Molecule

Molecule()

Molecule (Molecule mol)

Molecule (Molecule mol, List<Bool> atoms,

List<Bool> bonds, Bool adjustHydrogens)

Int numAtoms()

Iter<Atom> atoms()

Atom atom(Int index)

Atom addAtom()

Void removeAtom(Atom atom)

Int numBonds()

Iter<Bond> bonds()

Bond bond(Int index)

Bond bond(Atom source, Atom target)

Bond addBond()

Void removeBond(Bond bond)

Void clear()

Bond

Int index()

Atom source()

Atom target()

Atom other(Atom atom)

Int order()

Bool isAromatic()

Void setAromatic(Bool value)

Void setOrder(Int value)

Ring

Int size()

List<Atom> atoms()

Atom atom(Int index)

Bool containsAtom(Atom atom)

List<Bond> bonds()

Bond bond(Int index)

Bool containsBond(Bond bond)

Bool isAromatic()

Fingerprint [python]

Fingerprint(Int numWords)

String bin(Bool spaces = true)

String **hex**()

Void zero()

Int count()

Int count(Int begin, Int end)

Bool __getitem__(Int index)
Void __setitem__(Int index, Bool value)

Int numWords()

Atom

Int index() Bool isOxygen()

Int degree() Bool isPhosphorus()

Iter<Bond> bonds() Bool isSulfur()

Iter<Atom> nbrs() Int heavyDegree() Bool isAromatic()

Int boSum() Int element() Int valence()

Int mass() Int connectivity()

Int hydrogens() Void **setAromatic**(Bool **value**) Int charge()

Void **setElement**(Int **value**) Bool isHydrogen() Void **setMass**(Int **value**)

Bool isCarbon() Void setHydrogens(Int value)

Bool isNitrogen() Void **setCharge**(Int **value**)

Element

String element(Int element)

Int averageMass(Int element)

Bool addHydrogens(Int element)

Int valence(Int element, Int charge,

int degree)

RingSet

RingSet(Molecule mol)

Int size()

List<Ring> rings()

Ring ring(Int index)

Bool isAtomInRing(Atom atom)

Bool isAtomInRingSize(Atom atom, Int size)

Int numRingNbrs(Atom atom)

Int numRings(Atom atom)

Bool isBondInRing(Bond bond)

Bool isBondInRingSize(Bond bond, Int size)

DistanceMatrix

String __str__()

Int dim()

Int infinity()

Int operator()(Int i, Int j)

Int __call__(Int i, Int j)

Error

String what() [C++]

operator bool() [C++]

Bool __nonzero__() [python]

String __str__() [python]

Smiles

Enum Flags: None, Mass, Charge, Hydrogens, Order, All

Smiles()

Bool **read**(String **smiles**, Molecule **mol**)

String write(Molecule mol, Flags flags = All)

String write(Molecule mol, List<Index> order, Flags flags = All)

String writeCanonical(Molecule mol)

Error error()

Smarts

Smarts()

Mapping implementations: NoMapping.match, CountMapping.count,

SingleMapping.map, MappingList.maps

Error error()

Bool requiresCycles()

Bool **init**(String **smarts**)

Bool requiresExplicitHydrogens()

Bool findMapping(Molecule mol, RingSet rings, Mapping mapping, Bool uniqueComponents = false)

Bool findMapping(Molecule mol, Mapping mapping, Bool uniqueComponents = false)

Bool find(Molecule mol, RingSet rings, Bool uniqueComponents = false)

Bool find(Molecule mol, Bool uniqueComponents = false)

Smirks

Smirks()

Void setFixMass(Bool value)

Void setFixHydrogens(Bool value)

Bool **init**(String **smirks**)

Bool init(String reactants, String products)

SmirksError error()

Bool requiresCycles()

Bool requiresExplicitHydrogens()

Bool apply(Molecule mol, RingSet rings)

Bool apply(Molecule mol)

List<Molecule **mol**, RingSet **rings**,

Int min = 1, Int max = 1)

List<Molecule> react(Molecule mol,

Int min = 1, Int max = 1)

Memory Mapped Molecule File

MemoryMappedMoleculeFile()

MemoryMappedMoleculeFile(String filename)

Void **load**(String **filename**)

Int numMolecules()

Bool readMolecule(Int index, Molecule mol)

Void error()

SmirksError

Enum Type: None, NoReaction, ReactantSmarts,

ProductSmarts, AtomClassPairWise,

ProductContainsOr, ProductContainsNot, InvalidProductBond, ProductConflict

Type type()

