

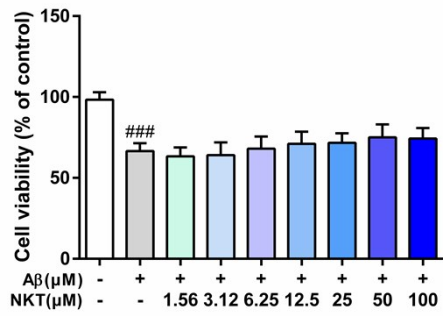
Supplementary Data

Table

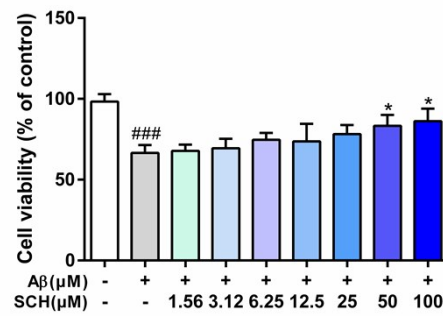
Table S1 Primer sequences used for qPCR analysis.

Gene	Forward	Reverse
PI3K	5'-ATGTGTATGGACCCGGAAGG-3'	5'-AGCCATCTGCCTCCACGTTAG-3'
AKT	5'-ACTCATTCCAGACCCACGAC-3'	5'-CCGGTACACCACGTTCTTCT-3'
Caspase3	5'-TGA CTGGAAAGCCGAAACT-3'	5'-GGGTGCGGTAGAGTAAGCAT-3'
p62	5'-ATCAGCTTCTGGTCCATCGG-3'	5'-GCTTCTTTTCCCTGTGCT-3'
Atg5	5'-CCCAGTATCCCTCACTTAC-3'	5'-GCTAGGGTGAGTCCATTAT-3'
Beclin1	5'-ATCCTGGACCGTGTCACCATCCAGG- 3'	5'-GTTGAGCTGAGTGTCCAGCTGG-3'
β -actin	5'-CACCCGCGAGTACAACCTTC-3'	5'-CCCATACCCACCATCACACC-3'

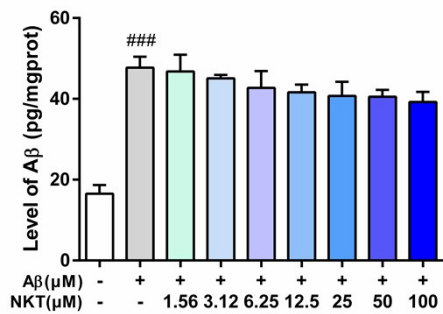
Figures



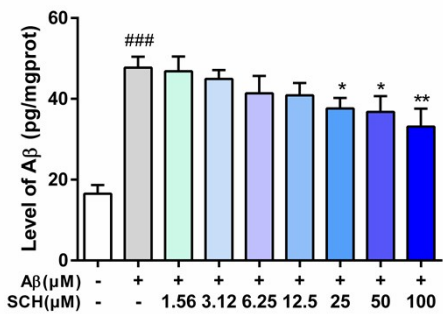
A



B



C



D

Fig. S1 PC12 cells were pretreated with different concentrations of NKT/ SCH for 4 h, and then incubated with 20μM Aβ for 20 h. The cell viability (A and B) was determined using MTT assay and the inhibition of Aβ level was determined using ELISA assay (C and D). The values represent the mean ± SD (n = 3 in each group), *p < 0.05, **p < 0.01, ***p < 0.001 versus the Aβ group; ###p < 0.001 versus the control group.