properties

Name	Reference	Computable description	
atom-bond connectivity index	<u>Mordred</u>	[Descriptors] index 0	
Graovac-Ghorbani atom-bond connectivity index	<u>Mordred</u>	[Descriptors] index 1	
[acidic/basic] group count	Mordred	[Descriptors] index [2 3]	
Sp[Abs/max/Diam/AD/MAD] of adjacency matrix	Mordred	[Descriptors] index [4- 8]	
LogEE of adjacency matrix	Mordred	[Descriptors] index 9	
V[E/R][1-3] of adjacency matrix	Mordred	[Descriptors] index	
Aromatic [atoms/bonds]	Mordred	[Descriptors] index	
Number of [all/heavy/spiro] atoms	Mordred	[Descriptors] index	
Number of [\bridgehead/hetero] atoms	<u>Mordred</u>	[Descriptors] index	
Number of [\H/B/C/N/O/S/P/F/Cl/Br/I] atoms	<u>Mordred</u>	[Descriptors] index [23-33]	
Number of halogen atoms	<u>Mordred</u>	[Descriptors] index 34	
moreau-broto autocorrelation of lag [0-8] weighted by	<u>Mordred</u>	[Descriptors] index	
valence electrons	<u>Mordred</u>	[Descriptors] index	
sigma electrons	<u>Mordred</u>	[Descriptors] index [44-52]	
intrinsic state	<u>Mordred</u>	[Descriptors] index [53-61]	
atomic number	Mordred	[Descriptors] index [62-70]	

Name	Reference	Computable	description
mass	<u>Mordred</u>	[Descriptors] index [71-79]	
vdw volume	<u>Mordred</u>	[Descriptors] index [80-88]	
sanderson EN	<u>Mordred</u>	[Descriptors] index [89-97]	
pauling EN	<u>Mordred</u>	[Descriptors] index [98-106]	
allred-rocow EN	<u>Mordred</u>	[Descriptors] index [107-115]	
polarizability	<u>Mordred</u>	[Descriptors] index [116-124]	
ionization potential	Mordred	[Descriptors] index [125-133]	
averaged moreau-broto autocorrelation of lag [0-8] weighted by	Mordred	[Descriptors] index [134-232]	
valence electrons	<u>Mordred</u>	[Descriptors] index [134-142]	
sigma electrons	<u>Mordred</u>	[Descriptors] index [143-151]	
intrinsic state	<u>Mordred</u>	[Descriptors] index [152-160]	
atomic number	Mordred	[Descriptors] index [161-169]	
mass	<u>Mordred</u>	[Descriptors] index [170-178]	
vdw volume	<u>Mordred</u>	[Descriptors] index [179-187]	
sanderson EN	Mordred	[Descriptors] index [188-196]	•••

		h the see	
Name	Reference	Computable	description
pauling EN	Mordred	[Descriptors] index [197-205]	
allred-rocow EN	Mordred	[Descriptors] index [206-214]	
polarizability	Mordred	[Descriptors] index [215-223]	
ionization potential	Mordred	[Descriptors] index [224-232]	
centered moreau-broto autocorrelation of lag [0-8] weighted by	Mordred	[Descriptors] index [233-340]	•••
gasteiger charge	Mordred	[Descriptors] index [233-340]	•••
valence electrons	Mordred	[Descriptors] index [242-250]	•••
sigma electrons	Mordred	[Descriptors] index [251-259]	
intrinsic state	Mordred	[Descriptors] index [260-268]	
atomic number	Mordred	[Descriptors] index [269-277]	
mass	Mordred	[Descriptors] index [278-286]	
vdw volume	Mordred	[Descriptors] index [287-295]	
sanderson EN	Mordred	[Descriptors] index [296-304]	
pauling EN	Mordred	[Descriptors] index [305-313]	
allred-rocow EN	Mordred	[Descriptors] index [314-322]	

Name	Reference	Computable	description
polarizability	Mordred	[Descriptors] index [323-331]	
ionization potential	Mordred	[Descriptors] index [332-340]	
averaged and centered moreau-broto autocorrelation of lag [0-8] weighted by	Mordred	[Descriptors] index [341-331]	
gasteiger charge	Mordred	[Descriptors] index [341-349]	
valence electrons	Mordred	[Descriptors] index [350-358]	
sigma electrons	Mordred	[Descriptors] index [359-367]	
intrinsic state	Mordred	[Descriptors] index [368-376]	
atomic number	Mordred	[Descriptors] index [377-385]	
mass	Mordred	[Descriptors] index [386-394]	
vdw volume	<u>Mordred</u>	[Descriptors] index [395-403]	
sanderson EN	Mordred	[Descriptors] index [404-412]	
pauling EN	Mordred	[Descriptors] index [413-421]	
allred-rocow EN	Mordred	[Descriptors] index [422-430]	
polarizability	Mordred	[Descriptors] index [431-439]	
ionization potential	Mordred	[Descriptors] index [440-448]	

Name	Reference	Computable	description
moran coefficient of lag [1-8] weighted by	Mordred	[Descriptors] index [449-544]	
gasteiger charge	Mordred	[Descriptors] index [449-456]	
valence electrons	Mordred	[Descriptors] index [457-464]	
sigma electrons	Mordred	[Descriptors] index [465-472]	
intrinsic state	Mordred	[Descriptors] index [473-480]	
atomic number	Mordred	[Descriptors] index [481-488]	
mass	Mordred	[Descriptors] index [489-496]	
vdw volume	Mordred	[Descriptors] index [497-504]	
sanderson EN	Mordred	[Descriptors] index [505-512]	
pauling EN	Mordred	[Descriptors] index [513-520]	
allred-rocow EN	Mordred	[Descriptors] index [521-528]	
polarizability	Mordred	[Descriptors] index [529-536]	
ionization potential	Mordred	[Descriptors] index [537-544]	
geary coefficient of lag [1-8] weighted by	Mordred	[Descriptors] index [545-640]	
gasteiger charge	Mordred	[Descriptors] index [545-562]	

