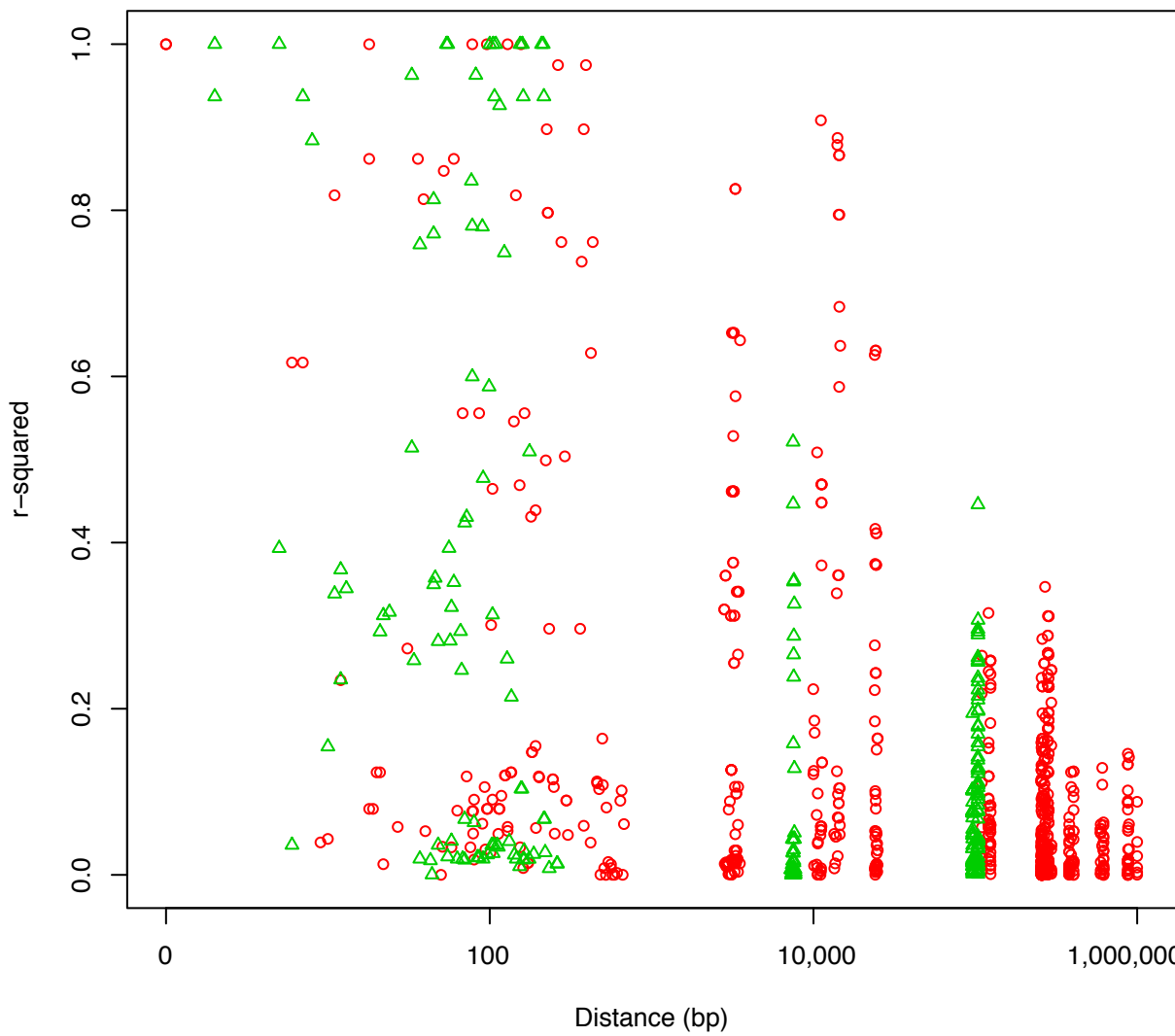


a) All sites



b) Excluding sites associated with phenotype

