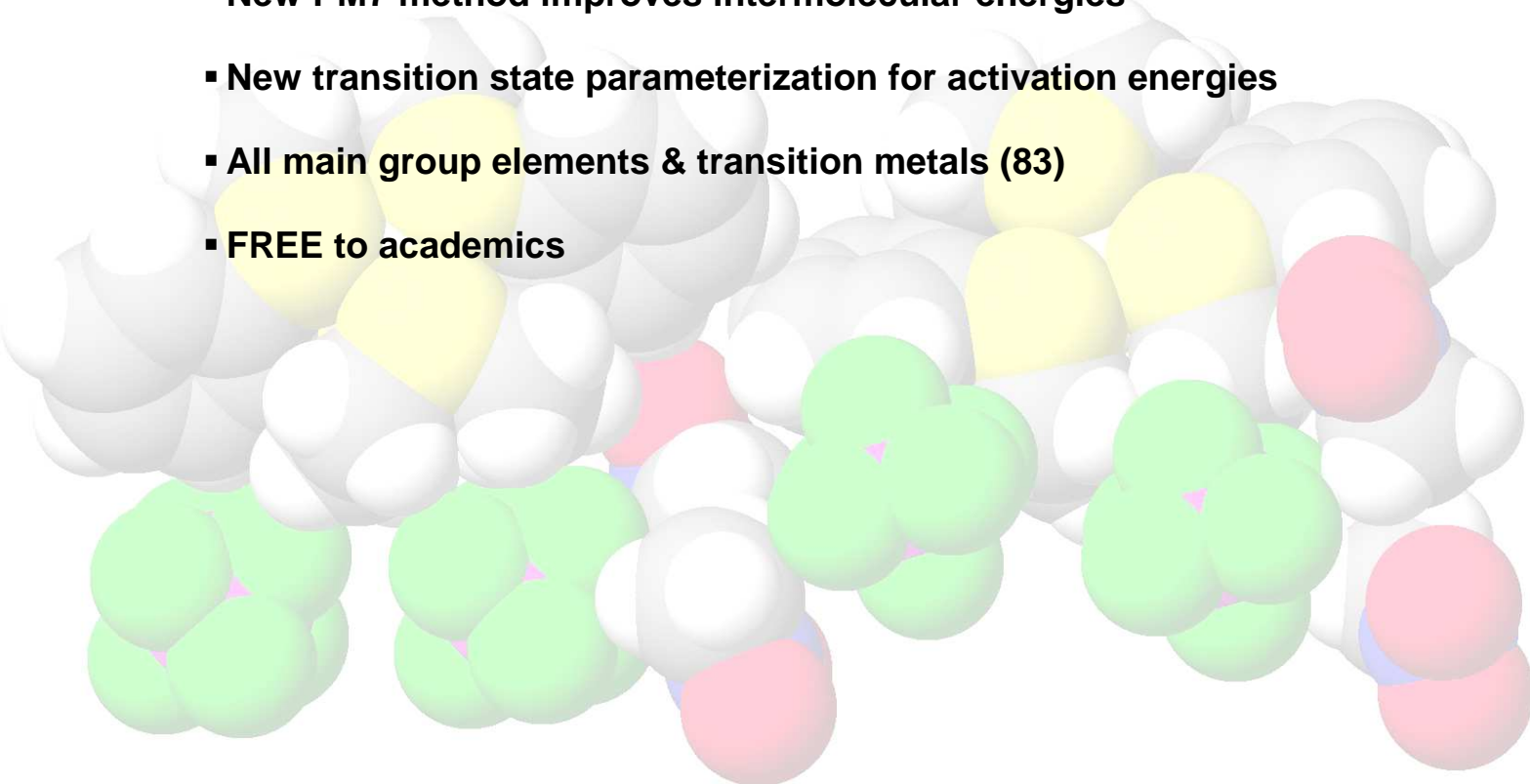




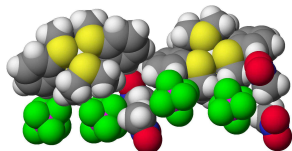
MOPAC2016TM

A practical quantum chemistry tool for modeling biological systems and crystals

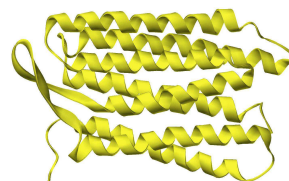
- Improved handling for co-crystals & biological systems
 - Fast optimizations up to 15,000 atoms, e.g. proteins, crystals
 - New PM7 method improves intermolecular energies
 - New transition state parameterization for activation energies
 - All main group elements & transition metals (83)
 - FREE to academics
- 

