

# SINDO 4.0 User's Manual: SINDO

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# 1 Sindo

## 1.1 &mol group

- Character(80) :: minfoFile  
The name of the .minfo file, in which the information of molecule is written.
- Integer :: Nat  
The number of atoms
- Real(8), dimension(Nat) :: Mass  
The mass of each atoms (in atomic mass unit)
- Real(8), dimension(3,Nat) :: x  
The reference (equilibrium) geometry (in Angstrom)
- Real(8), dimension(Nfree) :: omega  
The frequencies for the HO basis sets (in  $\text{cm}^{-1}$ )
- Real(8), dimension(Nat\*3,Nfree) :: L  
The vibrational displacement vectors

\* Note that 'minfoFile' is mutually exclusive from others.

## 1.2 &sys group

- Integer(8) :: Maxmem  
Maximum size of memory (MB)

## 1.3 &mrpes group

- Integer :: MR  
Mode representation (MR=1-4)
- Real(8) :: mcs\_cutoff  
Cutoff of QFF based on MCS in  $\text{cm}^{-1}$  (default = 1.d-04)
- Logical :: au  
The grid data in atomic unit (default = true)
- Character(80) :: mopFile  
The name of the mop file.

## 1.4 &vib group

- Integer :: Nfree  
Number of degrees of freedom (default = 3Nat - 6)
- Integer :: MR  
Mode representation (MR=1-4)

- Integer, dimension(Nfree) :: vmax  
Number of basis functions for each mode (default=10)
- Integer :: vmaxALL  
Number of basis functions for all modes (default=10)
- Integer :: vmax\_base  
same as vmaxALL
- Logical :: vscf, ocvscf, vci, vpt, vqdpt  
invoke vscf/ocvscf/vci/vpt/vqdpt
- Logical :: prpt  
invoke property calculation
- Logical :: readBasis  
read the basis functions from cho.basis

## 1.5 &states group

- Integer :: Nstate  
Number of states to calculate
- Integer, dimension(Nfree,Nstate) :: target\_state  
Labels of the target states
- Logical :: fund  
Compute fundamentals

## 1.6 &vscf group

- Logical :: state\_specific  
State specific VSCF if true (default = .false.)
- Logical :: restart  
Restart from vscf\_xxx.wfn (default = .false.)
- Integer :: Maxitr  
Maximum number of iteration (default = 10)
- Real(8) :: Ethresh  
Threshold of convergence (default = 1e-03 cm<sup>-1</sup>)

## 1.7 &ocvscf group

- Integer :: maxOptIter  
Maximum number of iteration (default = 30)
- Real(8) :: ethresh  
Threshold of the energy (default = 1e-06 cm<sup>-1</sup>)

- Real(8) :: gthresh  
Threshold of the gradient (default =  $1\text{e-}06 \text{ cm}^{-1} \text{ rad}^{-1}$ )
- Integer :: pfit  
Order of the Fourier fitting (default = 2)
- Character(80) :: mopFile  
The name of the mopfile
- Character(80) :: u1File  
The name of the file to write the transformation matrix (default = u1.dat)
- Integer :: icff  
Switch on CFF when icff = 1 and QFF when icff = 0 (default = 0)
- Integer :: iscreen  
Switch off/on pair selection when iscreen=0/1 (default = 1)
- Real(8) :: eta12thresh  
Threshold value for the pair screening (default =  $500 \text{ cm}^{-1}$ )

## 1.8 &vci group

- Integer :: Nstate  
Number of states to calculate
- Integer :: nCI  
Max CI dimension (cutoff based on the energy)
- Integer(Nfree) :: maxEx  
Max quantum number to excite for each mode
- Integer :: maxExALL  
Max quantum number to excite for all the modes
- Integer :: maxSum  
Max sum of quantum number
- Integer :: nCUP  
Max number of modes to excite
- Logical :: geomAv  
If true, calculate vibrationally averaged geometry
- Logical :: dump  
If true, dump the vci wavefunction to vci-w.wfn
- Real(8) :: printWeight  
Print the configuration with the weight larger than this threshold
- Logical :: readCIbasis  
If true, read CI basis from vci-w.wfn

- Logical :: dumpHmat  
If true, write the VCI hamiltonian matrix
- Logical :: noDiag  
If true, the diagonalization is skipped

## 1.9 &vpt group

- Integer :: maxSum  
Max sum of quantum number to excite (default = -1)
- Integer :: maxEx  
Max quantum number to excite (default = -1)
- Integer :: nCUP  
Max number of modes to excite (default = MR)
- Real(8) :: thresh\_ene  
Threshold energy to avoid divergence (default=1e-04 Hartree)
- Logical :: dump  
Dump the information to vmp-w.wfn

## 1.10 &vqdpt group

- Integer :: nGen  
The generation of P space (default=3)
- Real(8) :: thresh\_p0  
E0 pruning (default=500 cm<sup>-1</sup>)
- Real(8) :: thresh\_p1  
VPT based pruning (default=0.1)
- Real(8) :: thresh\_p2  
VCI pruning (default=0.05)
- Real(8) :: thresh\_p3  
VCI pruning (default=0.9)
- Integer :: pset  
Combine the p-space generated from several target states  
=0 when the target states have an overlap (default)  
=1 when the p-space components have an overlap
- Integer :: maxSum  
Max sum of quantum number to excite (default = -1)
- Integer :: nCUP  
Max number of modes to excite (default = MR)

- Integer :: pqSum  
P/Q interaction scheme  
> 0 prune the interaction when  $\lambda_{pq} > \text{maxSum}$  (default)  
< 0 full interaction
- Integer :: vqdpt2\_loop  
=0 loop over q, then p, p' (default)  
=1 loop over p, then p', q
- Real(8) :: thresh\_ene  
Threshold energy to avoid divergence (default=1e-04 Hartree)
- Real(8) :: printWeight  
Print the configuration with the weight larger than this threshold (default=0.001)
- Logical :: dump  
Dump the information to vqdpt-w.wfn (default=true)

### 1.11 &prpt group

- Logical :: vscfprpt, vciprpt, vptprpt, vqdptprpt  
Invoke property calculation for vscf, vci, vpt, vqpd wavefunction
- Integer :: MR  
Mode representation (default = 3)
- Character :: extn(\*)  
The extension of the property files
- Integer :: matrix(\*)  
= 0 calculate only the average  
> 0 calculate the matrix
- Logical :: infrared  
If true, calculate the IR intensity.

### 1.12 &prptvci group

- Integer :: Nstate  
The number of states

### 1.13 &IRspectrum group

- Real(8) :: minOmega, maxOmega  
Min/Max value of the spectrum (default = 100 - 4000  $\text{cm}^{-1}$ )
- Real(8) :: delOmega  
Interval of the data (default = 1  $\text{cm}^{-1}$ )



- Real(8) :: fwhm  
Full-width half maximum of the Lorentz function for convolutions (default =  $20 \text{ cm}^{-1}$ )
- Real(8) :: cutoff  
Cutoff of the band (default =  $-1 \text{ km mol}^{-1}$ )

## **2 Interface with Quantum Chemistry Programs**

**2.1 Gaussian**

**2.2 Molpro**

**2.3 ACESII**

**2.4 QChem**

### 3 Files

# References