9813	0.9357	0.9553	0.9051	0.0075	0.3798	0.6793	0.5335	1.3787	4.703	2.0599	0.9688	0.0210	28.7581	-113.3556	29.682	-95.7111	28.6299	-120.5318
11	$E_{CORR(B3LYP)}$, $H_{CORR(B3LYP)}$	0.8549	0.1049	0.9436	0.0161	0.9597	0.087	0.4343	0.8244	0.7223	59.5691	5	0.8831	0.874	0.8886	0.9794	0.9286	0.9411	0.8321	0.0593	0.3995	0.6991	0.5685	1.5087	5.437	2.6083	0.9649	0.0658	26.7928	-100.2076	28.451	-108.2405	30.1534	-89.2488
12	$E_{CORR(B3LYP)}$, E_{LMO} , $E_{CORR(B3LYP)}$	0.8514	0.0777	0.9007	0.0284	0.9291	0.0695	0.5762	0.8342	0.8242	32.7528	5	0.8478	0.836	0.8549	0.9632	0.9242	0.9193	0.7774	0.0885	0.4695	0.7105	0.5564	2.6564	5.5676	3.3966	0.9305	-0.2365	28.5725	-93.5056	29.2535	-101.2577	29.4615	-86.6797
13	$E_{CORR(B3LYP)}$, $H_{CORR(B3LYP)}$	0.8419	0.1027	0.9224	0.0222	0.9446	0.0223	0.5094	0.8603	0.6913	42.6178	5	0.8929	0.8846	0.8979	0.9715	0.9215	0.9434	0.8404	0.0642	0.4344	0.7419	0.497	2.0756	5.9212	2.3897	0.9488	-0.1774	27.8207	-96.2285	27.5899	-119.1871	28.3649	-93.9013
14**	$E_{CORR(HFX+LYP)}$, $\omega_{CORR(HFX+LYP)}$	0.7610	0.1867	0.9303	0.0174	0.9477	0.0998	0.4869	1.0410	0.9304	54.3871	3	0.7027	0.7026	0.8091	0.9732	0.8728	0.8178	0.5004	0.2375	0.4152	0.7658	0.6919	2.1333	9.7535	5.1935	0.9575	-0.9896	24.5289	-91.8275	25.6803	-80.3081	25.1231	-90.1154
15**	$E_{CORR(B3LYP)}$, $\omega_{CORR(B3LYP)}$	0.7350	0.1805	0.8732	0.0423	0.9155	0.1149	0.5588	0.9895	0.8160	21.6589	1	0.5787	0.5754	0.8198	0.9559	0.8838	0.7169	0.6378	0.1582	0.5075	0.9460	0.6441	2.1861	6.8536	3.3292	0.9129	-17.3882	32.9202	-204.3347	33.6834	-155.1932	31.6185	-201.5612

**Computed using data set represented with (d) and (e) respectively in Supporting Information Table S4.

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S26. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(HFX+LYP), CORR(BLYP), CORR(B3LYP)) based methods through 30% (random) splitting method.

