### NAME

Bond

#### SYNOPSIS

use Bond;

use Bond qw(:all);

### **DESCRIPTION**

Bond class provides the following methods:

new, Copy, DeleteBond, GetAtoms, GetBondBeginAtom, GetBondEndAtom, GetBondFromAtom, GetBondToAtom, GetBondedAtom, GetCommonAtom, GetLargestRing, GetNumOfRings, GetNumOfRingsWithEvenSize, GetNumOfRingsWithOddSize, GetNumOfRingsWithSize, GetNumOfRingsWithSizeGreaterThan, GetNumOfRingsWithSizeLessThan, GetRingsWithEvenSize, GetRingsWithOddSize, GetRingsWithSize, GetRingsWithSizeGreaterThan, GetRingsWithSizeLessThan, GetSizeOfLargestRing, GetSizeOfSmallestRing, GetSmallestRing, IsAromatic, IsBondStereochemistrySpecified, IsBondTypeSpecified, IsCis, IsCisOrTrans, IsCoordinate, IsDative, IsDouble, IsDown, IsDownward, IsHash, IsInRing, IsInRingOfSize, IsIonic, IsNotInRing, IsOnlyInOneRing, IsQuadruple, IsQuintuple, IsSextuple, IsSingle, IsTautomeric, IsTrans, IsTriple, IsUp, IsUpOrDown, IsUpward, IsWedge, IsWedgeOrHash, SetAtoms, SetBondOrder, SetBondStereochemistry, SetBondType, StringifyBond, SwitchBondFromAndToAtoms

Bond class is derived from ObjectProperty base class which provides methods not explicitly defined in Atom or ObjectProperty class using Perl's AUTOLOAD functionality. These methods are generated on-the-fly for a specified object property:

```
Set<PropertyName>(<PropertyValue>);
$PropertyValue = Get<PropertyName>();
Delete<PropertyName>();
```

### **METHODS**

new

```
$NewBond = new Bond([%PropertyNameAndValues]);
```

Using specified *Bond* property names and values hash, new method creates a new object and returns a reference to newly created Bond object. By default, following properties are initialized:

```
ID = SequentialObjectID
@Atoms = ();
BondType = ""
BondOrder = ""
```

Except for *ID* property, all other default properties and other additional properties can be set during invocation of this method.

Examples:

Copy

```
$BondCopy = $Bond->Copy();
```

Copy Bond and its associated data using Storable::dclone and return a new Bond object.

DeleteBond

```
$Bond->DeleteBond();
```

Delete **Bond** between atoms in from a molecule.

GetAtoms

```
@BondedAtoms = $Bond->GetAtoms();
```

Returns an array containing Atoms invoved in Bond.

GetBondedAtom

```
$BondedAtom = $Bond->GetBondedAtom($Atom);
```

Returns BondedAtom bonded to Atom in Bond.

GetBondBeginAtom

```
$BeginAtom = $Bond->GetBondBeginAtom();
```

Returns BeginAtom corresponding to bond starting atom in Bond.

GetBondEndAtom

```
$EndAtom = $Bond->GetBondEndAtom();
```

Returns EndAtom corresponding to bond ending atom in Bond.

#### GetBondFromAtom

```
$FromAtom = $Bond->GetBondFromAtom();
```

Returns FromAtom corresponding to bond starting atom in Bond.

### GetBondToAtom

```
$ToAotm = $Bond->GetBondToAtom();
```

Returns ToAtom corresponding to bond ending atom in Bond.

#### GetCommonAtom

```
$CommonAtom = $Bond->GetCommonAtom($OtherBond);
```

Returns Atom common to both Bond and \$OtherBond.

### GetLargestRing

```
@RingAtoms = $Bond->GetLargestRing();
```

Returns an array of ring Atoms corresponding to the largest ring containing Bond. in a molecule

### GetNumOfRings

```
$NumOfRings = $Bond->GetNumOfRings();
```

Returns number of rings containing Bond in a molecule.

### GetNumOfRingsWithEvenSize

```
$NumOfRings = $Bond->GetNumOfRingsWithEvenSize();
```

Returns number of rings with even size containing Bond in a molecule.

## GetNumOfRingsWithOddSize

```
$NumOfRings = $Bond->GetNumOfRingsWithOddSize();
```

Returns number of rings with odd size containing *Bond* in a molecule.

### GetNumOfRingsWithSize

```
$NumOfRings = $Bond->GetNumOfRingsWithSize($RingSize);
```

Returns number of rings with specific RingSize containing Bond in a molecule.

# ${\tt GetNumOfRingsWithSizeGreaterThan}$

```
$NumOfRings = $Bond->GetNumOfRingsWithSizeGreaterThan($RingSize);
```

Returns number of rings with size greater than specific RingSize containing Bond in a molecule.