MOPAC2016™

- **Biological systems, crystals, co-crystals.....major improvements**



MOPAC2016 brings major improvements in the prediction of intermolecular interactions. This significantly improves geometries and energies of proteins, crystals, co-crystals, metal clusters, inorganics and other condensed phase systems.



- **NEW parameterization (PM7)**

The most widely used semiempirical quantum chemistry package, MOPAC®, has been updated with a new and more accurate parameterization (PM7) for all the main group elements and transition metals. The addition of Jurečka's dispersion function and integrated hydrogen-bond terms to PM7 significantly improves the prediction of intermolecular interactions. Experimental and *ab initio* data from over 9,000 compounds were used to develop the new PM7 method. MOPAC2016™ with PM7 represents only the second major improvement in methodology to MOPAC®, since PM3 was published in 1989; the first being PM6 in 2007.¹

Year	Method	# compounds used for parameterization	
1977	MNDO	39	experimental
1985	AM1	~200	experimental
1989	PM3	~500	experimental
2007	PM6	> 9,000	experimental & <i>ab initio</i>
2016	PM7	> 9,000	experimental & <i>ab initio</i>

- **More accurate heats of formation from PM7 for simple organic compounds**

Method	Average unsigned error (kcal/mol)	Root Mean Square Error (Kcal/mol)	Largest error (kcal/mol)
PM7	4.01	5.89	-44.4
PM6	4.42	6.16	-42.2
B3LYP 6-31G(d)	5.14	7.36	36.1
PM3	6.23	9.44	-135.6
HF 6-31G(d)	7.34	10.64	72.1
AM1	10.00	14.65	200.4

Comparison of errors in heats of formation for a set of [1,366 compounds](#) containing only C, H, O, N, F, Cl, S, P and Br.

MOPAC® is the most cited semiempirical program and has far more published accuracy data than any comparable program.¹ MOPAC® has been used to check for and correct errors in the published experimental heats of formation tables on the NIST website.^{2,3}

- **Giant molecule capability**

MOPAC2016's linear-scaling algorithm, MOZYME, allows geometry optimizations on closed shell systems of up to 15,000 atoms (e.g. proteins). Conventional MOPAC is limited to about 1,500 atoms.

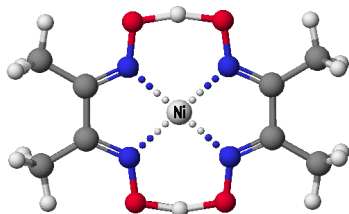
No. of atoms	Time for 1 SCF (minutes)		Memory (megabytes)	
	MOZYME	MOPAC	MOZYME	MOPAC
400	0.2	2.3	17	101
1,500	2.3	222.6	78	1,424
15,000	230.3	*222,600.0	1,026	*142,400

2.9 GHz machine

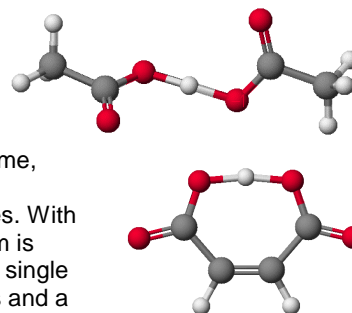
*estimated

- **More accurate hydrogen bonds from PM7**

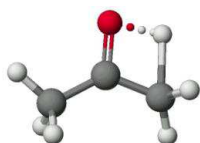
PM7 positions the bridging hydrogen approximately equidistant between the oxygen atoms in dicarboxylic acid anions such as



hydrogen diacetate and hydrogen maleate anions and, similarly, in nickel dimethylglyoxime, consistent with experimental observations and DFT geometries. With AM1 and PM3 the hydrogen atom is incorrectly displaced indicating a single bond to one of the oxygen atoms and a normal hydrogen bond to the other.



