


Within region correlation (after pruning & before recoding), SNPs ordered by LDbin

chr16.46925179.C.T
chr16.46901935.G.A
chr16.46920478.G.A
chr16.46889594.G.A
chr16.46928310.A.C
chr16.46907494.T.A
chr16.46871111.G.C
chr16.46869350.T.C
chr16.46886416.A.T
chr16.46858021.T.G
chr16.46880510.A.G
chr16.46855220.G.A
chr16.46855442.C.A
chr16.46942582.C.T
chr16.46934337.A.T
chr16.46918119.T.C
chr16.46908235.G.T
chr16.46905220.C.T
chr16.46896754.G.A
chr16.46884996.T.C
chr16.46882815.A.T
chr16.46864282.A.C
chr16.46859371.C.A
chr16.46876607.A.G

Correlation



-1.0 -0.5 0.0 0.5 1.0