Model	Variables	Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	$RMSE_{TR}$	$RMSE_{CV}$	$RMSE_{EXT}$	F	N_{ExtOK}	Q^2_{E1}	Q^2_{E2}	Q^2_{E3}	CCC_{TR}	CCC_{CV}	CCC_{EXT}	\bar{F}_{int}	ΔF_{int}	MAE_{TR}	MAE_{CV}	MAE_{EXT}	RSS_{TR}	$PRESS_{CV}$	$PRESS_{EXT}$	R^2_{LMO}	Q^2_{LMO}	R^2_{YSTR}	Q^2_{YSTR}	R^2_{Xrand}	Q^2_{Xrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{CORR(HFX+LYP)} / \chi_{CORR(HFX+LYP)}$	0.8736	0.0569	0.9151	0.0154	0.9305	0.4397	0.5149	0.6944	0.7580	60.2690	5	0.8808	0.8807	0.8494	0.9640	0.9241	0.9249	0.7162	0.0843	0.4128	0.5658	0.5025	3.1815	5.7858	2.2984	0.9335	0.6838	17.6814	-49.8828	17.5334	-50.6731	18.9238	-47.8202
2	$E_{CORR(HFX+LYP)} / E_{LUMO_{CORR(HFX+LYP)}}$	0.8562	0.0687	0.9082	0.0167	0.9249	0.3625	0.5353	0.7406	0.7399	55.4195	5	0.8865	0.8864	0.8565	0.9610	0.9278	0.9382	0.7201	0.0803	0.4548	0.6264	0.4428	3.4390	6.5826	2.1896	0.9275	0.6527	17.8824	-48.5402	17.7493	-49.9506	18.1734	-49.9844
3	$E_{CORR(B3LYP)} / \omega_{CORR(B3LYP)}$	0.8472	0.0644	0.8919	0.0196	0.9116	0.2910	0.5808	0.7636	0.6637	46.4003	5	0.9086	0.9086	0.8845	0.9538	0.9214	0.9517	0.7748	0.0609	0.4535	0.6037	0.4006	4.0483	6.9971	1.7623	0.9138	0.6126	17.3479	-50.7034	18.5792	-49.5018	18.3787	-49.7993
4	$E_{CORR(HFX+LYP)} / \eta_{CORR(HFX+LYP)}$	0.8424	0.0617	0.8828	0.0213	0.9041	0.1990	0.6048	0.7755	0.8583	42.4461	5	0.8472	0.8471	0.8069	0.9497	0.9211	0.9217	0.7318	0.0912	0.4896	0.6551	0.5769	4.3893	7.2160	2.9467	0.9039	0.5415	17.0629	-57.3638	17.1314	-51.2724	19.0504	-53.6329
5	$E_{CORR(B3LYP)} / \eta_{CORR(B3LYP)}$	0.8420	0.0859	0.9119	0.0160	0.9279	0.0831	0.5245	0.7764	0.9447	57.9217	5	0.8149	0.8148	0.7661	0.9626	0.9207	0.9084	0.6759	0.0979	0.4262	0.5994	0.7431	3.3011	7.2333	3.5701	0.9322	0.6899	18.2071	-51.2352	19.0721	-48.8447	18.3919	-50.2135
6	$E_{CORR(BLYP)} / E_{LUMO_{CORR(BLYP)}}$	0.8370	0.0731	0.8901	0.0200	0.9101	0.1148	0.5858	0.7887	0.7214	45.5462	5	0.8921	0.8920	0.8636	0.9529	0.9150	0.9447	0.7730	0.0656	0.4862	0.6588	0.5139	4.1174	7.4637	2.0817	0.9123	0.6062	18.8179	-49.4097	20.2050	-46.1911	17.5200	-49.9480
7	$E_{CORR(BLYP)} / \chi_{CORR(BLYP)}$	0.8342	0.0798	0.8949	0.0191	0.9140	0.2734	0.5729	0.7955	0.7951	47.8096	5	0.8689	0.8688	0.8343	0.9551	0.9137	0.9313	0.7239	0.0836	0.4589	0.6340	0.5142	3.9393	7.5933	2.5287	0.9191	0.6078	18.3289	-48.8464	19.2725	-45.6722	17.8342	-49.6133
