

SimulationType MolecularDynamics
NumberOfCycles 30000
NumberOfEquilibrationCycles 5000
NumberOfInitializationCycles 5000
PrintEvery 1000
RestartFile no

Ensemble NVT

ChargeMethod Ewald
CutOff 12.0
TimeStep 0.0005
Forcefield Dubbeldam2007FlexibleIRMOF-1
EwaldPrecision 1e-6

Framework 0
FrameworkName IRMOF-1
UnitCells 3 3 3
ExternalTemperature 298.0
ComputeMSD yes
PrintMSDEvery 10

FrameworkDefinitions Dubbeldam2007FlexibleIRMOF-1

Component 0 MoleculeName argon
MoleculeDefinition TraPPE
IdealGasRosenbluthWeight 1.0
TranslationProbability 1.0
RotationProbability 1.0
ReinsertionProbability 1.0
CreateNumberOfMolecules 16