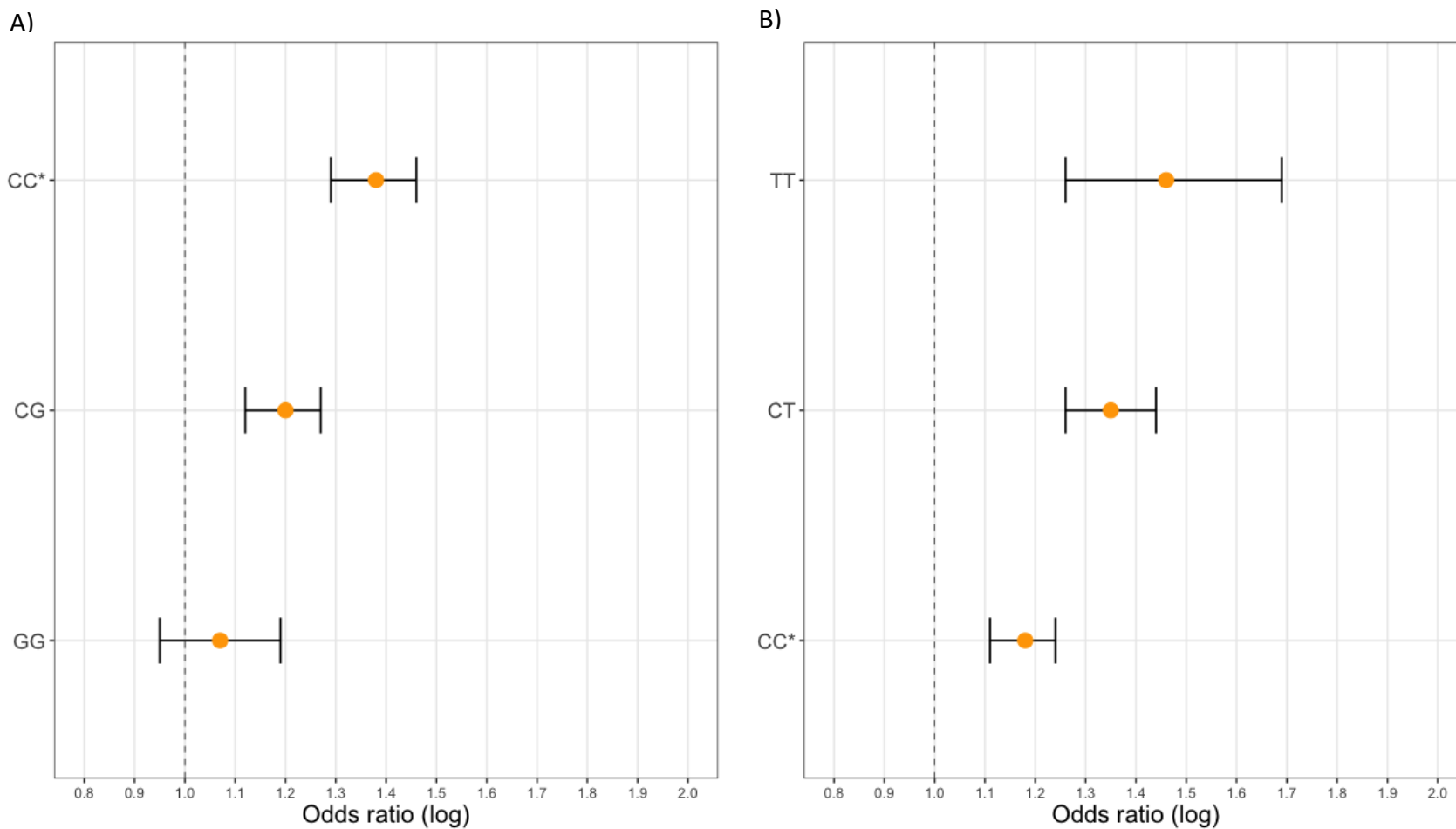


Supplemental Figure 3. Association of red meat intake and CRC risk stratified by variant genotypes.



Genotypes were defined by main results from genome-wide analysis (Table 3): A) rs4871179 (Chr 8, downstream *HAS2*), B) rs35352860 (Chr 18, intron *SMAD7*).
*Denotes reference allele.