Name	Reference	Computable	description
valence electrons	<u>Mordred</u>	[Descriptors] index [553-560]	
sigma electrons	<u>Mordred</u>	[Descriptors] index [561-568]	
intrinsic state	Mordred	[Descriptors] index [569-576]	
atomic number	<u>Mordred</u>	[Descriptors] index [577-584]	
mass	<u>Mordred</u>	[Descriptors] index [585-592]	
vdw volume	<u>Mordred</u>	[Descriptors] index [593-600]	
sanderson EN	<u>Mordred</u>	[Descriptors] index [601-608]	
pauling EN	<u>Mordred</u>	[Descriptors] index [609-616]	
allred-rocow EN	<u>Mordred</u>	[Descriptors] index [617-624]	
polarizability	<u>Mordred</u>	[Descriptors] index [625-632]	
ionization potential	<u>Mordred</u>	[Descriptors] index [633-640]	
first [heighest/lowest] eigenvalue of Burden matrix weighted by	<u>Mordred</u>	[Descriptors] index [545-640]	
gasteiger charge	<u>Mordred</u>	[Descriptors] index [641-642]	
valence electrons	<u>Mordred</u>	[Descriptors] index [643-644]	
sigma electrons	Mordred	[Descriptors] index [645-646]	

Name	Reference	Computable	description
intrinsic state	<u>Mordred</u>	[Descriptors] index [647-648]	
atomic number	Mordred	[Descriptors] index [649-650]	
mass	Mordred	[Descriptors] index [651-652]	
vdw volume	Mordred	[Descriptors] index [653-654]	•••
sanderson EN	Mordred	[Descriptors] index [655-656]	•••
pauling EN	Mordred	[Descriptors] index [657-658]	•••
allred-rocow EN	Mordred	[Descriptors] index [659-660]	•••
polarizability	Mordred	[Descriptors] index [661-662]	•••
ionization potential	Mordred	[Descriptors] index [663-664]	•••
Balaban's J index	<u>Mordred</u>	[Descriptors] index x	•••
0: graph energy	<u>Mordred</u>	[Descriptors] index x	•••
1: leading eigenvalue	<u>Mordred</u>	[Descriptors] index x	
2: spectral diamiter	<u>Mordred</u>	[Descriptors] index x	
3: spectral absolute diviation	<u>Mordred</u>	[Descriptors] index x	
4: spectral mean absolute diviation	<u>Mordred</u>	[Descriptors] index x	
5: Estrada-like index	<u>Mordred</u>	[Descriptors] index x	
6: spectral moment	Mordred	[Descriptors] index x	
7: coefficient sum of the last eigenvector	Mordred	[Descriptors] index x	
8: average coefficient of the last eigenvector	Mordred	[Descriptors] index x	•••
9: logarithmic coefficient sum of the last eigenvector	Mordred	[Descriptors] index x	•••
10: Randic-like eigenvector-based index	Mordred	[Descriptors] index x	

Name	Reference	Computable	description
11: normalized Randic-like eigenvector-based index	Mordred	[Descriptors] index x	
12: logarithmic Randic-like eigenvector-based index	Mordred	[Descriptors] index x	
from Barysz matrix weighted by atomic number	Mordred	[Descriptors] index [666-678]	
from Barysz matrix weighted by mass	Mordred	[Descriptors] index [679-691]	
from Barysz matrix weighted by vdw volume	Mordred	[Descriptors] index [692-704]	
from Barysz matrix weighted by sanderson EN	Mordred	[Descriptors] index [705-717]	
from Barysz matrix weighted by pauling EN	Mordred	[Descriptors] index [718-730]	
from Barysz matrix weighted by allred-rocow EN	Mordred	[Descriptors] index [731-743]	
from Barysz matrix weighted by polarizability	Mordred	[Descriptors] index [744-756]	
from Barysz matrix weighted by ionization potential	<u>Mordred</u>	[Descriptors] index [757-769]	
Bertz CT	<u>Mordred</u>	[Descriptors] index 770	
number of all bonds in non-kekulized structure	<u>Mordred</u>	[Descriptors] index 771	
number of bonds connecting to heavy atom in non-kekulized structure	Mordred	[Descriptors] index 772	
number of [single/double/triple] bonds in non-kekulized structure	Mordred	[Descriptors] index [773-775]	
number of [Aromatic/multiple] bonds in non-kekulized structure	Mordred	[Descriptors] index [776-777]	
number of [single/double] bonds in kekulized structure	Mordred	[Descriptors] index [778-779]	

Name	Reference	Computable	description
partial negative surface area (version [1-5])	<u>Mordred</u>	[Descriptors] index [780-784]	
partial positive surface area (version [1-5])	<u>Mordred</u>	[Descriptors] index [785-789]	
difference in charged partial surface area (version [1-5])	<u>Mordred</u>	[Descriptors] index [790-794]	
fractional charged partial negative surface area (version [1-5])	Mordred	[Descriptors] index [795-799]	
fractional charged partial positive surface area (version [1-5])	Mordred	[Descriptors] index [800-804]	
surface weighted charged partial negative surface area (version [1-5])	<u>Mordred</u>	[Descriptors] index [805-809]	
surface weighted charged partial positive surface area (version [1-5])	<u>Mordred</u>	[Descriptors] index [810-814]	
relative [negative/positive] charge	<u>Mordred</u>	[Descriptors] index [815-816]	
relative [negative/positive] charge surface area	<u>Mordred</u>	[Descriptors] index [816-817]	
total [hydrophobic/polar] surface area	<u>Mordred</u>	[Descriptors] index [818-819]	
relative [hydrophobic/polar] surface area	<u>Mordred</u>	[Descriptors] index [820-821]	
SP carbon bound to [1-2] other carbon	<u>Mordred</u>	[Descriptors] index [822-823]	
SP2 carbon bound to [1-3] other carbon	<u>Mordred</u>	[Descriptors] index [824-825]	
SP3 carbon bound to [1-4] other carbon	<u>Mordred</u>	[Descriptors] index [826-827]	
hybridization ratio	<u>Mordred</u>	[Descriptors] index 828	

