



Figure 5 – Results of biological annotation of genomic intervals of the first subset

*Note: a – association of genes within intervals of the first 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)*