# ${\sf GetNumOfRingsWithSizeLessThan}$

```
$NumOfRings = $Bond->GetNumOfRingsWithSizeLessThan($RingSize);
```

Returns number of rings with size less than specific RingSize containing Bond in a molecule.

## GetRings

```
@Rings = $Bond->GetRings();
```

Returns an array of references to arrays containing ring atoms corressponding to all rings containing *Bond* in a molecule.

## GetRingsWithEvenSize

```
@Rings = $Bond->GetRingsWithEvenSize();
```

Returns an array of references to arrays containing ring atoms corressponding to all rings with even size containing *Bond* in a molecule.

### GetRingsWithOddSize

```
@Rings = $Bond->GetRingsWithOddSize();
```

Returns an array of references to arrays containing ring atoms corressponding to all rings with odd size containing *Bond* in a molecule.

### GetRingsWithSize

```
@Rings = $Bond->GetRingsWithSize($RingSize);
```

Returns an array of references to arrays containing ring atoms corressponding to all rings with specific *RingSize* containing *Bond* in a molecule.

# ${\tt GetRingsWithSizeGreaterThan}$

```
@Rings = $Bond->GetRingsWithSizeGreaterThan($RingSize);
```

Returns an array of references to arrays containing ring atoms corressponding to all rings with size greater than specific *RingSize* containing *Bond* in a molecule.

## ${\sf GetRingsWithSizeLessThan}$

```
@Rings = $Bond->GetRingsWithSizeLessThan($RingSize);
```

Returns an array of references to arrays containing ring atoms corressponding to all rings with size less than specific *RingSize* containing *Bond* in a molecule.

### GetSizeOfLargestRing

```
$Size = $Bond->GetSizeOfLargestRing();
```

Returns size of the largest ring containing Bond in a molecule.

### GetSizeOfSmallestRing

```
$Size = $Bond->GetSizeOfSmallestRing();
```

Returns size of the smallest ring containing Bond in a molecule.

### GetSmallestRing

```
@RingAtoms = $Bond->GetSmallestRing();
```

Returns an array of ring Atoms corresponding to the largest ring containing Bond in a molecule.

#### **IsAromatic**

```
$Status = $Bond->IsAromatic();
```

Returns 1 or 0 based on whether it's an aromatic Bond.

#### IsBondStereochemistrySpecified

```
$Status = $Bond->IsBondStereochemistrySpecified();
```

Returns 1 or 0 based on whether Bond's sterochemistry is specified.

## IsBondTypeSpecified

```
$Status = $Bond->IsBondTypeSpecified();
```

Returns 1 or 0 based on whether Bond's type is specified.

IsCis

```
$Status = $Bond->IsCis();
```

Returns 1 or 0 based on whether it's a cis Bond.

### IsCisOrTrans

```
$Status = $Bond->IsCisOrTrans();
```

Returns 1 or 0 based on whether it's a cis or trans Bond.

### IsCoordinate

```
$Status = $Bond->IsCoordinate();
```

Returns 1 or 0 based on whether it's a coordinate or dative Bond.

## IsDative

```
$Status = $Bond->IsDative();
```

Returns 1 or 0 based on whether it's a coordinate or dative Bond.

## IsDouble

```
$Status =$Bond->IsDouble();
```

Returns 1 or 0 based on whether it's a double Bond.

### IsDown

```
$Status = $Bond->IsDown();
```

Returns 1 or 0 based on whether it's a hash or down single Bond.

```
IsDownward
           $Return = $Bond->IsDownward();
       Returns 1 or 0 based on whether it's a downward Bond.
IsHash
           $Status = $Bond->IsHash();
       Returns 1 or 0 based on whether it's a hash or down single Bond.
IsInRing
           $Status = $Bond->IsInRing();
       Returns 1 or 0 based on whether Bond is present in a ring.
IsInRingOfSize
           $Status = $Bond->IsInRingOfSize($Size);
       Returns 1 or 0 based on whether Bond is present in a ring of specific Size.
Islonic
           $Status = $Bond->IsIonic();
       Returns 1 or 0 based on whether it's an ionic Bond.
IsNotInRing
           $Status = $Bond->IsNotInRing();
       Returns 1 or 0 based on whether Bond is not present in a ring.
IsOnlyInOneRing
           $Status = $Bond->IsOnlyInOneRing();
       Returns 1 or 0 based on whether Bond is only present in one ring.
IsQuadruple
           $Status = $Bond->IsQuadruple();
       Returns 1 or 0 based on whether it's a quadruple Bond.
IsQuintuple
           $Status = $Bond->IsQuintuple();
       Returns 1 or 0 based on whether it's a quintuple Bond.
IsSextuple
           $Status = $Bond->IsSextuple();
       Returns 1 or 0 based on whether it's a sextuple Bond.
IsSingle
           $Status =$Bond->IsSingle();
       Returns 1 or 0 based on whether it's a single Bond.
IsTriple
           $Status =$Bond->IsTriple();
       Returns 1 or 0 based on whether it's a triple Bond.
IsTautomeric
           $Status = $Bond->IsTautomeric();
       Returns 1 or 0 based on whether it's a Bond.
IsTrans
           $Status = $Bond->IsTrans();
       Returns 1 or 0 based on whether it's a trans Bond.
IsUp
           $Status = $Bond->IsUp();
       Returns 1 or 0 based on whether it's a up Bond.
```

