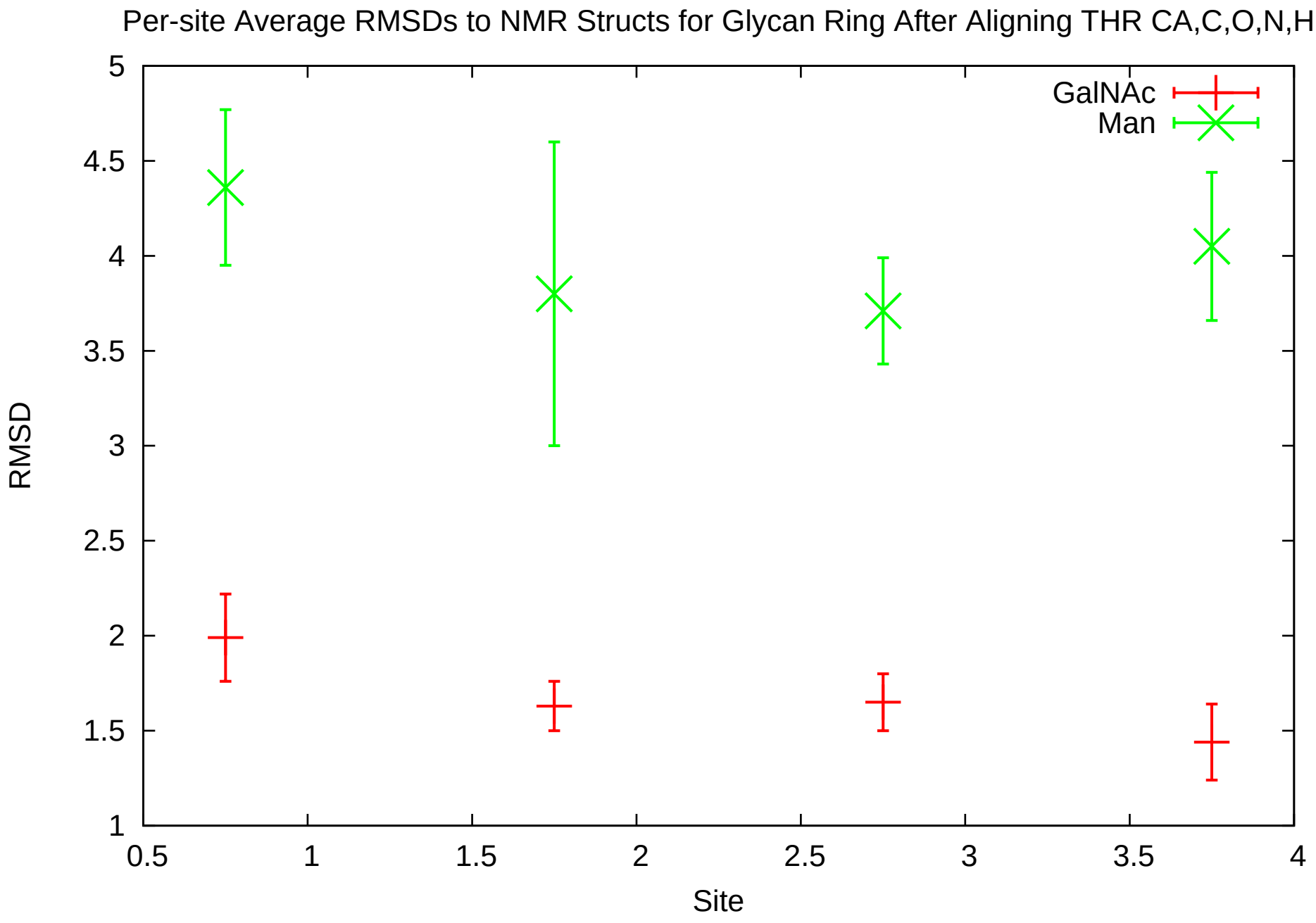
Per-site Average RMSDs to NMR Structs for Glycan Ring After Aligning THR CA,C,O,N,H
GalNAc
Man
RMSD
Site
5
4.5
4
3.5
3
2.5
2
1.5
1
0.5
1
1.5
2
2.5
3
3.5
4