8	$E_{CORR(BLYP)} / \eta_{CORR(BLYP)}$	0.8261	0.0734	0.8772	0.0223	0.8995	0.0519	0.6192	0.8146	0.7560	40.2908	5	0.8815	0.8814	0.8502	0.9471	0.9122	0.9403	0.7927	0.0692	0.5114	0.6884	0.5408	4.6005	7.9622	2.2862	0.9012	0.5336	18.3965	-54.0540	17.9292	-52.1666	19.6010	-49.5501
9	$E_{CORR(B3LYP)} / \chi_{CORR(B3LYP)}$	0.8183	0.0751	0.8697	0.0237	0.8934	0.3099	0.6377	0.8326	0.8291	37.7248	5	0.8574	0.8573	0.8199	0.9437	0.9059	0.9229	0.7135	0.0959	0.4918	0.6516	0.4884	4.8801	8.3194	2.7496	0.8958	0.5219	16.0694	-54.0524	18.2200	-49.5382	17.6263	-49.5754
10	$E_{CORR(B3LYP)} / E_{LUMO_{CORR(B3LYP)}}$	0.8060	0.0821	0.8632	0.0249	0.8880	0.0948	0.6536	0.8604	1.0093	35.6954	5	0.7888	0.7886	0.7331	0.9407	0.8972	0.8972	0.6949	0.1084	0.5492	0.7310	0.7251	5.1265	8.8845	4.0744	0.8888	0.4547	18.6311	-46.7816	17.5033	-50.1550	20.0132	-43.9804
11	$E_{CORR(B3LYP)} / E_{HOMO_{CORR(B3LYP)}}$	0.8058	0.1123	0.8999	0.0182	0.9181	0.1384	0.5590	0.8609	0.8133	50.4445	5	0.8628	0.8627	0.8267	0.9573	0.9045	0.9270	0.6969	0.0909	0.4169	0.6025	0.5318	3.7503	8.8943	2.6455	0.9259	0.5918	17.8543	-50.3190	18.5740	-49.2757	18.4400	-48.8594
12	$E_{CORR(BLYP)} / E_{HOMO_{CORR(BLYP)}}$	0.7845	0.1178	0.8805	0.0217	0.9022	0.4019	0.6108	0.9069	0.8656	41.5279	5	0.8446	0.8445	0.8036	0.9486	0.8917	0.9190	0.7048	0.0966	0.4817	0.6866	0.5659	4.4769	9.8695	2.9974	0.9094	0.5254	19.6301	-46.1700	18.0365	-49.6408	18.5387	-49.1631
13	$E_{CORR(HFX+LYP)} / E_{HOMO_{CORR(HFX+LYP)}}$	0.7022	0.1943	0.8735	0.0230	0.8965	0.1876	0.6285	1.0660	0.8352	38.9661	5	0.8553	0.8552	0.8172	0.9454	0.8620	0.9246	0.7274	0.0902	0.5080	0.7665	0.5366	4.7407	13.6364	2.7904	0.9062	0.4129	18.7263	-56.1730	18.7885	-48.3759	18.2294	-56.0729
14**	$E_{CORR(HFX+LYP)} / \omega_{CORR(HFX+LYP)}$	0.8190	0.0585	0.8469	0.0306	0.8775	0.1449	0.6932	0.8426	0.7238	28.6553	5	0.8669	0.8443	0.8665	0.9348	0.9038	0.9369	0.6714	0.0846	0.5140	0.6633	0.6290	5.2851	7.8104	2.0953	0.8720	0.1137	21.2369	-57.1164	18.6228	-59.5211	18.8275	-60.6302
15**	$E_{CORR(BLYP)} / \omega_{CORR(BLYP)}$	0.5313	0.3321	0.8087	0.0547	0.8634	0.1055	0.7011	1.2985	0.5013	15.7974	5	0.7976	0.7966	0.9301	0.9267	0.8196	0.8790	0.7862	0.0722	0.5608	1.0192	0.4867	3.9321	13.4888	1.0051	0.8741	-16.8911	29.9140	-146.7222	29.3147	-111.0799	28.2766	-156.3253

