“Characterisation of genetic variation in *ST8SIA2* and its interaction region in NCAM1 in patients with bipolar disorder”

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**Figure S1: Comparison of genotype quality score and minor allele frequency to identify potentially erroneous SNPs.** All points represent raw SNPs that were novel (not present in dbSNP 135). Allele frequency (AF) vs. Quality by Depth score (QD). Hollow diamonds – SNPs that failed filtering according to GATK recommended filtering, Black circles – SNPs that passed filtering. Points in dashed oval have low QD and high AF, suggesting they are not bona-fide SNPs.