



Figure 6 – Results of biological annotation of genomic intervals of the second subset

Note: a – association of genes within intervals of the second subset with diseases according to <https://diseases.jensenlab.org/portal> (terms of gene-associated diseases were selected when z-score > 5); b – associations of genes within intervals with predicted age-associated and tissue-specific expression changes according to VoyAger web tool (adjusted P-value ≤ 0.05)