**Computed using data set represented with (d) and (e) respectively in Supporting Information Table S4.

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chicco, E. Papa, S. Kowarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotol. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_{ExtOK} Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S27. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N.																																
		Q^2_{LOO}	$R^2_{Q^2_{LOO}}$	R^2_{adj}	R^2_{adj}	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Xrand}	Q^2_{Xrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{CORR(M06-L)}$, $E_{CORR(M06-L)}^{HOMO}$	0.7274	0.0792	0.7882	0.0184	0.8066	0.3152	0.8304	0.9859	0.8642	43.7948	3	0.7161	0.7124	0.7906	0.893	0.8545	0.8111	0.5982	0.1386	0.6279	0.7393	0.7094	16.5513	23.3294	15.6824	0.8157	0.7222	8.5983	-20.3654	8.4089	-20.2828	8.8788	-20.0677
2	$E_{CORR(M06)}$, $E_{CORR(M06)}^{HOMO}$	0.723	0.0856	0.7904	0.0182	0.8087	0.3042	0.826	0.9938	0.8062	44.3782	3	0.7529	0.7497	0.8177	0.8942	0.8529	0.8397	0.6298	0.1259	0.6554	0.777	0.6464	16.3754	23.7044	13.6486	0.8191	0.7132	8.5041	-20.9155	8.4297	-19.9952	8.3211	-21.124
3	$E_{CORR(M06-2X)}$, $E_{CORR(M06-2X)}^{HOMO}$	0.6861	0.0852	0.7495	0.0218	0.7713	0.3115	0.9031	1.0581	0.7464	35.405	5	0.7882	0.7855	0.8438	0.8709	0.8287	0.8659	0.7351	0.0946	0.7496	0.8728	0.5921	19.5763	26.8679	11.6993	0.7784	0.6782	8.8287	-19.4964	8.6815	-19.6684	8.3277	-20.4056
4	$E_{CORR(M06-L)}$, $\chi_{CORR(M06-L)}$	0.6695	0.0904	0.7371	0.0229	0.7599	0.241	0.9253	1.0856	0.9136	33.2364	1	0.6826	0.6786	0.7659	0.8636	0.8192	0.8005	0.6188	0.102	0.7607	0.8855	0.6909	20.547	28.2853	17.5282	0.7694	0.6687	8.8857	-19.4851	8.7441	-19.5772	8.447	-20.6445
5	$E_{CORR(M06-L)}$, $\eta_{CORR(M06-L)}$	0.6694	0.0905	0.737	0.0229	0.7599	0.2345	0.9253	1.0857	0.9114	33.2309	1	0.6842	0.6801	0.7671	0.8636	0.8192	0.8027	0.6188	0.1047	0.756	0.8802	0.6891	20.5496	28.2921	17.4438	0.7684	0.6552	8.391	-20.3677	8.2215	-20.3838	8.5237	-19.8401
6	$E_{CORR(M06-L)}$, $\omega_{CORR(M06-L)}$	0.6606	0.0938	0.7311	0.0234	0.7544	0.2341	0.9358	1.1001	0.9231	32.2604	1	0.676	0.6719	0.7611	0.86	0.8141	0.7936	0.6117	0.1147	0.7582	0.8835	0.6849	21.016	29.0443	17.8928	0.7641	0.6600	8.9533	-19.5899	8.57	-20.1556	8.6003	-20.0929
7	$E_{CORR(M06)}$, $\chi_{CORR(M06)}$	0.6513	0.0953	0.7225	0.0241	0.7466	0.1828	0.9506	1.1152	0.9493	30.9351	1	0.6574	0.653	0.7473	0.8549	0.8076	0.7846	0.5909	0.0852	0.8021	0.9337	0.7317	21.6881	29.8475	18.9236	0.7581	0.6389	8.6126	-19.9997	8.7449	-19.6801	8.6409	-20.0249
8	$E_{CORR(M06)}$, $\eta_{CORR(M06)}$	0.6483	0.0968	0.7209	0.0243	0.7452	0.1819	0.9533	1.1199	0.9448	30.7042	1	0.6606	0.6563	0.7497	0.854	0.8059	0.7873	0.5942	0.0849	0.8106	0.9439	0.7261	21.8097	30.0983	18.7452	0.7573	0.6307	8.7074	-19.7769	8.6103	-19.9712	8.9118	-19.7475
9	$E_{CORR(M06)}$, $\omega_{CORR(M06)}$	0.623	0.1053	0.7024	0.0259	0.7283	0.1965	0.9843	1.1595	0.9104	28.1441	1	0.6849	0.6809	0.7676	0.8428	0.7911	0.7974	0.6325	0.086	0.8507	0.989	0.7023	23.2546	32.2671	17.4038	0.7384	0.6227	9.0155	-19.4652	8.7895	-19.6078	9.0004	-19.8773
10	$E_{CORR(M06-2X)}$, $\eta_{CORR(M06-2X)}$	0.6188	0.1118	0.7049	0.0257	0.7305	0.2631	0.9803	1.166	0.88	28.4672	4	0.7055	0.7018	0.7828	0.8443	0.79	0.8137	0.6618	0.0493	0.8277	0.9717	0.6909	23.0617	32.6282	16.2639	0.7446	0.6063	8.8405	-20.0973	8.5221	-20.224	8.3951	-20.8453
11	$E_{CORR(M06-2X)}$, $\chi_{CORR(M06-2X)}$	0.6152	0.1115	0.7007	0.026	0.7267	0.2568	0.9872	1.1714	0.8773	27.9249	4	0.7073	0.7036	0.7842	0.8417	0.7876	0.8137	0.6626	0.0659	0.8317	0.976	0.6874	23.3872	32.934	16.1639	0.7397	0.6057	8.8554	-20.2998	8.7956	-19.7671	8.7678	-20.1372
12	$E_{CORR(M06-2X)}$, $\omega_{CORR(M06-2X)}$	0.6081	0.113	0.6946	0.0266	0.7212	0.266	0.9972	1.1822	0.8982	27.1551	2	0.6932	0.6893	0.7738	0.838	0.7827	0.8002	0.6532	0.074	0.8291	0.972	0.6962	23.8653	33.5397	16.9426	0.7334	0.5916	8.6783	-19.9128	8.995	-19.4032	8.0754	-21.0687
13	$E_{CORR(M06-2X)}$, $E_{CORR(M06-2X)}^{LUMO}$	0.5315	0.1223	0.6208	0.033	0.6537	0.0027	1.1112	1.2926	1.1335	19.8246	0	0.5115	0.5053	0.6397	0.7906	0.7276	0.644	0.5047	0.0598	0.9233	1.0646	0.9344	29.6343	40.0998	26.979	0.6715	0.5171	8.8175	-20.0737	8.3812	-20.1623	8.52	-20.5053
14	$E_{CORR(M06-L)}$, $E_{CORR(M06-L)}^{LUMO}$	0.5227	0.134	0.624	0.0327	0.6567	-0.002	1.1064	1.3046	1.2451	20.0891	0	0.4106	0.4031	0.5653	0.7928	0.7266	0.5574	0.3797	0.0779	0.9312	1.0896	1.0321	29.3782	40.8468	32.5554	0.6744	0.5009	8.561	-20.8725	8.4949	-19.9891	8.2863	-22.2119
15	$E_{CORR(M06)}$, $E_{CORR(M06)}^{LUMO}$	0.494	0.1452	0.6048	0.0344	0.6392	-0.0325	1.1343	1.3434	1.2058	18.6019	0	0.4472	0.4401	0.5923	0.7799	0.7039	0.5859	0.4352	0.0744	0.9476	1.1042	0.999	30.8794	43.3104	30.5339	0.6589	0.4860	8.5851	-20.6376	8.9148	-19.2392	8.4583	-21.0518

