Validation of astrocytic cAMP signalling to study therapeutic targets for Alzheimer's disease



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4. cAMP activation in rodent primary astrocytes

reduces the expression of inflammatory genes

1. Introduction

Astrocytes play a fundamental role in pathological processes associated with neurodegenerative diseases including neuroinflammation, impaired glutamate uptake, reduced neurotrophic support and defective metabolism. Activation of cAMP signalling in astrocytes elevates glycolytic rate, increases glutamate transporter and neurotrophic factor expression, and suppresses the immune response. Molecules that modulate astrocytic cAMP signalling are therefore potential therapeutic targets for neurodegenerative diseases such as Alzheimer's disease (AD). To support the identification and validation of astrocyte-centric targets, we have optimised a suite of in vitro assays including cAMP, Ca2+, and metabolic assays, multielectrode array (MEA) and RNA-seg in rodent, human foetal and hiPSC-derived control and familial AD (fAD) astrocyte models. We have tested the adenylyl cyclase activator forskolin and a GPCR agonist in these assays to validate and increase our confidence in astrocytic cAMP signalling as an AD drug target.



GPCR RNA expression in control and AD hippocampus. A-B) RNAscope analysis demonstrating GPCR co localisation with EAAT2 (astrocytic marker; (A)) and MAP2 (neuronal marker; (B)) in human control hippocampus. C) qPCR analysis of GPCR RNA expression relative to TBP and UBE2D2 in human hippocampus, n=7-17 biological replicates. qPCR data shown as mean \pm SEM and analysed using one-way ANOVA, ns= non-significant.

3. Forskolin and a GPCR agonist induce cAMP activation in astrocytes in vitro





8 or 24h. (A) PCA analysis, (B) Gene Ontology (GO) analysis for biological processes carried out on the 200 significantly up-regulated genes (UP) and the 200 significantly down-regulated genes (DOWN), with the greatest log, fold change at the 4h and 8h timepoints. Pathways below dashed line have a false-discovery rate (FDR)>0.05. (C) Volcano plots of significantly upregulated genes (red; FDR<0.05) and downregulated genes (blue; FDR<0.05) at 4h and 8h post-GPCR agonist treatment, (D) FPKM values of individual genes, N=3 biological replicates per condition



Glycolytic rate was measured in rat primary astrocytes (A and B), 2-week matured hiPSC-derived control and fAD astrocytes (C), human foetal astrocytes (D) using the Seahorse XFe96 analyser (Agilent) and olycolytic rate assay kit, Representative mean ± SEM traces, 3-10 technical replicates per condition, Data normalised to cell number (Hoechst stain; A, B, D), and number of live cells (Hoescht and PI stain; C). Vehicle = H₂O (B and C), 0.1% DMSO (A) and 0.2% DMSO (D). IBMX = 3-lsobutyl-1-methylxanthine

6. A GPCR agonist enhances neuronal activity in rodent co-cultures of astrocytes and neurons



Neuronal activity in rat primary co-cultures of astrocytes and neurons treated with a GPCR agonist. (A) Agonist or vehicle was administered at DIV14 at 0.15h; mean firing rate was recorded for 15 minutes at the indicated timepoints using the Maestro Pro multielectrode array (MEA) system (Axion Biosystems), n=6 technical replicates. (B) Percentage change in firing rate vs baseline at 2h post-agonist or vehicle treatment, n=4 biological replicates. (C) Raster plot of neuronal activity in a single well 2 hours post-agonist or vehicle treatment. Vehicle = cell culture media.

7. Summary

We have demonstrated that RNA levels of a G_s -coupled GPCR are not altered in the AD brain Both forskolin and the GPCR agonist elevate glycolytic rate in rat and human foetal astrocytes, suggesting that the mechanism is cAMP-mediated. Interestingly, the GPCR agonist enhances neuronal firing in vitro in neuron-astrocyte co-cultures. Finally, RNA-seq analysis identified cAMP pathway and inflammatory genes as major targets of both forskolin and GPCR agonist administration. Overall, our data suggest that activation of astrocytic cAMP signalling has exciting therapeutic potential as a treatment for AD



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Data



