The dynamic mechanism of presenilin-function: Sensitive gate dynamics and loop unplugging control protein access - DTU Orbit (08/11/2017)

The dynamic mechanism of presenilin-function: Sensitive gate dynamics and loop unplugging control protein access There is no molecular explanation for the many presenilin 1 (PSEN1) mutations causing Alzheimer's disease, but both gain of function relating to amyloid production and loss of isolated PSEN1 function have been implied. We report here the first detailed dynamic all-atom model of mature PSEN1 from molecular dynamics in an explicit membrane with particular account of the as yet unexplored loop dynamics. We find that mature PSEN1 contains multiple distinct conformational states whereas non-mature PSEN1 is a typical one-state protein. We confirm a previously suggested gating mechanism, and find that the 106-131 loop acts as a "hinge" for the TM2 and TM6 "doors". More importantly, we identify an unplugging mechanism of the Exon 9 loop associated only with mature PSEN1. Proper opening of both the "gate" and "plug" in the membrane produces channel-like morphologies and access to the catalytic aspartates. Dynamically, these features seem linked. The long-range sensitivity of this gate-plug system to subtle conformational changes can explain why so many PSEN1 mutants cause disease. Reduced access and imprecise substrate cleavage associated with impaired gate-plug dynamics is directly illustrated by the effect of maturation in our work and could explain the overall reduction in Aß levels revealing PSEN1-only dynamics relating to e.g. its role as membrane channel. Thus, our identified gate-plug mechanism is relevant for designing PSEN1 modulating therapies for treatment of Alzheimer's disease within both the amyloid/ysecretase hypothesis and within the PSEN1 loss of function paradigm.

General information

State: Published

Organisations: Department of Chemistry Authors: Somavarapu, A. K. (Intern), Kepp, K. P. (Intern) Number of pages: 10 Pages: 147-156 Publication date: 2016 Main Research Area: Technical/natural sciences

Publication information

Journal: Neurobiology of Disease Volume: 89 ISSN (Print): 0969-9961 Ratings: BFI (2017): BFI-level 2 Web of Science (2017): Indexed Yes BFI (2016): BFI-level 2 Scopus rating (2016): SJR 2.707 SNIP 1.176 CiteScore 5.2 Web of Science (2016): Indexed yes BFI (2015): BFI-level 2 Scopus rating (2015): SJR 2.883 SNIP 1.269 CiteScore 5.17 BFI (2014): BFI-level 2 Scopus rating (2014): SJR 2.896 SNIP 1.278 CiteScore 5.19 BFI (2013): BFI-level 2 Scopus rating (2013): SJR 3.087 SNIP 1.414 CiteScore 5.93 BFI (2012): BFI-level 2 Scopus rating (2012): SJR 2.874 SNIP 1.397 CiteScore 5.93 BFI (2011): BFI-level 1 Scopus rating (2011): SJR 2.781 SNIP 1.306 CiteScore 5.67 BFI (2010): BFI-level 1 Scopus rating (2010): SJR 2.539 SNIP 1.161 BFI (2009): BFI-level 1 Scopus rating (2009): SJR 2.687 SNIP 1.122 BFI (2008): BFI-level 2 Scopus rating (2008): SJR 2.705 SNIP 1.057 Scopus rating (2007): SJR 2.42 SNIP 1.097 Scopus rating (2006): SJR 2.271 SNIP 1.091 Scopus rating (2005): SJR 2.343 SNIP 1.071 Scopus rating (2004): SJR 2.552 SNIP 1.093 Scopus rating (2003): SJR 2.695 SNIP 1.012

Scopus rating (2002): SJR 2.826 SNIP 1.1 Scopus rating (2001): SJR 2.582 SNIP 1.08 Scopus rating (2000): SJR 2.966 SNIP 1.181 Scopus rating (1999): SJR 3.859 SNIP 1.095 Original language: English Neurology, Alzheimer's disease, APP, Calcium, Conformation change, Protein structure, PSEN1 DOIs: 10.1016/j.nbd.2016.02.008 Source: FindIt Source-ID: 2291830392 Publication: Research - peer-review > Journal article – Annual report year: 2016