

Stable Expression of Tau Fluorescent Mutants as a Model to Screen Different Modulators of AD-related Kinases

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Abstract

The microtubule associated protein Tau is the main component of the neurofribillary tangles (NFT), aberrant structures that appear in the brain of Alzheimer's disease patients and other tauopathies, such as FTDP-17 or corticobasal degeneration. Tau protein binds to and stabilizes microtubules (MTs) but in pathological states, it aggregates and loses its important functions. These Tau aggregates are composed basically by hyperphosphorylated and truncated forms of tau. Multiple Tau gene mutations are pathogenic for hereditary FTDP-17 disease. These mutations have similar effects to hyperphosphorylation in Tau in AD and result in NFT formation.

We generated stably transfected cell lines that expressed different forms and mutants of Tau protein fused to turboGFP in order to study Tau behaviour in these conditions and perform a fluorescence-based cell assay for the screening of kinase modulators that affect the behaviour or location of Tau protein.

Results





Fig. 1: Different cell lines were stably transfected with the 0N3R, 0N4R and single and triple mutants isoforms of Tau fused to turboGFP. Tau showed a filamentous distribution, characteristic of microtubule-binding location. Bundling of microtubules induced by Tau transfection are also evident.







Fig. 2: A) U2OS cells stably transfected with the 0N4R isoform Wt (wild-type), SM (single mutant) and TM (triple mutant). Depending on the mutation, MT-bundle (arrows) amount varies.

B) Binding of Tau to MTs and consecuent bundle-formation is phosphorylation dependent. These Tau mutations have similar effects to hyperphosphorylation: the triple mutant has less affinity for microtubules and consequently forms less bundles. Wt and SM shows more bundling cells than TM at rest, so their capacity to form bundles after LiCI addition (GSK3 inhibitor) is limited.



Fig. 4: Screening in Tau-TM U2OS cells of representative and less toxic compounds from a library of 1200. Negative and positive (LiCl 30 mM) controls are shown in green and cyan, respectively. Some of the compounds have effect on Tau distribution: C3 and C26 shows no binding to MTs and in some cells Tau appears in nucleus; C9 and C49 shows more MT bundles; C22 shows an odd punctuate pattern around the nucleus.

References

Wagner et al., Journal of Cell Science, 1996. -Lu and Kosik, Molecular Biology of the Cell, 2001. -Tatebayashi et al., Journal of Cell Science, 2004. -Samsonov et al., Journal of Cell Science, 2004. -Ding et al., The Journal of Biological Chemistry, 2006. -Gendron and Petrucelli, Molecular neurodegeneration, 2009. -Tian et al., J Alzheimer Dis, 2009. -Kaufmann eta al., Cell Commun Signal, 2011.



Fig. 3: A) Not only kinase inhibitors like LiCl, but also PP2A modulators (Forskolin) and oxidative stress inducers (Na Arsenite) affect Tau-TM binding to MTs in U2OS cells.
B) Dose-response curve for the GSK3 inhibitor LiCl. IC50 value for LiCl was determined by treating U2OS Tau-TM model cells with concentrations from 100 mM to 100M during 2h. The intracellular bundle formation is quantified with a BD Pathway 855 High-Content Bioimager and Attovision software. Ic50 for LiCl is 11.75 mM and z' for this experiment was 0,80 +/-0,02.

Conclusions

- 1. Stably Tau transfected cells are viable.
- 2. Tau-turboGFP shows a filamentous distribution, characteristic of microtubule-binding location.
- 3. Fusion of turboGFP does not affect the functional properties of Tau.
- 4. Transfected Tau construct induces bundles of microtubules.
- 5. Mutations in Tau have different effects on bundle formation.
- 6. Distinct compounds over Tau-tGFP cells affect bundle formation.
- 7. BD Pathway 855 High-Content Bioimager is able to quantify number and intensity of bundles and therefore, study Tau behaviour and location.
- 8. According to the previos point, our assay could be valid for the screening of new modulators of AD-related kinases.