Name	Reference	Computable	description
the fraction of C atoms that are SP3 hybridized	<u>Mordred</u>	[Descriptors] index 829	
[3-7]-ordered Chi chain weighted by sigma electrons	<u>Mordred</u>	[Descriptors] index [834-838]	
[3-7]-ordered Chi chain weighted by valence electrons	<u>Mordred</u>	[Descriptors] index [839-843]	
[3-6]-ordered Chi cluster weighted by sigma electrons	<u>Mordred</u>	[Descriptors] index [844-847]	
[3-6]-ordered Chi cluster weighted by valence electrons	<u>Mordred</u>	[Descriptors] index [848-851]	
[4-6]-ordered Chi path-cluster weighted by sigma electrons	<u>Mordred</u>	[Descriptors] index [852-854]	
[4-6]-ordered Chi path-cluster weighted by valence electrons	<u>Mordred</u>	[Descriptors] index [855-857]	
[0-7]-ordered Chi path weighted by sigma electrons	<u>Mordred</u>	[Descriptors] index [858-865]	
[0-7]-ordered averaged Chi path weighted by sigma electrons	<u>Mordred</u>	[Descriptors] index [866-873]	
[0-7]-ordered Chi path weighted by sigma electrons	<u>Mordred</u>	[Descriptors] index [874-881]	
[0-7]-ordered averaged Chi path weighted by valence electrons	<u>Mordred</u>	[Descriptors] index [882-889]	
sum of constitutional weighted by	<u>Mordred</u>	[Descriptors] index [890-897]	
atomic number	<u>Mordred</u>	[Descriptors] index 890	
mass	<u>Mordred</u>	[Descriptors] index 891	
vdw volume	<u>Mordred</u>	[Descriptors] index 892	

Deference	Computable	description
Mordred	[Descriptors] index	
<u>Mordred</u>	[Descriptors] index 894	
Mordred	[Descriptors] index 895	
<u>Mordred</u>	[Descriptors] index 896	
<u>Mordred</u>	[Descriptors] index 897	
<u>Mordred</u>	[Descriptors] index [898-905]	
<u>Mordred</u>	[Descriptors] index 898	•••
<u>Mordred</u>	[Descriptors] index 899	•••
Mordred	[Descriptors] index 900	•••
<u>Mordred</u>	[Descriptors] index 893	•••
<u>Mordred</u>	[Descriptors] index 894	•••
Mordred	[Descriptors] index 895	•••
Mordred	[Descriptors] index 896	•••
<u>Mordred</u>	[Descriptors] index 897	•••
<u>Mordred</u>	[Descriptors] index 897	
	Mordred Mordred	Mordred Mordre

Name	Reference	Computable	description
graph energy	<u>Mordred</u>	[Descriptors] index 906	•••
leading eigenvalue	<u>Mordred</u>	[Descriptors] index 907	•••
spectral diamiter	<u>Mordred</u>	[Descriptors] index 908	•••
spectral absolute diviation	<u>Mordred</u>	[Descriptors] index 909	•••
spectral mean absolute diviation	<u>Mordred</u>	[Descriptors] index 910	•••
Estrada-like index	<u>Mordred</u>	[Descriptors] index 911	
spectral moment	Mordred	[Descriptors] index 912	
coefficient sum of the last eigenvector	Mordred	[Descriptors] index 913	•••
average coefficient of the last eigenvector	<u>Mordred</u>	[Descriptors] index 914	
logarithmic coefficient sum of the last eigenvector	<u>Mordred</u>	[Descriptors] index 915	
Randic-like eigenvector-based index	Mordred	[Descriptors] index 916	
normalized Randic-like eigenvector-based index	<u>Mordred</u>	[Descriptors] index 917	•••
logarithmic Randic-like eigenvector-based index	<u>Mordred</u>	[Descriptors] index 918	•••
from detourn matrix	<u>Mordred</u>	[Descriptors] index [906-918]	•••
detour index	<u>Mordred</u>	[Descriptors] index 919	

Name	Reference	Computable	description
graph energy	<u>Mordred</u>	[Descriptors] index 920	
leading eigenvalue	<u>Mordred</u>	[Descriptors] index 921	
spectral diamiter	<u>Mordred</u>	[Descriptors] index 922	
spectral absolute diviation	<u>Mordred</u>	[Descriptors] index 923	
spectral mean absolute diviation	<u>Mordred</u>	[Descriptors] index 924	
Estrada-like index	<u>Mordred</u>	[Descriptors] index 925	
coefficient sum of the last eigenvector	<u>Mordred</u>	[Descriptors] index 926	
average coefficient of the last eigenvector	<u>Mordred</u>	[Descriptors] index 927	
logarithmic coefficient sum of the last eigenvector	<u>Mordred</u>	[Descriptors] index 928	
Randic-like eigenvector-based index	<u>Mordred</u>	[Descriptors] index 929	
normalized Randic-like eigenvector-based index	<u>Mordred</u>	[Descriptors] index 930	
logarithmic Randic-like eigenvector-based index	<u>Mordred</u>	[Descriptors] index 931	
from detourn matrix	<u>Mordred</u>	[Descriptors] index [920-931]	
number of [sLi/ssBe/ssssBe/ssBH/sssB/ssssB/sCH3/dCH2/ssCH2/tch]	<u>Mordred</u>	[Descriptors] index [932-941]	
number of [dsCH/aaCH/sssCH/ddC/tsC/dssC/aasC/aaaC/ssssC/sNH3/sNH2]	<u>Mordred</u>	[Descriptors] index [942-952]	