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N_Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S28. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA100), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through 30% (random) splitting method.

Model	Variables	N																																
		Q^2_{LO}	$R^2-Q^2_{LO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ex.OK	Q^2_{T1}	Q^2_{T2}	Q^2_{T3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{F}^2	Δr^2	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Vint}	Q^2_{Vint}	R^2_{Vintd}	Q^2_{Vintd}	R^2_{Vintd}	Q^2_{Vintd}
1	$E_{CORR(M06)}^{HOMO}$	0.7232	0.0543	0.7617	0.0159	0.7775	0.2798	0.8119	0.9056	0.7592	48.9348	5	0.8235	0.8175	0.8055	0.8749	0.8464	0.8892	0.8156	0.0106	0.673	0.7509	0.585	20.4338	25.4217	7.4932	0.7812	0.6946	6.528	-15.1181	6.7988	-14.4774	6.9398	-14.6168
2	$E_{CORR(M06-L)}^{HOMO}$	0.7216	0.0523	0.7578	0.0161	0.774	0.2845	0.8184	0.9082	0.849	47.9387	5	0.7794	0.7718	0.7568	0.8726	0.8451	0.8573	0.7515	0.0787	0.6648	0.7402	0.6746	20.7624	25.5704	9.3694	0.7775	0.6937	6.7815	-14.755	6.1939	-15.284	6.9448	-14.5312
3	$E_{CORR(M06-2X)}^{HOMO}$	0.7019	0.0575	0.7421	0.0172	0.7593	0.2901	0.8445	0.9399	0.6396	44.1733	5	0.8748	0.8705	0.8619	0.8632	0.8326	0.9264	0.8493	0.0489	0.6967	0.7734	0.4619	22.1063	27.3842	5.3183	0.7622	0.6784	6.7294	-14.7926	6.7132	-14.5327	6.8356	-14.6129
4	$E_{CORR(M06-L)}^{HOMO} / \eta_{CORR(M06-L)}$	0.6754	0.0602	0.7168	0.0189	0.7357	0.3326	0.885	0.9807	0.7907	38.9602	5	0.8086	0.802	0.789	0.8477	0.8155	0.8829	0.799	0.0726	0.7154	0.7933	0.5677	24.2823	29.8127	8.1281	0.7395	0.6480	6.5491	-14.7557	6.4427	-14.9318	6.433	-15.1223
5	$E_{CORR(M06-L)}^{HOMO} / \chi_{CORR(M06-L)}$	0.6753	0.0601	0.7165	0.0189	0.7354	0.3367	0.8854	0.9809	0.7992	38.9183	5	0.8044	0.7978	0.7844	0.8476	0.8154	0.8803	0.7937	0.0739	0.7168	0.7948	0.5739	24.3015	29.825	8.3039	0.7387	0.6416	6.6108	-14.7732	6.7026	-14.7912	6.53	-14.8213
6	$E_{CORR(M06)}^{HOMO} / \eta_{CORR(M06)}$	0.663	0.0617	0.705	0.0197	0.7247	0.2685	0.9033	0.9994	0.7982	36.8452	5	0.805	0.7983	0.785	0.8403	0.8066	0.8815	0.7806	0.0678	0.74	0.8196	0.5709	25.2924	30.9598	8.2816	0.7282	0.6256	6.8607	-14.5786	6.7568	-14.529	6.5069	-15.0691
7	$E_{CORR(M06)}^{HOMO} / \chi_{CORR(M06)}$	0.662	0.0617	0.7039	0.0197	0.7237	0.2617	0.9049	1.0008	0.7942	36.6623	5	0.8069	0.8003	0.7871	0.8397	0.806	0.8826	0.7817	0.066	0.7418	0.8215	0.5583	25.3837	31.0503	8.2004	0.7282	0.6216	6.6641	-14.6638	6.4698	-15.0797	6.8296	-14.6233
8	$E_{CORR(M06)}^{HOMO} / \omega_{CORR(M06)}$	0.6613	0.0648	0.7066	0.0196	0.7261	0.2755	0.9008	1.0018	0.8018	37.1222	5	0.8032	0.7965	0.7831	0.8413	0.8065	0.8818	0.7646	0.0661	0.7559	0.8389	0.59	25.1553	31.1107	8.3569	0.7310	0.6292	6.8367	-14.5961	6.8001	-14.572	7.0385	-14.5611
9	$E_{CORR(M06-L)}^{HOMO} / \omega_{CORR(M06-L)}$	0.6609	0.0648	0.7061	0.0196	0.7257	0.3244	0.9015	1.0024	0.7898	37.0447	5	0.809	0.8025	0.7895	0.8411	0.8065	0.8819	0.8079	0.0706	0.7296	0.8103	0.5737	25.1935	31.1497	8.1098	0.7292	0.6409	6.4979	-15.0033	6.6576	-14.5171	6.7142	-14.9102
10	$E_{CORR(M06-2X)}^{HOMO} / \chi_{CORR(M06-2X)}$	0.6564	0.0667	0.7032	0.0198	0.723	0.3365	0.9059	1.0091	0.7164	36.5454	5	0.8429	0.8375	0.8268	0.8392	0.8036	0.9054	0.8173	0.0533	0.759	0.8454	0.5295	25.4424	31.5655	6.6715	0.7284	0.6221	6.4501	-15.47	6.4223	-15.1167	6.7822	-14.9699
11	$E_{CORR(M06-2X)}^{HOMO} / \eta_{CORR(M06-2X)}$	0.656	0.0668	0.7029	0.0198	0.7227	0.3487	0.9064	1.0097	0.7178	36.4911	5	0.8422	0.8369	0.8261	0.839	0.8034	0.9051	0.8154	0.0538	0.7609	0.8471	0.5288	25.4698	31.6022	6.6986	0.7253	0.6259	6.3675	-15.4972	6.5863	-15.0296	6.2227	-15.5397
12	$E_{CORR(M06-2X)}^{HOMO} / \omega_{CORR(M06-2X)}$	0.6396	0.0698	0.6886	0.0208	0.7093	0.3612	0.928	1.0335	0.7395	34.1662	5	0.8326	0.8268	0.8154	0.83	0.7924	0.896	0.832	0.0511	0.7744	0.8616	0.5575	26.6991	33.1097	7.1096	0.7130	0.6059	6.6898	-15.1298	6.5104	-14.7607	6.7995	-14.9874
13	$E_{CORR(M06-2X)}^{LUMO}$	0.5755	0.0832	0.6343	0.0244	0.6587	0.0378	1.0056	1.1215	0.9235	27.0222	4	0.7389	0.73	0.7122	0.7943	0.7465	0.8103	0.7924	0.0762	0.8302	0.9239	0.7376	31.3488	38.9937	11.0877	0.6667	0.5487	6.5367	-15.3422	6.402	-14.9494	6.632	-14.9778
14	$E_{CORR(M06-L)}^{LUMO} / \chi_{CORR(M06-L)}$	0.5624	0.0938	0.6316	0.0246	0.6561	0.0137	1.0094	1.1387	1.0665	26.7135	0	0.6518	0.6399	0.6161	0.7924	0.7416	0.7254	0.5871	0.1498	0.8563	0.9616	0.8713	31.5865	40.1981	14.7868	0.6651	0.5290	6.6064	-15.6886	6.7866	-14.4637	6.8157	-15.2508
15	$E_{CORR(M06)}^{LUMO} / \omega_{CORR(M06)}$	0.5461	0.0972	0.6179	0.0255	0.6434	-0.0012	1.028	1.1597	1.0389	25.2549	0	0.6696	0.6583	0.6358	0.783	0.7267	0.7437	0.6479	0.1244	0.864	0.9679	0.8556	32.7601	41.6903	14.0301	0.6506	0.5347	6.484	-15.4977	6.7901	-14.4595	6.6852	-15.0387

Models are arranged according to Q^2_{LO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecotox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N. Ex.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S29. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through activity sampling (ordered response) splitting method.

