



SUPPLEMENTARY FIG. S5. DMF did not modify the expression of proteasome genes. qRT-PCR determination of mRNA levels of proteasome subunits coding *PsmB7*, *PsmC3*, and *PsmC4*, normalized by β -actin. (A), dopaminergic MN9D. (B), microglial BV2 and (C), astrocytic IMA2.1 cell lines were treated with DMF ($20 \mu M$) for 4, 8, and 24 h. Differences among groups were assessed by two-way ANOVA followed by Bonferroni's test. Asterisks denote significant differences with $**p < 0.01$ comparing the basal group.