Name	Reference	Computable	description
number of [ssNH2/dNH/ssNH/aaNH/tN/sssNH/dsN/aaN/sssN/ddsN/aasN]	Mordred	[Descriptors] index [953-963]	
number of [ssssN/sOH/dO/ssO/aaO/sF/sSiH3/ssSiH2/sssSiH/ssssSi]	Mordred	[Descriptors] index [964-973]	
number of [sPH2/ssPH/sssP/dsssP/sssssP/sSH/dS/ssS/aaS/dssS/ddssS]	Mordred	[Descriptors] index [974-984]	
number of [sCl/sGeH3/ssGeH2/sssGeH/ssssGe/sAsH2//ssAsH/sssAs]	Mordred	[Descriptors] index [985-992]	
number of [sssdAs/sssssAs/sSeH/dSe/ssSe/aaSe/dssSe/ddssSe/sBr]	Mordred	[Descriptors] index [993-1001]	
number of [sSnH3/ssSnH2/sssSnH/ssssSn/sI/sPbH3/ssPbH2/sssPbH/ssssPb]	Mordred	[Descriptors] index [1002-1010]	
sum of [sLi/ssBe/ssssBe/ssBH/sssB/ssssB/sCH3/dCH2/ssCH2/tch]	Mordred	[Descriptors] index [1011-1020]	
sum of [dsCH/aaCH/sssCH/ddC/tsC/dssC/aasC/aaaC/ssssC/sNH3/sNH2]	Mordred	[Descriptors] index [1021-1031]	
sum of [ssNH2/dNH/ssNH/aaNH/tN/sssNH/dsN/aaN/sssN/ddsN/aasN]	Mordred	[Descriptors] index [1032-1042]	
sum of [ssssN/sOH/dO/ssO/aaO/sF/sSiH3/ssSiH2/sssSiH/ssssSi]	Mordred	[Descriptors] index [1043-1052]	
sum of [sPH2/ssPH/sssP/dsssP/sssssP/sSH/dS/ssS/aaS/dssS/ddssS]	Mordred	[Descriptors] index [1053-1065]	
sum of [sCl/sGeH3/ssGeH2/sssGeH/ssssGe/sAsH2//ssAsH/sssAs]	Mordred	[Descriptors] index [1066-1071]	
sum of [sssdAs/sssssAs/sSeH/dSe/ssSe/aaSe/dssSe/ddssSe/sBr]	Mordred	[Descriptors] index [1072-1082]	
sum of [sSnH3/ssSnH2/sssSnH/ssssSn/sI/sPbH3/ssPbH2/sssPbH/ssssPb]	Mordred	[Descriptors] index [1083-1090]	
max of [sLi/ssBe/ssssBe/ssBH/sssB/ssssB/sCH3/dCH2/ssCH2/tch]	<u>Mordred</u>	[Descriptors] index [1090-1100]	

Name	Reference	Computable	description
max of [dsCH/aaCH/sssCH/ddC/tsC/dssC/aasC/aaaC/ssssC/sNH3/sNH2]	Mordred	[Descriptors] index [1101-1110]	
max of [ssNH2/dNH/ssNH/aaNH/tN/sssNH/dsN/aaN/sssN/ddsN/aasN]	Mordred	[Descriptors] index [1111-1121]	
max of [ssssN/sOH/dO/ssO/aaO/sF/sSiH3/ssSiH2/sssSiH/ssssSi]	Mordred	[Descriptors] index [1121-1131]	
max of [sPH2/ssPH/sssP/dsssP/sssssP/sSH/dS/ssS/aaS/dssS/ddssS]	Mordred	[Descriptors] index [1132-1142]	
max of [sCl/sGeH3/ssGeH2/sssGeH/ssssGe/sAsH2//ssAsH/sssAs]	Mordred	[Descriptors] index [1143-1150]	
max of [sssdAs/sssssAs/sSeH/dSe/ssSe/aaSe/dssSe/ddssSe/sBr]	<u>Mordred</u>	[Descriptors] index [1151-1159]	
max of [sSnH3/ssSnH2/sssSnH/ssssSn/sI/sPbH3/ssPbH2/sssPbH/ssssPb]	Mordred	[Descriptors] index [1160-1168]	
min of [sLi/ssBe/ssssBe/ssBH/sssB/ssssB/sCH3/dCH2/ssCH2/tch]	<u>Mordred</u>	[Descriptors] index [1169-1178]	
min of [dsCH/aaCH/sssCH/ddC/tsC/dssC/aasC/aaaC/ssssC/sNH3/sNH2]	<u>Mordred</u>	[Descriptors] index [1179-1189]	
min of [ssNH2/dNH/ssNH/aaNH/tN/sssNH/dsN/aaN/sssN/ddsN/aasN]	<u>Mordred</u>	[Descriptors] index [1190-1200]	
min of [ssssN/sOH/dO/ssO/aaO/sF/sSiH3/ssSiH2/sssSiH/ssssSi]	<u>Mordred</u>	[Descriptors] index [1202-1210]	
min of [sPH2/ssPH/sssP/dsssP/sssssP/sSH/dS/ssS/aaS/dssS/ddssS]	<u>Mordred</u>	[Descriptors] index [1211-1221]	
min of [sCl/sGeH3/ssGeH2/sssGeH/ssssGe/sAsH2//ssAsH/sssAs]	<u>Mordred</u>	[Descriptors] index [1222-1229]	
min of [sssdAs/sssssAs/sSeH/dSe/ssSe/aaSe/dssSe/ddssSe/sBr]	<u>Mordred</u>	[Descriptors] index [1230-1238]	
min of [sSnH3/ssSnH2/sssSnH/ssssSn/sI/sPbH3/ssPbH2/sssPbH/ssssPb]	Mordred	[Descriptors] index [1239-1247]	

Name	Reference	Computable	description
eccentric connectivity index	<u>Mordred</u>	[Descriptors] index 1248	
[/averaged] ETA core count	<u>Mordred</u>	[Descriptors] index [1249-1250]	ETA stands for Ethanolamine
ETA shape index (type: [p/y/x])	<u>Mordred</u>	[Descriptors] index [1251-1253]	ETA stands for Ethanolamine
	<u>Mordred</u>	[Descriptors] index 1254	
averaged	<u>Mordred</u>	[Descriptors] index 1255	
sigma contribution to	<u>Mordred</u>	[Descriptors] index 1256	
averaged sigma contribution to	<u>Mordred</u>	[Descriptors] index 1257	
nonsigma contribution to	<u>Mordred</u>	[Descriptors] index 1258	
averaged nonsigma contribution to	<u>Mordred</u>	[Descriptors] index 1259	
delta contribution to	<u>Mordred</u>	[Descriptors] index 1260	
averaged delta contribution to	<u>Mordred</u>	[Descriptors] index 1261	
valence electron mobile count	<u>Mordred</u>	[Descriptors] index [1254-1261]	
[/averaged/local/averaged local] ETA composite index for reference graph	<u>Mordred</u>	[Descriptors] [index 1262-1269]	ETA stands for Ethanolamine
[/averaged/local/averaged local] ETA functionality index	<u>Mordred</u>	[Descriptors] [index 1270-1273]	ETA stands for Ethanolamine
[/averaged] ETA branching index [/(use ring count)]	Mordred	[Descriptors] [index 1274-1277]	ETA stands for Ethanolamine

