Hemisphere Asymmetry of Response to Pharmacologic Treatment in an Alzheimer’s Disease Mouse Model

Antigoni Manousopouloua,b, Satoshi Saitoc, Yumi Yamamotoe, Nasser M. Al-Daghrid,d,e, Masafumi Ichra, Roxana O. Carareb,1 and Spiros D. Garbisab,g,1,*

aInstitute for Life Sciences, University of Southampton, UK
bClinical and Experimental Sciences, Faculty of Medicine, University of Southampton, UK
cDepartment of Regenerative Medicine and Tissue Engineering, National Cerebral and Cardiovascular Center, Japan
dBiomarkers Research Program, Department of Biochemistry College of Science, King Saud University, KSA
ePrince Mutaib Chair for Biomarkers of Osteoporosis, Department of Biochemistry King Saud University, KSA
fDepartment of Neurology, National Cerebral and Cardiovascular Center, Japan
gCancer Sciences, Faculty of Medicine, University of Southampton, UK

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Abstract. The aim of this study was to examine hemisphere asymmetry of response to pharmacologic treatment in an Alzheimer’s disease mouse model using cilostazol as a chemical stimulus. Eight-month-old mice were assigned to vehicle or cilostazol treatment for three months and hemispheres were analyzed using quantitative proteomics. Bioinformatics interpretation showed that following treatment, aggregation of blood platelets significantly decreased in the right hemisphere whereas neurodegeneration significantly decreased and synaptic transmission increased in the left hemisphere only. Our study provides novel evidence on cerebral laterality of pharmacologic activity, with important implications in deciphering regional pharmacodynamic effects of existing drugs thus uncovering novel hemisphere-specific therapeutic targets.

Keywords: Alzheimer’s disease, cilostazol, hemisphere asymmetry, mass spectrometry pharmacology, pharmacodynamics, pharmacoproteomics

INTRODUCTION

Structural asymmetry in the human brain was first described by Broca in the 19th century [1]. Brain asymmetries have also been described in other mammals, including mice [2]. The asymmetry between cerebral hemispheres encompasses functional and neurochemical aspects [3]. Data from gene expression studies in humans demonstrated lateralization of gene transcription involved in synaptic transmission and neuronal electrophysiology [4].

Cerebral laterality has been implicated in the pathophysiology of Alzheimer’s disease [5]. However, whether therapeutic agents used for Alzheimer’s disease exert hemisphere-specific pharmacologic effects has yet to be elucidated.

Pharmacoproteomics, as a subdiscipline of proteomics, examines how a pharmacologic agent
perturbs a system of proteins that map to canonical or novel biological pathways. Such a discourse provides a more unbiased and holistic approach to understanding pharmacologic on-target or off-target effects at the protein level [6–9].

We have shown that treatment with cilostazol, a cyclic nucleotide phosphodiesterase III inhibitor, prevented cognitive decline as a result of amyloid-β (Aβ) deposition in the brain of Tg-SwDI mice [10]. Our hypothesis was that pharmacologic treatment of Alzheimer’s disease exerts hemisphere asymmetry.

MATERIALS AND METHODS

Transgenic mouse model

To ensure reproducibility of the pharmacologic treatment and subsequent proteomic analysis the experiment was performed in duplicate. For experiments A and B, male, eight-month-old, homozygous Tg-SwDI mice (n = 6 per experiment, n = 12 in total) on a pure C57BL/6 background were obtained from Jackson Laboratories. These mice express low levels of human Swedish/Dutch/Iowa mutant AβPP in neurons under the control of the mouse Thy1.2 promoter. Mice were randomly assigned to two groups and fed with pelleted chow containing 0.3% cilostazol (n = 3 per experiment, n = 6 in total) or standard pelleted chow only (n = 3 per experiment, n = 6 in total) for 3 months. Cilostazol was supplied by Otsuka Pharmaceutical Co. Ltd.

A previous study reported that in rats treated with 0.1% cilostazol, the plasma cilostazol concentration was approximately 1 μM [11]. Based on the in vitro IC50 value, this concentration is estimated to sufficiently inhibit phosphodiesterase III activity [12]. Additionally, studies have shown that treatment with 0.1-0.3% cilostazol in rats positively affected vascular function [12, 13]. Based on this evidence, a 0.3% cilostazol treatment was used in the present study.