### IsUpOrDown

```
$Status = $Bond->IsUpOrDown();
```

Returns 1 or 0 based on whether it's an up or down Bond.

#### **IsUpward**

```
$Status = $Bond->IsUpward();
```

Returns 1 or 0 based on whether it's an upward Bond.

#### IsWedge

```
$Status = $Bond->IsWedge();
```

Returns 1 or 0 based on whether it's a wedge Bond.

## IsWedgeOrHash

```
$Status = $Bond->IsWedgeOrHash();
```

Returns 1 or 0 based on whether it's a wedge or hash Bond.

#### SetAtoms

```
$Bond->SetAtoms($AtomsRef);
$Bond->SetAtoms(@Atoms);
```

Set atoms of Bond to atoms in Atoms array or in a reference to an array of atoms and return Bond.

#### SetBondOrder

```
$Bond->SetBondOrder($BondOrder);
```

Sets bond order of Bond to specified BondOrder and returns Bond. Possible bond order values: 1 = Single, 1.5 = Aromatic, 2 = Double, 3 = Triple, 4 = Quadruple, 5 = Quintuple, 6 = Sextuple, 7 = Septuple

#### Notes:

- . BondType property is automatically assigned using default BondType values for specified BondOrder.
- . BondType values can also be explicit set.
- . To make bonds aromatic in a ring, explicitly set "Aromatic" property for bond/atoms and make sure appropriate BondOrder values are assigned.
- . Dative or coordinate bond types are treated as single bond types with explicit formal charge of + and on first and second bond atoms.

### SetBondType

```
$Bond->SetBondType($BondType);
```

Sets bond type for *Bond* to specified *BondType* and returns *Bond*. Possible bond type values for different bond orders are:

```
0: None, Ionic, Unspecified
```

- 1 : Single, Dative, Coordinate, SingleOrDouble, SingleOrAromatic, Tautomeric
- 2 : Double, SingleOrDouble, DoubleOrAromatic, Tautomeric
- 3 : Triple
- 4 : Quadruple
- 5 : Quintuple
- 6 : Sextuple
- 7 : Septuple
- 1.5 : Aromatic, Resonance, SingleOrAromatic, DoubleOrAromatic

### Notes:

- o BondType Any is valid for all BondOrders.
- o BondOrder property is automatically assigned using default BondOrder values for specified BondType.

Possible bond stereochemistry values for different bond orders are:

```
0 : None, Unspecified
```

- 1 : Wedge, Up, Hash, Down, Wavy, WedgeOrHash, UpOrDown, Upward, Downward, None, Unspecified
- 2 : Cis, Trans, Z, E, DoubleCross, CisOrTrans, None, Unspecified

## SetBondStereochemistry

```
$Bond = $Bond->SetBondStereochemistry($BondStereochemistry);
```

Sets bond stereochemistry of *Bond* to specified *BondStereochemistry* and returns *Bond*. Possible *BondStereoChemistry* values for different bond orders are:

BondOrder: 1

```
None, Unspecified: Not a stereo bond or unspecified
    Wedge, Up : Wedge end pointing up
    Hash, Down: Wedge end pointing down
    Wavy, WedgeOrHash, UpOrDown: Wedge end up or down
    Upward: Single bond around cis/trans double bonds pointing upward
    Downward: Single bond around cis/trans double bonds pointing upward
Notes:
    o Wedge starts at begin atom of a bond making wedge pointed end always
      at this atom.
    o Upward/downward bonds start at atoms involved in cis/trans double bonds.
BondOrder: 2
   None, Unspecified: Not a stereo bond or unspecified
    Z, cis: Similar groups on same side of double bond
    E, trans: Similar groups on different side of double bond
    CisOrTrans, DoubleCross: cis or trans
    $BondString = $Bond->StringifyBond();
```

## StringifyBond

Returns a string containing information about *bond* object.

### SwitchBondFromAndToAtoms

\$Bond = \$Bond->SwitchBondFromAndToAtoms();

Swaps bond from and to atoms in Bond and returns Bond.

### **AUTHOR**

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### SEE ALSO

Atom.pm, Molecule.pm

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