String what()

MoleculeFile

MoleculeFile()

MoleculeFile(String filename)

Void load(String filename)

Int numMolecules()

Bool readMolecule (Molecule mol)

Bool readMolecule(Int index, Molecule mol)

Void close()

Void error()

MoleculeOutputFile

MoleculeOutputFile(String filename) Bool writeMolecule (Molecule mol)

Void error()

Free Functions

HYDROGENS

Void make hydrogens explicit(Molecule mol)

Void make hydrogens implicit(Molecule mol)

Void reset_implicit_hydrogens(Molecule mol)

Void reset_implicit_hydrogens(Molecule mol, Atom atom)

FINGERPRINTS

Int BitePerWord

Int bitvec num words for bits(Int bits)

Void **bitvec_zero**(Fingerprint **fp**)

Fingerprint bitvec copy(Fingerprint fp)

Bool **bitvec_get**(Int **index**, Fingerprint **fp**)

Void **bitvec_set**(Int **index**, Fingerprint **fp**)

Void **bitvec_reset**(Int **index**, Fingerprint **fp**)

Int bitvec_count(Fingerprint fp)

Int bitvec_count(Fingerprint fp,

Int begin, Int end)

String **bitvec_to_binary**(Fingerprint **fp**)

String **bitvec_to_hex**(Fingerprint **fp**)

Fingerprint bitvec from binary(String bin)

Fingerprint **bitvec_from_hex**(String **hex**)

Int bitvec union count(Fingerprint fp1, Fingerprint fp2)

Bool bitvec is subset superset(Fingerprint subset, Fingerprint superset)

Real bitvec tanimoto(Fingerprint fp1, Fingerprint fp2) Real bitvec tanimoto(Fingerprint fp1, Fingerprint fp2, Int count1, Int count2)

Real bitvec_cosine(Fingerprint fp1, Fingerprint fp2) Real bitvec_cosine(Fingerprint fp1, Fingerprint fp2,

Int count1, Int count2)

Real **bitvec_hamming**(Fingerprint **fp1**, Fingerprint **fp2**)

Real bitvec_hamming(Fingerprint fp1, Fingerprint fp2, Int count1, Int count2)

Real bitvec forbes(Fingerprint fp1, Fingerprint fp2) Real bitvec_forbes(Fingerprint fp1, Fingerprint fp2,

Int count1, Int count2)

Real bitvec russell rao(Fingerprint fp1, Fingerprint fp2)

DFSVisitor [base class]

Void initialize(Molecule mol)

Void component(Int component)

Void atom(Molecule mol, Atom prev, Atom atom)

Void bond(Molecule mol, Atom prev, Bond bond)

Void **backtrack**(Molecule **mol**, Atom **atom**)

Void back_bond(Molecule mol, Bond bond)

Bool stop()

Default implementations: DFSAtomOrderVisitor,

DFSBondOrderVisitor, DFSDebugVisitor,

DFSClosureRecorderVisitor

AtomInvariant [base class]

Int operator()(Molecule mol, Atom atom) [C++]
Int __call__(Molecule mol, Atom atom) [python]

DefaultAtomInvariant

Enum Invariants: None, Element, Mass, Charge, Degree, Aromatic, All

DefaultAtomInvariant(Invariants invariants = All)

Dijkstra

Dijkstra(Molecule mol, Atom source,

Bool preferSmallerIndexPaths = false)

Int infinity()

Int distance(Atom target)

List<Atom> path(Atom target)

BFSVisitor [base class]

Void initialize(Molecule mol)

Void component(Int component)

Void depth(Int depth)

Void atom(Molecule mol, Atom prev, Atom atom)

Void **bond**(Molecule **mol**, Atom **prev**, Bond **bond**)

Void back_bond(Molecule mol, Bond bond)

Bool stop()

Default implementations: BFSAtomOrderVisitor,

BFSBondOrderVisitor, BFSDebugVisitor,

BFSClosureRecorderVisitor

BondInvariant

Int operator()(Molecule mol, Bond bond)
Int __call__(Molecule mol, Bond bond)

DefaultBondInvariant

Enum Invariants: None, Order, Aromatic, All DefaultAtomInvariant(Invariants invariants = All)

SubgraphCallback

Void operator()(Subgraph subgraph)
Void __call__(Subgraph subgraph)

Subgraph

List<Bool> hashable()
List<Bool> atoms, bonds

Algorithms

DEPTH-FIRST SEARCH

Void depth_first_search(Molecule mol, DFSVisitor visitor)