Name	Reference	Computable	description
ETA delta alpha (type: [A/B])	<u>Mordred</u>	[Descriptors] [index 1278-1279]	ETA stands for Ethanolamine
ETA epsilon (type: [1-5])	Mordred	[Descriptors] [index 1280-1284]	ETA stands for Ethanolamine
ETA delta epsilon (type: [A/B/C/D])	<u>Mordred</u>	[Descriptors] [index 1285-1288]	ETA stands for Ethanolamine
[/averaged] ETA delta beta	<u>Mordred</u>	[Descriptors] [index 1289-1290]	ETA stands for Ethanolamine
ETA psi	<u>Mordred</u>	[Descriptors] index 1291	ETA stands for Ethanolamine
ETA delta psi (type: [A/B])	<u>Mordred</u>	[Descriptors] [index 1292-1293]	ETA stands for Ethanolamine
fragment complexity	<u>Mordred</u>	[Descriptors] index 1294	
molecular framework ratio	<u>Mordred</u>	[Descriptors] index 1295	
geometric [diameter/radius]	<u>Mordred</u>	[Descriptors] [index 1296-1297]	
[geometrical shape/geometric Petitjean] index	<u>Mordred</u>	[Descriptors] [index 1298-1299]	
[heavy atom/heavy atom pair/pair] gravitational index	Mordred	[Descriptors] [index 1300-1303]	
number of hydrogen bond [acceptor/donor]	<u>Mordred</u>	[Descriptors] [index 1304-1305]	
[0-5]-ordered neighborhood [/total]	<u>Mordred</u>	[Descriptors] [index 1306-1317]	
[0-5]-ordered structural	<u>Mordred</u>	[Descriptors] [index 1318-1323]	
[0-5]-ordered bonding	<u>Mordred</u>	[Descriptors] [index 1324-1329]	

Name	Reference	Computable	description
[0-5]-ordered complementary	Mordred	[Descriptors] [index 1330-1335]	
[0-5]-ordered modified	<u>Mordred</u>	[Descriptors] [index 1336-1341]	
[0-5]-ordered Z-modified	<u>Mordred</u>	[Descriptors] [index 1342-1347]	
information content	<u>Mordred</u>	[Descriptors] index [1306-1347]	
kappa shape index [1-3]	<u>Mordred</u>	[Descriptors] [index 1348-1350]	
Lipinski rule of five	<u>Mordred</u>	[Descriptors] index 1351	
Ghose filter	<u>Mordred</u>	[Descriptors] index 1352	
Filter-it™ LogS	<u>Mordred</u>	[Descriptors] index 1353	•••
McGowan volume	<u>Mordred</u>	[Descriptors] index 1354	•••
3D-MoRSE (distance = [1-32])	<u>Mordred</u>	[Descriptors] [index 1355-1386]	
3D-MoRSE weighted by [mass/vdw volume](distance = [1-32])	<u>Mordred</u>	[Descriptors] [index 1387-1450]	
3D-MoRSE weighted by [sanderson EN/polarizability](distance = [1-32])	<u>Mordred</u>	[Descriptors] [index 1451-1514]	
Labute's Approximate Surface Area	<u>Mordred</u>	[Descriptors] index 1515	
MOE Charge VSA Descriptor [1-13]	Mordred	[Descriptors] [index 1516-1528]	
$(-\inf < x < -0.3)$	<u>Mordred</u>	[Descriptors] index 1516	

00.0 1.202 1 20.01		proportion	
Name	Reference	Computable	description
$(-0.3 \le x < -0.25)$	<u>Mordred</u>	[Descriptors] index 1517	
(-0.25 <= x <-0.2)	<u>Mordred</u>	[Descriptors] index 1518	
MOE MR VSA Descriptor [1-9]	Mordred	[Descriptors] [index 1529-1537]	
$(-\inf < x < 1.29)$	<u>Mordred</u>	[Descriptors] index 1529	
(1.29 <= x < 1.82)	Mordred	[Descriptors] index 1530	
(1.82 <= x < 2.24)	<u>Mordred</u>	[Descriptors] index 1531	
($2.24 \le x \le 2.45$)	<u>Mordred</u>	[Descriptors] index 1532	
(2.45 <= x < 2.75)	<u>Mordred</u>	[Descriptors] index 1533	
($2.75 \le x \le 3.05$)	<u>Mordred</u>	[Descriptors] index 1534	
($3.05 \le x \le 3.63$)	<u>Mordred</u>	[Descriptors] index 1535	
($3.63 \le x \le 3.80$)	<u>Mordred</u>	[Descriptors] index 1536	
($3.80 \le x \le 4.00$)	<u>Mordred</u>	[Descriptors] index 1537	
MOE logP VSA Descriptor [1-11]	<u>Mordred</u>	[Descriptors] [index 1538-1548]	
$(-\inf < x < -0.40)$	<u>Mordred</u>	[Descriptors] index 1538	
$(-0.40 \le x < -0.20)$	Mordred	[Descriptors] index 1539	

