

Figure S4: Overview of the SROs and probability profiling of genomic regions linked to selected traits



Legend: : The probability profile was calculated taking into account both the maximum size of the deletions determined by array-CGH analysis

Figure S5: Overview of the SROs and probability profiling of genomic regions linked to selected traits



Legend : The probability profile was calculated taking into account both the minimum size of the deletions determined by array-CGH analysis