

Single marker analysis

- The patients are classified as WORST, MIDDLE, LEAST.
- WORST and MIDDLE are also combined as SICK to compare with LEAST.
- Use chi-square test to test an association between a SNP marker and CFS for allele frequencies in case (W, M, S) and control (L) group.
- Raw p-values are corrected by BH.
- No association is observed for case and control group.