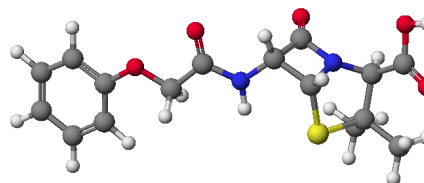
- **New activation energy parameterization for transition states**



Reaction barrier heights have been parameterized using published coupled cluster high-level *ab initio* calculations on the PM7 transition state and related ground state geometries (reactants and products). The new method, PM7-TS reduces activation energy errors by a factor of approximately 3, relative to PM6 (Average Unsigned Error for PM7-TS: 3.8 Kcal/mol; PM6: 12.3 Kcal/mol)

- **Fast descriptor generation for QSAR**

New algorithms in MOPAC2016™ facilitate fast property prediction for screening libraries of drug-sized molecules for a wide range of properties, including pKa. The speed of MOPAC2016™ and improved accuracy of PM7 are particularly valuable for generating electronic descriptors for quantitative structure-property relationships (QSAR).

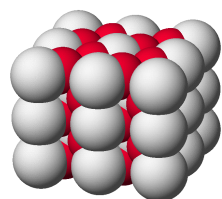


- **ALL main group elements & transition metals parameterized with PM7**

PM7 in MOPAC2016™ includes parameters for a wider range of elements than any other semiempirical quantum chemistry program. All main group elements and transition metals up to Bismuth are now parameterized in PM7. Further, MOPAC2016™ includes new elements for PM3 & AM1.

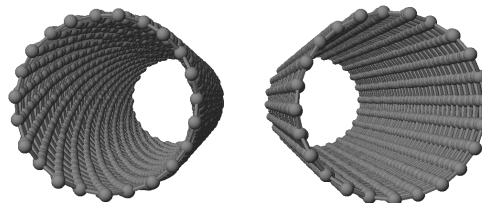
PM7: (83) H, He, Li, Be, B, C, N, O, F, Ne, Na, Mg, Al, Si, P, S, Cl, Ar, K, Ca, Sc, Ti, V, Cr, Mn, Fe, Co, Ni, Cu, Zn, Ga, Ge, As, Se, Br, Kr, Rb, Sr, Y, Zr, Nb, Mo, Tc, Ru, Rh, Pd, Ag, Cd, In, Sn, Sb, Te, I, Xe, Cs, Ba, La, Lu, Hf, Ta, W, Rh, Os, Ir, Pt, Au, Hg, Tl, Pb, Bi, + 15 lanthanide sparkles⁴

- **Co-crystals, crystals, surfaces & polymers, with periodic boundaries**



MOPAC2016™ can handle extended solids including straight-chain polymers such as bucky tubes (one-dimensional), surfaces (two-dimensional), and crystals (three-dimensional) with periodic boundary conditions. This approach eliminates the problems of edge effects and facilitates the calculation of

properties on extended systems that cannot be handled adequately by other quantum chemistry packages.



• MOPAC2016™ specifications

Properties

Thermodynamic properties (heats of formation, entropies, free energies, heat capacities), vibrational (IR) spectra (including isotope effects), dipole moments, ionization potentials, partial charges, bond orders, static and frequency-dependent polarizabilities and hyperpolarizabilities, pKa, molecular orbitals, band gaps, electron densities, electrostatic potentials, transition states, intrinsic reaction coordinates, dynamic reaction coordinates, excited state geometries and energetics, compressibility, density, and thermodynamic properties of crystals.