Model	Variables	N																																				
		Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Ystr}	Q^2_{Ystr}	R^2_{Yrand}	Q^2_{Yrand}	R^2_{Yrand}	Q^2_{Yrand}				
1	$E_{CORR(M06-2X)}^{HOMO}$	0.9611	0.0198	0.9733	0.0076	0.9809	0.2325	0.2937	0.4194	0.4926	128.4472	5	0.8998	0.8978	0.9463	0.9904	0.9806	0.9508	0.8575	0.0754	0.2205	0.3249	0.42	0.69	1.4072	1.2131	0.9819	0.6700	27.3059	-109.8418	28.4018	-108.0133	29.1242	-105.3201				
2	$E_{CORR(M06-L)}^{HOMO}$	0.9559	0.0225	0.9697	0.0086	0.9784	0.1867	0.3125	0.4463	0.4916	113.1597	5	0.9002	0.8982	0.9465	0.9891	0.978	0.9484	0.8666	0.0434	0.2566	0.3811	0.3949	0.7812	1.5932	1.2081	0.9792	0.6044	29.605	-106.4775	29.0671	-107.032	28.6567	-113				
3	$E_{CORR(M06-L)}^{HOMO}$	0.9539	0.0238	0.9688	0.0089	0.9777	0.1881	0.3175	0.4563	0.4764	109.5577	5	0.9063	0.9044	0.9498	0.9887	0.977	0.9523	0.8764	0.0705	0.2713	0.4095	0.3817	0.8063	1.6657	1.1349	0.9789	0.5603	28.4443	-126.6977	29.2881	-106.2399	28.3569	-119.7613				
4	$E_{CORR(M06-L)}^{HOMO}$	0.9438	0.032	0.9661	0.0097	0.9758	0.4404	0.331	0.5039	0.6715	100.6098	5	0.8138	0.8101	0.9002	0.9877	0.9719	0.9094	0.7557	0.123	0.2497	0.4003	0.5515	0.8763	2.0312	2.2544	0.9779	0.4794	29.0988	-94.1846	27.7599	-109.4249	27.7401	-100.3493				
5	$E_{CORR(M06-L)}^{HOMO}$	0.9359	0.036	0.9606	0.0113	0.9718	0.3715	0.3567	0.5382	0.7111	86.2733	5	0.7912	0.787	0.8881	0.9857	0.9676	0.8958	0.7419	0.1119	0.2466	0.3837	0.5937	1.0178	2.3175	2.5286	0.9749	0.3744	28.4071	-98.586	29.9315	-100.8189	27.3732	-101.3424				
6	$E_{CORR(M06-L)}^{HOMO}$	0.9338	0.0372	0.9594	0.0116	0.971	0.3918	0.3621	0.547	0.7249	83.6362	5	0.7831	0.7787	0.8837	0.9853	0.9665	0.8911	0.7333	0.1109	0.2523	0.3956	0.598	1.049	2.3934	2.6271	0.9739	0.3444	29.6038	-95.005	28.0895	-109.1424	29.8759	-92.1366				
7	$E_{CORR(M06-2X)}^{HOMO}$	0.932	0.0376	0.9574	0.0122	0.9696	0.4162	0.3707	0.5545	0.7322	79.6938	5	0.7787	0.7742	0.8813	0.9846	0.9656	0.8864	0.7211	0.0729	0.3056	0.478	0.541	1.0993	2.4593	2.6804	0.9720	0.3794	28.7329	-100.8902	28.9748	-101.9151	28.0158	-102.8402				
8	$E_{CORR(M06-2X)}^{HOMO}$	0.9308	0.038	0.9563	0.0125	0.9688	0.4878	0.3756	0.5593	0.7085	77.536	5	0.7928	0.7886	0.8889	0.9841	0.9651	0.8965	0.7389	0.0904	0.2871	0.4424	0.5737	1.1289	2.5021	2.5095	0.9716	0.4423	28.7792	-97.627	29.3015	-107.527	29.0874	-94.9439				
9	$E_{CORR(M06-2X)}^{HOMO}$	0.9287	0.0398	0.9559	0.0126	0.9685	0.4364	0.3774	0.5675	0.7146	76.7836	5	0.7891	0.7849	0.887	0.984	0.964	0.8938	0.7376	0.1002	0.2799	0.4302	0.5661	1.1396	2.5767	2.5535	0.9710	0.3673	29.2764	-94.4813	27.779	-111.1206	29.375	-94.7392				
10	$E_{CORR(M06)}^{HOMO}$	0.9256	0.0426	0.9556	0.0127	0.9683	0.4757	0.3787	0.5796	0.7536	76.246	5	0.7655	0.7608	0.8743	0.9839	0.9625	0.8802	0.7115	0.0929	0.2976	0.478	0.6161	1.1474	2.6879	2.8397	0.9714	0.4077	28.3468	-95.6521	28.0674	-110.4506	29.5381	-93.032				
11	$E_{CORR(M06)}^{HOMO}$	0.9226	0.0447	0.9543	0.0131	0.9674	0.4747	0.3839	0.5912	0.7411	74.1171	5	0.7732	0.7686	0.8784	0.9834	0.9609	0.8833	0.7185	0.0825	0.2931	0.472	0.5956	1.1793	2.7957	2.7464	0.9710	0.2739	29.7615	-90.2826	28.193	-108.2182	29.3719	-92.5418				
12	$E_{CORR(M06)}^{HOMO}$	0.9032	0.0612	0.9501	0.0142	0.9644	0.4063	0.4011	0.6614	0.6777	67.6918	5	0.8104	0.8066	0.8983	0.9819	0.9514	0.9016	0.7843	0.1195	0.2948	0.4877	0.5544	1.2872	3.4999	2.2963	0.9696	0.3628	28.5542	-91.4535	28.1268	-108.4975	27.707	-92.6319				
13	$E_{CORR(M06-2X)}^{LUMO}$	0.8823	0.0682	0.9307	0.0198	0.9505	0.1528	0.473	0.7293	0.8589	47.9911	1	0.6955	0.6893	0.8367	0.9746	0.9408	0.8449	0.623	0.0467	0.3896	0.6139	0.6646	1.7895	4.2555	3.6882	0.9538	-0.0319	28.4266	-95.5974	28.2882	-104.9962	28.451	-95.8422				
14	$E_{CORR(M06)}^{LUMO}$	0.8808	0.0646	0.9235	0.0219	0.9453	0.1002	0.4969	0.734	0.9187	43.2416	1	0.6515	0.6445	0.8132	0.9719	0.9403	0.8214	0.5774	0.0483	0.4186	0.6304	0.7235	1.9753	4.3096	4.2201	0.9486	-0.1326	28.5311	-100.8503	28.382	-109.0724	28.4172	-100.3383				
15	$E_{CORR(M06-L)}^{LUMO}$	0.8734	0.0744	0.927	0.0209	0.9479	0.0484	0.4854	0.7561	0.8966	45.4447	1	0.6681	0.6614	0.822	0.9732	0.9366	0.8308	0.5941	0.0445	0.4056	0.6353	0.7042	1.8845	4.5738	4.0196	0.9520	0.1360	28.0617	-91.414	29.1216	-104.2578	28.2849	-91.9444				

