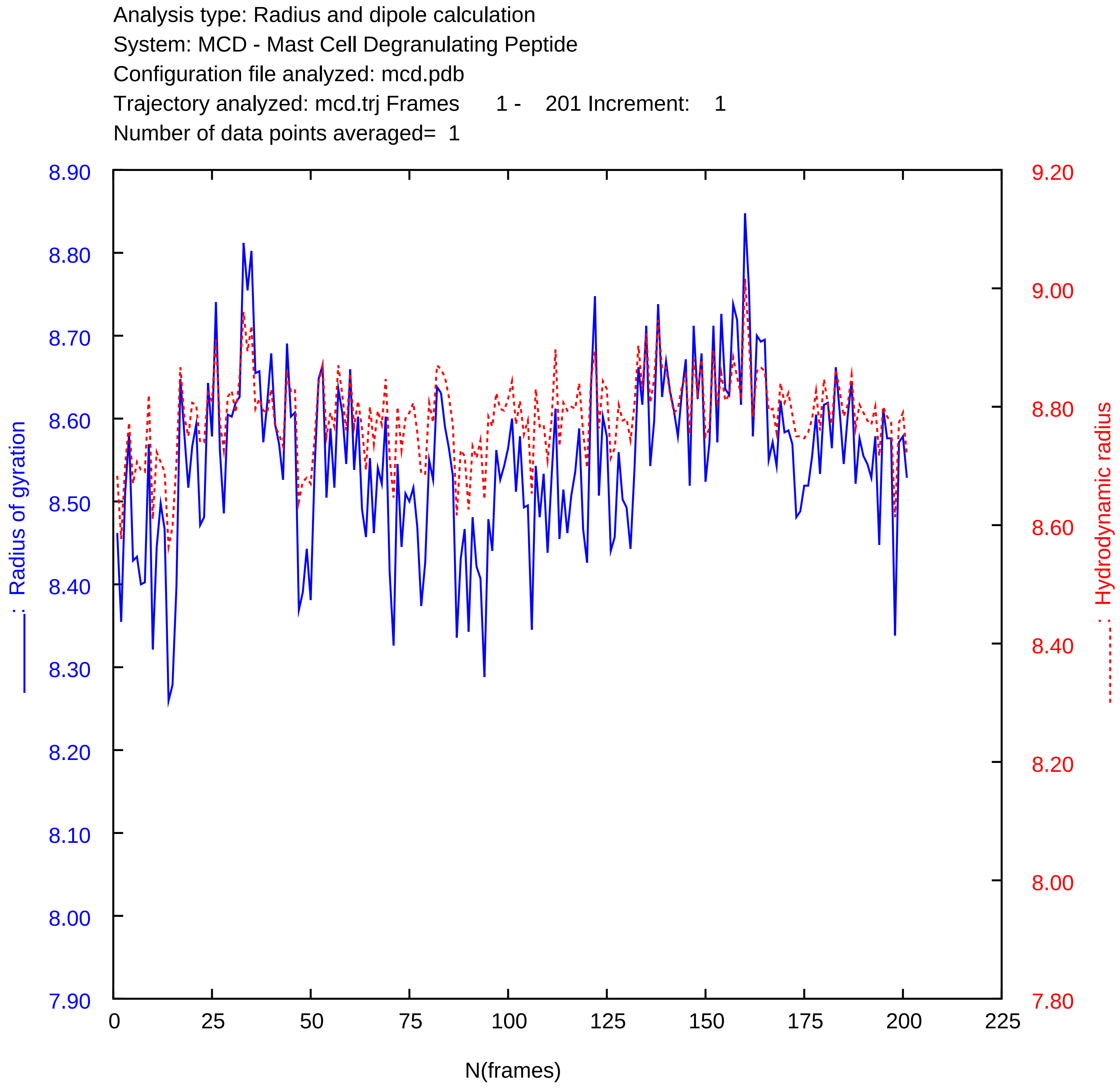
Analysis type: Radius and dipole calculation
System: MCD - Mast Cell Degranulating Peptide
Configuration file analyzed: mcd.pdb
Trajectory analyzed: mcd.trj Frames 1 - 201 Increment: 1
Number of data points averaged= 1
8.90
8.80
8.70
8.60
8.50
8.40
8.30
8.20
8.10
8.00
7.90
9.20
9.00
8.80
8.60
8.40
8.20
8.00
7.80
0
25
50
75
100
125
150
175
200
225
N(frames)
Radius of gyration
Hydrodynamic radius

Analysis type: Radius and dipole calculation
System: MCD - Mast Cell Degranulating Peptide
Configuration file analyzed: mcd.pdb
Trajectory analyzed: mcd.trj Frames 1 - 201 Increment: 1
Number of data points averaged= 1