Mice were perfused intracardially with phosphate buffer saline, cerebral hemispheres removed and snap frozen in liquid nitrogen. All experimental procedures were approved by the Institutional Animal Care and Use Committee at the National Cerebral and Cardiovascular Center, Japan.

Proteomic analysis

For experiments A and B, respectively, cerebral hemispheres were suspended in dissolution buffer (0.5 M triethylammonium bicarbonate, 0.05% sodium dodecyl sulfate) and homogenized using the FastPrep system (Savant Bio, Cedex, Fr) followed by pulsed probe sonication (Misonix, Farmingdale, NY, USA). Lysates were subjected to centrifugation (16,000 g, 10 min, 4°C) and supernatants measured for protein content using the Direct Detect™ system (Merck Millipore, Darmstadt, Germany). For experiments A and B separately, protein extracts were pooled from each hemisphere of the three vehicle treated mice (33.3 μg from each lysate giving 100 μg final protein content). Left and right hemispheres of cilostazol treated mice were individually analyzed (100 μg from each lysate). All samples were subjected to reduction, alkylation, trypsin proteolysis, eight-plex iTRAQ labeling and two-dimensional liquid chromatography, tandem mass spectrometry analysis as described previously [8, 14, 15] (Fig. 1A).

Unprocessed raw files were submitted to Proteome Discoverer 1.4 for target decoy searching with SequestHT [8, 14]. Quantification ratios were median-normalized and log2 transformed. After treatment, a protein was considered differentially expressed in the left compared to the right hemisphere relative to its respective control when its two-group T-Test p-value across both experiments was equal to or below 0.05.

Principal component analysis (using the iTRAQ ratios of all analyzed proteins for experiments A and B) was performed using BioConductor-R (version 2.15.1) and g-plots in R (version 3.1.2). Heatmap construction of differentially expressed proteins between the two hemispheres was generated using Gene Cluster (version 3.0) and Java Treeview (version 1.6r4). Ingenuity Pathway Analysis (Qiagen, Venlo, Netherlands) was applied to identify canonical pathways and biological processes enriched in the differentially expressed proteins between right and left cerebral hemispheres. An activation z-score ≥ 2.0 and a p-value ≤ 0.05 were considered significant.

RESULTS

A total of 9,116 proteins were quantitatively analyzed in experiment A (p < 0.05) and another 7,418 proteins in experiment B (p < 0.05), of which 6,137 were reproducibly profiled in both experiments. The average coefficient variation (CV) for the iTRAQ ratios of all proteins profiled across biological replicates was determined to be 24% and 23% for the right and left hemispheres, respectively, of cilostazol treated mice compared to control in experiment A.
Fig. 1. A) Experimental workflow. B) Principal component analysis for each experiment showed separate clustering for the right and left hemispheres of cilostazol treated mice compared to controls suggesting hemisphere asymmetry of pharmacologic response to cilostazol treatment. C) Hierarchical clustering analysis of differentially expressed proteins between the right and left hemispheres in both experiments visualized in heatmap format. D) Canonical pathway analysis of differentially expressed proteins between right and left hemisphere following cilostazol treatment showed that the G beta-gamma (G<sub>Bgamma</sub>) signaling pathway significantly decreased in the right hemisphere of cilostazol treated mice (p = 1.16E-2; activation z-score = -2.646) whereas it significantly increased in the left hemisphere of cilostazol treated mice (p = 1.16E-2; activation z-score = 2.646).

and 15% and 16% for the right and left hemispheres, respectively, of cilostazol treated mice compared to control in experiment B. Analogous CV values between biological replicates were reported by the authors using similar proteomic methodologies [14, 15]. Principal component analysis of the reporter ion ratios of proteins profiled in each experiment showed that the right hemispheres clustered separately from the left hemispheres of cilostazol treated mice (Fig. 1B). Following treatment with cilostazol, 765 proteins were differentially expressed between the right and left cerebral hemispheres across both proteomic experiments (p ≤ 0.05) (Fig. 1C) (Supplementary Table 1).

**Bioinformatics interpretation of proteomic results**

Ingenuity Pathway Analysis (IPA) using the average reporter ion ratios across both experiments of differentially expressed proteins between right and left hemisphere following cilostazol treatment showed that the G beta-gamma (G<sub>Bgamma</sub>) signaling pathway significantly decreased in the right hemisphere of cilostazol treated mice (p = 1.16E-2; activation z-score = -2.646) whereas it significantly increased in the left hemisphere of cilostazol treated mice (p = 1.16E-2; activation z-score = 2.646) (Fig. 1D).