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecolox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N.Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value

Table S30. Value of internal and external validation parameters of two-descriptor QSAR models, for mutagenicity (LogTA98), developed using electron-correlation (CORR(M06), CORR(M06-L), CORR(M06-2X)) based methods through activity sampling 30% (random) splitting method.

Model	Variables	N																																
		Q^2_{LOO}	$R^2-Q^2_{LOO}$	R^2_{adj}	$R^2-R^2_{adj}$	R^2	ΔK	RMSE _{TR}	RMSE _{CV}	RMSE _{EXT}	F	Ext.OK	Q^2_{F1}	Q^2_{F2}	Q^2_{F3}	CCC _{TR}	CCC _{CV}	CCC _{EXT}	\bar{r}^2_{int}	Δr^2_{int}	MAE _{TR}	MAE _{CV}	MAE _{EXT}	RSS _{TR}	PRESS _{CV}	PRESS _{EXT}	R^2_{LMO}	Q^2_{LMO}	R^2_{Year}	Q^2_{Year}	R^2_{Nrand}	Q^2_{Nrand}	R^2_{Yrand}	Q^2_{Yrand}
1	$E_{CORR(M06-2X)}$, $E_{CORR(M06-2X)}^{HOMO}$	0.8891	0.0546	0.9276	0.0161	0.9437	0.2141	0.4696	0.659	0.3061	58.6196	5	0.9745	0.9741	0.9761	0.971	0.9434	0.9874	0.899	0.0141	0.3975	0.562	0.2843	2.2053	4.3422	0.3748	0.9462	0.7427	22.6424	-59.2023	22.1038	-67.8088	22.7615	-59.8664
2	$E_{CORR(M06-L)}$, $E_{CORR(M06-L)}^{HOMO}$	0.8799	0.0588	0.9211	0.0175	0.9386	0.2153	0.4901	0.6858	0.3138	53.5449	5	0.9732	0.9728	0.9748	0.9684	0.9388	0.9877	0.8535	0.018	0.3977	0.5631	0.2669	2.4015	4.7028	0.3938	0.9405	0.7126	22.6056	-60.9789	23.2717	-64.3708	21.9391	-62.4158
3	$E_{CORR(M06)}$, $E_{CORR(M06)}^{HOMO}$	0.8798	0.0594	0.9219	0.0174	0.9392	0.2166	0.4876	0.6858	0.3258	54.1107	5	0.9711	0.9706	0.9729	0.9687	0.9388	0.9866	0.8489	0.0182	0.3965	0.5594	0.2738	2.3779	4.7032	0.4246	0.9433	0.6687	22.1341	-61.5401	21.7539	-69.0825	23.2127	-59.496
4	$E_{CORR(M06)}$, $\omega_{CORR(M06)}$	0.8361	0.0768	0.888	0.0249	0.9129	0.3879	0.5839	0.801	0.5497	36.6812	5	0.9177	0.9164	0.9228	0.9545	0.9134	0.965	0.7415	0.0495	0.4704	0.6593	0.4495	3.4094	6.4154	1.2087	0.9177	0.5801	22.2309	-61.4706	21.8487	-68.0915	22.9083	-60.7558
5	$E_{CORR(M06-L)}$, $\omega_{CORR(M06-L)}$	0.8297	0.0644	0.8638	0.0303	0.894	0.3397	0.644	0.8165	0.5152	29.5338	5	0.9277	0.9266	0.9322	0.9441	0.9114	0.9661	0.7907	0.0326	0.4723	0.6456	0.504	4.1471	6.666	1.0616	0.8928	0.5006	21.938	-65.0466	22.0054	-68.7132	22.4464	-65.3091
6	$E_{CORR(M06-2X)}$, $\chi_{CORR(M06-2X)}$	0.819	0.0852	0.8768	0.0274	0.9042	0.2938	0.6125	0.8417	0.6359	33.0221	5	0.8898	0.8881	0.8967	0.9497	0.907	0.9463	0.7828	0.0366	0.486	0.6889	0.6234	3.751	7.0844	1.6177	0.9071	0.5675	21.9262	-67.103	21.5217	-69.9175	22.9683	-64.6405
7	$E_{CORR(M06-2X)}$, $\eta_{CORR(M06-2X)}$	0.8165	0.0786	0.865	0.03	0.895	0.2906	0.641	0.8476	0.5799	29.843	5	0.9084	0.907	0.9141	0.9446	0.9043	0.9573	0.7685	0.0406	0.4835	0.6721	0.5623	4.1086	7.1835	1.3453	0.8943	0.4570	22.4365	-64.9872	22.298	-67.0859	21.8294	-67.7045