00.0 1.202 1 20.01		proportion	
Name	Reference	Computable	description
$(-0.20 \le x \le 0.00)$	<u>Mordred</u>	[Descriptors] index 1540	
($0.00 \le x \le 0.10$)	<u>Mordred</u>	[Descriptors] index 1541	
($0.10 \le x \le 0.15$)	<u>Mordred</u>	[Descriptors] index 1542	
($0.15 \le x \le 0.20$)	<u>Mordred</u>	[Descriptors] index 1543	
($0.20 \le x \le 0.25$)	<u>Mordred</u>	[Descriptors] index 1544	
($0.25 \le x \le 0.30$)	<u>Mordred</u>	[Descriptors] index 1545	
($0.30 \le x \le 0.40$)	<u>Mordred</u>	[Descriptors] index 1546	
($0.40 \le x < 0.50$)	<u>Mordred</u>	[Descriptors] index 1547	
($0.50 \le x \le 0.60$)	<u>Mordred</u>	[Descriptors] index 1548	
EState VSA Descriptor [1-10]	<u>Mordred</u>	[Descriptors] [index 1549-1558]	
$(-\inf < x < -0.39)$	<u>Mordred</u>	[Descriptors] index 1549	
(-0.39 <= x < -0.29)	<u>Mordred</u>	[Descriptors] index 1550	
$(-0.29 \le x \le 0.72)$	<u>Mordred</u>	[Descriptors] index 1551	
(0.72 <= x < 1.17)	<u>Mordred</u>	[Descriptors] index 1552	
(1.17 <= x < 1.54)	<u>Mordred</u>	[Descriptors] index 1553	

30.04.2024 10.37		properties1.11tm	
Name	Reference	Computable	description
(1.54 <= x < 1.81)	<u>Mordred</u>	[Descriptors] index 1554	
(1.81 <= x < 2.05)	<u>Mordred</u>	[Descriptors] index 1555	
($2.05 \le x \le 4.69$)	<u>Mordred</u>	[Descriptors] index 1556	
($4.69 \le x \le 9.17$)	<u>Mordred</u>	[Descriptors] index 1557	
($9.17 \le x \le 15.00$)	<u>Mordred</u>	[Descriptors] index 1558	
VSA EState Descriptor [1-9]	<u>Mordred</u>	[Descriptors] [index 1559-1567]	
$(-\inf < x < 4.78)$	<u>Mordred</u>	[Descriptors] index 1559	
($4.78 \le x \le 5.00$)	<u>Mordred</u>	[Descriptors] index 1560	
($5.00 \le x \le 5.41$)	<u>Mordred</u>	[Descriptors] index 1561	
($5.41 \le x \le 5.74$)	<u>Mordred</u>	[Descriptors] index 1562	
($5.74 \le x \le 6.00$)	<u>Mordred</u>	[Descriptors] index 1563	
($6.00 \le x \le 6.07$)	<u>Mordred</u>	[Descriptors] index 1564	
($6.07 \le x \le 6.45$)	<u>Mordred</u>	[Descriptors] index 1565	
($6.45 \le x \le 7.00$)	<u>Mordred</u>	[Descriptors] index 1566	
($7.00 \le x \le 11.00$)	Mordred	[Descriptors] index 1567	

Name	Reference	Computable	description
molecular distance edge between	<u>Mordred</u>	[Descriptors] [index 1568-1586]	
primary C and [primary C/secondary C/tertiary C/quaternary C]	Mordred	[Descriptors] [index 1568-1571]	
secondary C and [secondary C/tertiary C/quaternary C]	<u>Mordred</u>	[Descriptors] [index 1572-1574]	
tertiary C and [tertiary C/quaternary C]	<u>Mordred</u>	[Descriptors] [index 1575-1576]	
quaternary C and quaternary C	<u>Mordred</u>	[Descriptors] index 1577	
primary O and [primary O/secondary O]	<u>Mordred</u>	[Descriptors] [index 1578-1579]	
secondary O and secondary O	<u>Mordred</u>	[Descriptors] index 1580	
primary N and [primary N/secondary N/tertiary N]	<u>Mordred</u>	[Descriptors] [index 1581-1583]	
secondary N and [secondary N/tertiary N]	<u>Mordred</u>	[Descriptors] [index 1584-1585]	
tertiary N and tertiary N	<u>Mordred</u>	[Descriptors] index 1586	
[/averaged] molecular ID [/on [H/C/N/O] atoms]	<u>Mordred</u>	[Descriptors] [index 1587-1598]	
moment of inertia (axis = $[X/Y/Z]$)	<u>Mordred</u>	[Descriptors] [index 1599-1601]	
PBF	<u>Mordred</u>	[Descriptors] index 1602	
[2-10]-ordred patch count	<u>Mordred</u>	[Descriptors] [index 1603-1611]	
10-ordred total patch count	<u>Mordred</u>	[Descriptors] index 1612	
Marsili/Gasteiger partial charges	RDKit		Electron Distribution

Name	Reference	Computable	description
BalabanJ	<u>RDKit</u>	Graphdescriptor module	Atom indexing in molecule
BertzCT	RDKit	Graphdescriptor module	Tuple of two numbers. First number is correlated with complexity of bonds, second number is correlated with complexity of the distribution of heteroatoms.
IPC	RDKit	Graphdescriptor module	This returns the information content of the coefficients of the characteristic polynomial of the adjacency matrix of a hydrogen-suppressed graph of a molecule.
Kappa [1-3]	<u>RDKit</u>	Graphdescriptor module	Hallkier Kappa values. Note that these are NOT broken.
Phi	<u>RDKit</u>	NC	Kier Phi value for a molecule From Quantitative Structure-Activity Relationships
Chi0n - Chi4n	RDKit	Graphdescriptor module	Similar to Hall Kier ChiXv, but uses nVal instead of valence.\ This makes a big difference after we get out of the first row.
Chi0v - Chi4v	<u>RDKit</u>	Graphdescriptor module	From equations (5),(9) and (10) of Rev. Comp. Chem. vol 2, 367-422, (1991)
MolLogP	<u>RDKit</u>	Crippen module	Wildman-Crippen LogP for a molecule
MolMR	<u>RDKit</u>	Crippen module	Wildman-Crippen MR for a molecule
MolWT	RDKit	Descriptors module	Sum of atomic weights in the molecule
ExactMolWT	<u>RDKit</u>	Descriptors module	More accurate molecular weight
Heavy Atom count	<u>RDKit</u>	Lipinski module	Counter of atoms with atomic weight above some threshold in the molecule
Heavy Atom MolWT	RDKit	Descriptors module	We assume this means the the ratio between the sum of heavy atom weights and the weight of the whole molecule
NHOH group count	<u>RDKit</u>	Lipinski module	Counts the number of NHOH groups in the molecule
NOCount	<u>RDKit</u>	Descriptors module	Nitrogen Oxide Grp counter
NumHAcceptors	<u>RDKit</u>	Lipinski module	The number of proton acceptors in the molecule
NumHDonators	<u>RDKit</u>	Lipinski module	Number of bonds in the molecule that are weak enough to give off a proton.