IPA also showed that aggregation of blood platelets significantly decreased in the right hemisphere only of cilostazol treated mice (p = 3.98E-3; activation z-score = -2.231) (Fig. 2A) whereas neurodegeneration significantly decreased (p = 7.7E-6; activation z-score = -2.071) (Fig. 2B) and synaptic transmission significantly increased (p = 8.62E-6; activation z-score = 2.028) (Fig. 2C) specifically in the left hemisphere of cilostazol treated mice.

**DISCUSSION**

This study provided novel protein level evidence on the hemisphere-specific pharmacologic effects of cilostazol, a drug that has been reported to ameliorate the symptoms of Alzheimer’s disease [16]. This observation may have important implications for an ongoing clinical trial assessing the efficacy of cilostazol in mild cognitive impairment (COMCID
Fig. 2. Bioinformatics interpretation of the results showed that (A) aggregation of blood platelets significantly decreased in the right hemisphere ($p = 3.98 \times 10^{-3}$; activation z-score $= -2.231$) whereas (B) neurodegeneration significantly decreased ($p = 7.7 \times 10^{-6}$; activation z-score $= -2.071$) and (C) synaptic transmission significantly increased ($p = 8.62 \times 10^{-6}$; activation z-score $= 2.028$) in the left hemisphere of cilostazol treated mice.

trial, ClinicalTrials.gov identifier: NCT02491268) but may also be extended to other pharmacologic agents.

There are only few studies examining cerebral laterality in the pathophysiology of Alzheimer’s disease. Douaud et al. [17] suggested that neurodegenerative pathology may develop asymmetrically in humans, with left hemisphere atrophy being an important predictor of progression from mild cognitive impairment to Alzheimer’s disease. Singh et al. [5] have found that atrophy of the cerebral cortex occurred earlier and progressed faster in the left compared to the right hemisphere in patients with Alzheimer’s disease. Their findings confirmed previous observations by Chetelat et al. [18] on left lateralization of temporal changes occurring during mild cognitive impairment to Alzheimer’s disease transition. Thompson et al. [19] examined the dynamics of grey matter loss in Alzheimer’s disease and concluded that shifting deficits were asymmetric and predominantly occurring in the left compared to the right hemisphere.

Our results showed that cilostazol asymmetrically affected the two hemispheres of a transgenic Alzheimer’s disease mouse model. In the right hemisphere, aggregation of blood platelets was found to significantly decrease following cilostazol treatment (Fig. 2A). Cerebral amyloid angiopathy (CAA), characterized by the deposition of amyloid-beta in the walls of cerebral arteries plays an important role in the pathogenesis of Alzheimer’s disease. Recent evidence suggests that platelet activation may also contribute to the development of CAA [20, 21]. Targeting blood platelets has been examined as a new venue for the treatment of Alzheimer’s disease and cilostazol is a well-established antiplatelet agent currently used for the secondary prevention of cerebral infarction [22, 23].

In the left hemisphere, cilostazol decreased neurodegeneration (Fig. 2B) and increased synaptic transmission (Fig. 2C), both suggestive of an improvement to the neurodegenerative phenotype. Furthermore, Gβγ signaling was asymmetrically increased in the left and decreased in the right cerebral hemisphere following cilostazol treatment (Fig. 1D). G-protein coupled receptors are highly diverse membrane proteins that participate in the transduction of external signals to various subcellular compartments via trimeric G-proteins. G-protein signaling in the central nervous system has been implicated in nuclear gene expression and cytoskeletal reorganization, processes that significantly contribute to synaptic plasticity and memory [24]. To further investigate whether these processes were on- or off-target effects of cilostazol was beyond the scope of the present proof-of-concept study.

In conclusion, our pharmacoproteomic study provides novel endophenotypic evidence on the hemisphere-specific pharmacologic effects of cilostazol. Future studies should account for hemisphere laterality with important implications in deciphering regional pharmacodynamic effects of
existing drugs thus uncovering new hemisphere-specific therapeutic targets.

Some limitations of the present study include the non-validated proteomic findings using an alternative technique (e.g., immunohistochemistry). Furthermore, sub-regions of the cerebral hemispheres (e.g., hippocampus, cortex, cerebellum) were not studied separately to identify regional asymmetric patterns of pharmacologic response to treatment. These constitute objectives in future studies.

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SUPPLEMENTARY MATERIAL

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