Void depth_first_search_mask (Molecule mol, DFSVisitor visitor, List<Bool> atomMask)

Void depth_first_search_mask(Molecule mol, DFSVisitor visitor, List<Bool> atomMask,

List<Bool> bondMask)

Void depth_first_search (Molecule mol, Atom atom, DFSVisitor visitor)

Void depth_first_search_mask(Molecule mol, Atom atom, DFSVisitor visitor, List<Bool> atomMask)

Void depth_first_search_mask (Molecule mol, Atom atom, DFSVisitor visitor, List<Bool> atomMask, List<Bool> bondMask)

Void ordered_depth_first_search(Molecule mol, List<Int> order, DFSVisitor visitor)

Void exhaustive depth_first_search(Molecule mol, Atom atom, DFSVisitor visitor)

BREADTH-FIRST SEARCH

Void breadth_first_search(Molecule mol, BFSVisitor visitor)

Void breadth_first_search_mask (Molecule mol, BFSVisitor visitor, List<Bool> atomMask)

Void breadth_first_search_mask(Molecule mol, BFSVisitor visitor, List<Bool> atomMask,

List<Bool> bondMask)

Void breadth first search (Molecule mol, Atom atom, BFS Visitor visitor)

Void breadth_first_search_mask (Molecule mol, Atom atom, BFSVisitor visitor, List<Bool> atomMask)

Void breadth_first_search_mask(Molecule mol, Atom atom, BFSVisitor visitor, List<Bool> atomMask,

List<Bool> bondMask)

COMPONENTS

List<Int> connected_bond_components(Molecule mol)

List<Int> connected_atom_components(Molecule mol)

Int num_connected_components(Molecule mol)

Algorithms (contd.)

CYCLES (RINGS)

Int cyclomatic number(Molecule mol)

Int cyclomatic_number(Molecule mol, Int numComponents)

(List<Int, List<Int>) cycle_membership(Molecule mol) [python]

Void cycle_membership(Molecule mol, List<Bool> cyclicAtoms, List<Bool> cyclicBonds) [C++]

RingSet relevant_cycles(Molecule mol)

RingSet relevant_cycles(Molecule mol, List<Bool> cyclicAtoms, List<Bool> cyclicBonds)

FLOYD-WARSHALL SHORTEST PATH

DistanceMatrix floyd warshall(Molecule mol)

MORGAN'S EXTENDED CONNECTIVITIES

List<Int> extended connectivities (Molecule mol, AtomInvariant invariant)

CANONICALIZATION

(List<Int>, List<Int>) canonicalize component(Molecule mol, List<Int> symmetry,

AtomInvariant atomInvariant, BondInvariant bondInvariant)

(List<Int>, List<Int>) canonicalize(Molecule mol, List<Int> symmetry,

AtomInvariant atomInvariant, BondInvariant bondInvariant,

List<Int> atomComponents, List<Int> bondComponents)

ENUMERATION

List<List<Int>> enumerate_paths(Molecule mol, Int maxSize)

Void enumerate_subgraphs(Molecule mol, SubgraphCallback callback, Int maxSize, Bool onlyTrees)

FINGERPRINTS

Fingerprint path_fingerprint(Molecule mol, Int maxSize = 7, Int numWords = 16, Int hashPrime = 1021)
Fingerprint tree_fingerprint(Molecule mol, Int maxSize = 7, Int numWords = 16, Int hashPrime = 1021)
Fingerprint subgraph_fingerprint(Molecule mol, Int maxSize = 7, Int numWords = 16,

Int hashPrime = 1021)

Int hashPrime = 1021)

2D COORDINATE GENERATION

List<(Real, Real)> generate_diagram(Molecule mol)

Depict

Enum Options: BlackWhiteAtoms, NoMargin,

DrawTermC, DrawAllC, NoWedgeHashGen,

AsymmetricDoubleBond, AromaticCircle,

AromaticHash

Depict(Painter painter)

Bool drawMolecule (Molecule mol, RingSet rings,

List<(Real, Real)> coordinates)

Void **setBondLength**(Real **length**)

Real bondLength()

Void setPenWidth(Real width)

Real penWidth()

Void **setBondSpacing**(Real **spacing**)

Real bondSpacing()

Void setBondWidth(Real width)

Real bondWidth()

Void setOptions(Int options)

Int options()

Void clearOptions()

Void **setFontFamily**(String **family**)

String fontFamily()

Void **setFontSize**(Real **size**)

Real fontSize()

Void setBondColor(String color)

Painter implementations: SVGPainter