

SimulationType	MolecularDynamics
NumberOfCycles	10000
NumberOfEquilibrationCycles	5000
NumberOfInitializationCycles	5000
PrintEvery	100
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 298.15
 ComputeMSD yes
 PrintMSDEvery 10

FrameworkDefinitions Dubbeldam2007FlexibleIRMOF-1

Component 0	MoleculeName	argon
	MoleculeDefinition	TraPPE
	MolFraction	0.10
	IdealGasRosenbluthWeight	1.0
	TranslationProbability	1.0
	RegrowProbability	0.5
	IdentityChangeProbability	1.0
	NumberOfIdentityChanges	2
	IdentityChangeList	0 1
	RotationProbability	1.0
	ReinsertionProbability	1.0
	CreateNumberOfMolecules	16

Component 1	MoleculeName	methane
	MoleculeDefinition	TraPPE
	MolFraction	0.90
	IdealGasRosenbluthWeight	1.0
	TranslationProbability	1.0
	RegrowProbability	0.5
	IdentityChangeProbability	1.0
	NumberOfIdentityChanges	2
	IdentityChangeList	0 1
	RotationProbability	1.0
	ReinsertionProbability	1.0
	CreateNumberOfMolecules	16