Types of calculations

Open and closed-shell Hartree-Fock methods: Restricted (RHF), Unrestricted (UHF), Restricted open-shell (ROHF), singlet, doublet, triplet, quartet, quintet, sextet, etc. (ground and excited states)
Geometry optimization: Eigenvector following (EF), BFGS, NLLSQ, SIGMA
Locating Transition states: Saddle, Path, Grid
Refining Transition states: Transition State Eigenvector Following (TS), NLLSQ, SIGMA
Vibrational frequency calculation
Solvent effects: Conductor-like Screening Model (COSMO) including d-orbitals and excited states
Configuration interaction (CI) includes dynamic and static Jahn-Teller, symmetry analysis
Periodic boundary conditions: 1, 2, & 3 dimensional translation vectors (Tv)

Hamiltonians

PM7: (83) H, He, Li, Be, B, C, N, O, F, Ne, Na, Mg, Al, Si, P, S, Cl, Ar, K, Ca, Sc, Ti, V, Cr, Mn, Fe, Co, Ni, Cu, Zn, Ga, Ge, As, Se, Br, Kr, Rb, Sr, Y, Zr, Nb, Mo, Tc, Ru, Rh, Pd, Ag, Cd, In, Sn, Sb, Te, I, Xe, Cs, Ba, La, Lu, Hf, Ta, W, Rh, Os, Ir, Pt, Au, Hg, Tl, Pb, Bi +15 lanthanide sparkles⁴ (uses PM6 sparkles)
PM6: (83) H, He, Li, Be, B, C, N, O, F, Ne, Na, Mg, Al, Si, P, S, Cl, Ar, K, Ca, Sc, Ti, V, Cr, Mn, Fe, Co, Ni, Cu, Zn, Ga, Ge, As, Se, Br, Kr, Rb, Sr, Y, Zr, Nb, Mo, Tc, Ru, Rh, Pd, Ag, Cd, In, Sn, Sb, Te, I, Xe, Cs, Ba, La, Lu, Hf, Ta, W, Rh, Os, Ir, Pt, Au, Hg, Tl, Pb, Bi +15 lanthanide sparkles⁴
PM3: (57) H, He, Li, Be, B, C, N, O, F, Ne, Na, Mg, Al, Si, P, S, Cl, Ar, K, Ca, Zn, Ga, Ge, As, Se, Br, Kr, Rb, Sr, Cd, In, Sn, Sb, Te, I, Xe, Cs, Ba, Hg, Tl, Pb, Bi, + 15 lanthanide sparkles⁴
AM1: (57) H, He, Li, Be, B, C, N, O, F, Ne, Na, Mg, Al, Si, P, S, Cl, Ar, K, Ca, Zn, Ga, Ge, As, Se, Br, Kr, Rb, Sr, Mo, In, Sn, Sb, Te, I, Xe, Cs, Ba, Hg, Tl, Pb, Bi, + 15 lanthanide sparkles⁴
MNDO: (17) H, B, C, N, O, F, Na, Al, Si, P, S, Cl, Zn, Br, Cd, I, Hg
RM1²: (10) H, C, N, O, P, S, F, Cl, Br, I

User Interface

MOPAC2016™ is command driven and is designed to work with a wide range of third-party graphical user interfaces (GUI)s. To allow GUIs to read MOPAC results, a special file, <test>.aux, written in human readable markup language is now written (keyword AUX). MOPAC2016™ also reads and writes other file formats such as PDB and Gaussian®.

Platforms

MOPAC2016™ is available to download for Windows®, LINUX and Macintosh. For source code, please contact MrMOPAC@OpenMOPAC.net

• FREE to academics

MOPAC2016™ is available for free to bona-fide academics for teaching <http://www.openmopac.net/MOPAC2016.html> and not-for-profit R&D. Academics may download MOPAC2016 at

MOPAC2012™ prices (Windows®)	Academic	Government	Industry
1 permanent license	free	\$2,500	\$5,000
5 permanent licenses	-	\$5,000	\$10,000
Annual site license	-	\$5,000	\$10,000

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2. Stewart, J. J. P. (2004). J. Mol. Modelling 10: 6-12.
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5. Rocha, G. B. et al (2006). J. Comp. Chem. 27(10): 1101-1111