Name	Reference	Computable	description
NumHeteroatoms	<u>RDKit</u>	Lipinski module	Number of atoms that are NOT carbon or hydrogen
NumRotatableBonds	<u>RDKit</u>	Lipinski module	Counter of bonds in the molecule that can rotate. Usually single electron bonds.
NumValenceElectrons	<u>RDKit</u>	Descriptors module	Number of electrons in the outer most shell of an atom modulo 8.
NumAmideBonds	RDKit	NC	Amide bond counter. Google what an amide is.
Num{Aromatic,Saturated,Aliphatic}Rings	<u>RDKit</u>	NC	Usually something like a benzene ring with delocalized electrons.
RingCount	RDKit	Lipinski module	Counter of cyclical rings in molecule
FractionCSP3	RDKit	NC	The fraction of C atoms that are SP3 hybridized
NumSpiroAtoms	RDKit	NC	Atoms shared between rings that share exactly one atom
NumBridgeheadAtoms	RDKit	NC	atoms shared between rings that share at least two bonds
TPSA	RDKit	MolSurf module	Topological Polar Surface Area
LabuteASA	RDKit	MolSurf module	Accessible Surface Area to a solvent
PEOE_VSA [1-14]	<u>RDKit</u>	MolSurf module	Acronym explanation: Partial Equalization of Orbital Electronegativity and Volume, Area and Shape
SMR_VSA [1-10]	<u>RDKit</u>	MolSurf module	Surface-Weighted Mean Refractivity. Captures information related to the polarizability
SlogP_VSA [1-12]	<u>RDKit</u>	MolSurf module	the SlogP_VSA descriptors help us understand how a molecule's lipophilicity influences its behavior, especially in terms of interactions with other molecules or surfaces
EState_VSA [1-11]	RDKit	EState.EState_VSA	Used to describe the electronic state of a molecule.
MQNs	RDKit	NC	Molecular Quantum Numbers
Topliss fragments	RDKit	NC	Could not find a good description for this descriptor
Autocorr2D	<u>RDKit</u>	NC	From Todeschini and Consoni "Descriptors from Molecular Geometry" Handbook of Chemoinformatics
BCUT2D	<u>RDKit</u>	NC	From Pearlman and Smith in "3D-QSAR and Drug design: Recent Advances" (1997)
fr_Al_COO	<u>RDKit</u>	fragments module	Number of aliphatic carboxylic acids

Name	Reference	Computable	description	
fr_Al_OH	RDKit	fragments module	Number of aliphatic hydroxyl groups	
fr_Al_OH_noTert	<u>RDKit</u>	fragments module	Number of aliphatic hydroxyl groups excluding tert-OH	
fr_ArN	<u>RDKit</u>	fragments module	Number of N functional groups attached to aromatics	
fr_Ar_COO	<u>RDKit</u>	fragments module	Number of Aromatic carboxylic acide	
fr_Ar_N	<u>RDKit</u>	fragments module	Number of aromatic nitrogens	
fr_Ar_NH	<u>RDKit</u>	fragments module	Number of aromatic amines	
fr_Ar_OH	<u>RDKit</u>	fragments module	Number of aromatic hydroxyl groups	
fr_COO	<u>RDKit</u>	fragments module	Number of carboxylic acids	
fr_COO2	RDKit	fragments module	Number of carboxylic acids	
fr_C_O	RDKit	fragments module	Number of carbonyl O	
fr_C_O_noCOO	RDKit	fragments module	Number of carbonyl O, excluding COOH	
fr_C_S	RDKit	fragments module	Number of thiocarbonyl	
fr_HOCCN	<u>RDKit</u>	fragments module	Number of C(OH)CCN-Ctert-alkyl or C(OH)CCNcyclic	
fr_Imine	<u>RDKit</u>	fragments module	Number of Imines	
fr_NH0	<u>RDKit</u>	fragments module	Number of Tertiary amines	
fr_NH1	<u>RDKit</u>	fragments module	Number of Secondary amines	
fr_NH2	<u>RDKit</u>	fragments module	Number of Primary amines	
fr_N_O	<u>RDKit</u>	fragments module	Number of hydroxylamine groups	
fr_Ndealkylation1	RDKit	fragments module	Number of XCCNR groups	
fr_Ndealkylation2	<u>RDKit</u>	fragments module	Number of tert-alicyclic amines (no heteroatoms, not quinine-like bridged N)	
fr_Nhpyrrole	<u>RDKit</u>	fragments module	Number of H-pyrrole nitrogens	
fr_SH	<u>RDKit</u>	fragments module	Number of thiol groups	
fr_aldehyde	<u>RDKit</u>	fragments module	Number of aldehydes	
fr_alkyl_carbamate	<u>RDKit</u>	fragments module	Number of alkyl carbamates (subject to hydrolysis)	
fr_alkyl_halide	<u>RDKit</u>	fragments module	Number of alkyl halides	
fr_allylic_oxid	<u>RDKit</u>	fragments module	Number of allylic oxidation sites excluding steroid dienone	