8	$E_{CORR(M06-L)}$, $\eta_{CORR(M06-L)}$	0.8099	0.0844	0.864	0.0302	0.8942	0.3713	0.6434	0.8626	0.5284	29.5934	5	0.9239	0.9228	0.9287	0.9442	0.9009	0.9646	0.783	0.0346	0.4931	0.6914	0.5189	4.1396	7.4413	1.1167	0.8978	0.5739	22.0247	-65.2178	22.6672	-66.1658	21.9127	-66.4633
9	$E_{CORR(M06-L)}$, $\chi_{CORR(M06-L)}$	0.8027	0.0823	0.8522	0.0328	0.885	0.3618	0.6708	0.8788	0.5297	26.9468	5	0.9235	0.9224	0.9283	0.939	0.8963	0.9646	0.7792	0.0358	0.5096	0.7018	0.517	4.4995	7.7224	1.1222	0.8827	0.5222	22.4299	-63.988	22.4202	-67.0032	21.816	-66.269
10	$E_{CORR(M06-2X)}$, $\omega_{CORR(M06-2X)}$	0.7985	0.0968	0.8654	0.0299	0.8953	0.2566	0.6401	0.8881	0.5891	29.9385	5	0.9054	0.904	0.9113	0.9448	0.8976	0.9527	0.8168	0.0318	0.4882	0.7107	0.5444	4.0969	7.8867	1.388	0.8965	0.5239	21.9545	-72.5352	21.6271	-69.1194	22.7447	-69.239
11	$E_{CORR(M06)}$, $\eta_{CORR(M06)}$	0.7958	0.0878	0.8503	0.0333	0.8836	0.3053	0.675	0.8941	0.5827	26.5669	5	0.9075	0.9061	0.9132	0.9382	0.8923	0.9548	0.7994	0.0298	0.5312	0.7339	0.5816	4.5563	7.9939	1.3584	0.8842	0.4748	22.5092	-63.633	22.8636	-67.0277	22.0439	-65.375
12	$E_{CORR(M06)}$, $\chi_{CORR(M06)}$	0.7936	0.0859	0.8451	0.0344	0.8795	0.3428	0.6866	0.8987	0.5744	25.5558	5	0.9101	0.9087	0.9157	0.9359	0.8912	0.9578	0.7739	0.0376	0.5247	0.7198	0.5686	4.7149	8.0772	1.3199	0.8804	0.5043	21.2928	-66.6456	22.8814	-65.4006	22.3948	-63.6
13	$E_{CORR(M06-2X)}$, $E_{CORR(M06-2X)}^{LUMO}$	0.6705	0.1492	0.7682	0.0515	0.8197	0.2581	0.8401	1.1357	0.6439	15.9117	5	0.887	0.8853	0.8941	0.9009	0.8217	0.956	0.7036	0.0647	0.664	0.9172	0.5402	7.0573	12.8982	1.6582	0.8271	0.0144	22.8463	-66.7218	22.6558	-67.4599	22.231	-68.1258
14	$E_{CORR(M06-L)}$, $E_{CORR(M06-L)}^{LUMO}$	0.665	0.1531	0.7661	0.052	0.8181	0.09	0.8438	1.1451	0.6628	15.7427	5	0.8803	0.8785	0.8878	0.9	0.8214	0.9548	0.7152	0.0623	0.6627	0.8979	0.5718	7.1193	13.1118	1.7571	0.8230	0.1211	22.0395	-66.0791	21.3707	-69.9536	22.0005	-66.2792
15	$E_{CORR(M06)}$, $E_{CORR(M06)}^{LUMO}$	0.6506	0.1511	0.745	0.0567	0.8016	0.1777	0.8811	1.1695	0.663	14.1452	5	0.8802	0.8784	0.8877	0.8899	0.8099	0.9543	0.7029	0.0659	0.7078	0.9499	0.5692	7.7638	13.6769	1.7585	0.8063	-0.0129	22.0868	-70.1	22.5876	-66.138	22.0817	-69.3175

Models are arranged according to Q^2_{LOO} values.

Parameters are obtained through QSARINS software. Ref. [54] N. Chirico, E. Papa, S. Kovarich, S. Cassani, P. Gramatica, QSAR Res. Unit in Environ. Chem. and Ecolox. University of Insubria, Varese, Italy, 2012. (<http://www.qsar.it>).

R^2_{adj} adjusted R^2

N.Ext.OK Number of external validation parameters meeting threshold values

F "F-test" statistic value