

Figure S3: Manhattan and Quantile-Quantile Plot of GWAS analysis of individuals affected up to two years of age. Identified variants could have an effect on the time and/or severity of the affection. One variant can be considered significant (chr21, BICF2G630640798, raw P-value = 5.01×10^{-7}), it reaches the significance threshold (Bonferroni correction = 5.8×10^{-7} ; shown as a purple line), and it lies in a gene ANO3. Lambda value shows that stratification correction worked well.