Name	Reference	Computable	description	
fr_amide	RDKit	fragments module	Number of amides	
fr_amidine	RDKit	fragments module	Number of amidine groups	
fr_aniline	RDKit	fragments module	Number of anilines	
fr_aryl_methyl	<u>RDKit</u>	fragments module	Number of aryl methyl sites for hydroxylation	
fr_azide	<u>RDKit</u>	fragments module	Number of azide groups	
fr_azo	RDKit	fragments module	Number of azo groups	
fr_barbitur	<u>RDKit</u>	fragments module	Number of barbiturate groups	
fr_benzene	<u>RDKit</u>	fragments module	Number of benzene rings	
fr_benzodiazepine	<u>RDKit</u>	fragments module	Number of benzodiazepines with no additional fused rings	
fr_bicyclic	RDKit	fragments module	Bicyclic	
fr_diazo	RDKit	fragments module	Number of diazo groups	
fr_dihydropyridine	<u>RDKit</u>	fragments module	Number of dihydropyridines	
fr_epoxide	<u>RDKit</u>	fragments module	Number of epoxide rings	
fr_ester	<u>RDKit</u>	fragments module	Number of esters	
fr_ether	<u>RDKit</u>	fragments module	Number of ether oxygens (including phenoxy)	
fr_furan	<u>RDKit</u>	fragments module	Number of furan rings	
fr_guanido	<u>RDKit</u>	fragments module	Number of guanidine groups	
fr_halogen	<u>RDKit</u>	fragments module	Number of halogens	
fr_hdrzine	RDKit	fragments module	Number of hydrazine groups	
fr_hdrzone	<u>RDKit</u>	fragments module	Number of hydrazone groups	
fr_imidazole	RDKit	fragments module	Number of imidazole rings	
fr_imide	<u>RDKit</u>	fragments module	Number of imide groups	
fr_isocyan	RDKit	fragments module	Number of isocyanates	
fr_isothiocyan	RDKit	fragments module	Number of isothiocyanates	
fr_ketone	<u>RDKit</u>	fragments module	Number of ketones	
fr_ketone_Topliss	<u>RDKit</u>	fragments module	Number of ketones excluding diaryl, a,b-unsat. dienones, heteroatom on Calpha	
fr_lactam	<u>RDKit</u>	fragments module	Number of beta lactams	

Name	Reference	Computable	description	
fr_lactone	RDKit	fragments module	Number of cyclic esters (lactones)	
fr_methoxy	RDKit	fragments module	Number of methoxy groups -OCH3	
fr_morpholine	RDKit	fragments module	Number of morpholine rings	
fr_nitrile	RDKit	fragments module	Number of nitriles	
fr_nitro	<u>RDKit</u>	fragments module	Number of nitro groups	
fr_nitro_arom	<u>RDKit</u>	fragments module	Number of nitro benzene ring substituents	
fr_nitro_arom_nonortho	<u>RDKit</u>	fragments module	Number of non-ortho nitro benzene ring substituents	
fr_nitroso	<u>RDKit</u>	fragments module	Number of nitroso groups, excluding NO2	
fr_oxazole	<u>RDKit</u>	fragments module	Number of oxazole rings	
fr_oxime	<u>RDKit</u>	fragments module	Number of oxime groups	
fr_para_hydroxylation	<u>RDKit</u>	fragments module	Number of para-hydroxylation sites	
fr_phenol	RDKit	fragments module	Number of phenols	
fr_phenol_noOrthoHbond	<u>RDKit</u>	fragments module	Number of phenolic OH excluding ortho intramolecular Hbond substituents	
fr_phos_acid	<u>RDKit</u>	fragments module	Number of phosphoric acid groups	
fr_phos_ester	<u>RDKit</u>	fragments module	Number of phosphoric ester groups	
fr_piperdine	RDKit	fragments module	Number of piperdine rings	
fr_piperzine	RDKit	fragments module	Number of piperzine rings	
fr_priamide	RDKit	fragments module	Number of primary amides	
fr_prisulfonamd	RDKit	fragments module	Number of primary sulfonamides	
fr_pyridine	RDKit	fragments module	Number of pyridine rings	
fr_quatN	RDKit	fragments module	Number of quaternary nitrogens	
fr_sulfide	<u>RDKit</u>	fragments module	Number of thioether	
fr_sulfonamd	<u>RDKit</u>	fragments module	Number of sulfonamides	
fr_sulfone	<u>RDKit</u>	fragments module	Number of sulfone groups	
fr_term_acetylene	<u>RDKit</u>	fragments module	Number of terminal acetylenes	
fr_tetrazole	RDKit	fragments module	Number of tetrazole rings	
fr_thiazole	RDKit	fragments module	Number of thiazole rings	
fr_thiocyan	RDKit	fragments module	Number of thiocyanates	

Name	Reference	Computable	description
fr_thiophene	RDKit	fragments module	Number of thiophene rings
fr_unbrch_alkane	<u>RDKit</u>	fragments module	Number of unbranched alkanes of at least 4 members (excludes halogenated alkanes)
fr_urea	RDKit	fragments module	Number of urea groups
Dipole moment	<u>PyG</u> , <u>MolNet</u>	Not included	
Isotrope polarizability	<u>PyG</u> , <u>MolNet</u>	Not included	
Highest occupied molecular orbital energy	<u>PyG</u> , <u>MolNet</u>		
Lowest unoccupied molecular orbital energy	<u>PyG</u> , <u>MolNet</u>		
HOMO-LUMO gap	<u>PyG</u> , <u>MolNet</u>		
Electronic spatial extent	<u>PyG</u> , <u>MolNet</u>		
Zero point vibration energy	<u>PyG</u> , <u>MolNet</u>	autodE/ORCA	
Internal energy at 0K	<u>PyG</u> , <u>MolNet</u>		
Internal energy at 298.15K	<u>PyG</u> , <u>MolNet</u>	autodE/ORCA	
Enthalpy at 298.15K	<u>PyG</u> , <u>MolNet</u>	autodE/ORCA	
Free energy at 298.15K	<u>PyG</u> , <u>MolNet</u>	autodE/ORCA	
Heat capavity at 298.15K	<u>PyG</u> , <u>MolNet</u>		
Atomization energy at 0K	<u>PyG</u> , <u>MolNet</u>		

Name	Reference	Computable	description
Atomization energy at 298.15K	<u>PyG</u> , <u>MolNet</u>		
Atomization enthalpy at 298.15K	<u>PyG</u> , <u>MolNet</u>		
Atomization free energy at 298.15K	<u>PyG</u> , <u>MolNet</u>		

[Descriptors]: Calculator(descriptors, ignore_3D=False).descriptors