

1. **Research Unit:** Laboratory of Chemical Biology of Metal Ions
2. **Supervisor:** Prof. Dr hab. Wojciech Bal
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4. **Project title (English):** Establishment of partition of A β peptides related to Alzheimer's disease between the water and phospholipid phases and search for molecules modifying this partition.
5. **Project title (Polish):** Ustalenie partycji peptydów A β związanych z chorobą Alzheimera między fazą wodną i fosfolipidową i poszukiwanie cząsteczek modyfikujących tę partycję.
6. **Description of the project (up to 500 words):**

Alzheimer's Disease (AD) is a progressive and inevitably lethal dementia, caused by massive death of neurons in brain structures dealing with memory and emotions. The age is the main risk factor and no treatment can revert or even halt its progress. The best available recommendation is to maintain a healthy lifestyle. Although a number of aberrant processes were identified in AD brains and many drugs were developed to curb them, none has been successful in human clinical trials.

Soluble aggregated A β peptides are the best documented early toxic species in AD and many therapies were directed against these species. Those therapies, although successful in experimental animals, did not work in patients. This indicates that something crucial is missing in our understanding of A β physiology and pathology. The aim of this project is to clarify one such physiological aspect on the molecular level.

The A β peptides are soluble both in water and body fluids and in phospholipid bilayers constituting cellular membranes. When present in a membrane-like phospholipid environment, these peptides remain there and do not form toxic aggregates, but, once in aqueous solution, these aggregates are formed spontaneously within hours. In contrast, the disease progress takes many years, suggesting that until this time the A β peptides are not released to brain fluids in significant amounts. Based on prior evidence we propose that in normal brain the A β peptides reside mainly in the phospholipid environment and the molecules or processes affecting their biomembrane solubility may contribute to the progress of AD. These molecules or processes may be targeted by future AD therapies.

The aims of the project include determination how much of main A β peptides, A β 1-40, A β 1-42 and A β 4-42, can be dissolved in phospholipid bilayers of various compositions, modelling the physiological variability of cellular membranes. In such environment A β peptides remain monomeric, assume a partially α -helical conformation. Monomeric A β peptides in solution do not have a fixed shape, while the toxic A β aggregates contain β -sheet structures. We want to explain what causes the α -helical A β monomers to leave the membrane environment and get transformed into toxic aggregates. Such aggregates return to the membrane and damage it. Our further aim is to correlate the membrane leaving and membrane destructing propensity of A β peptides with its composition. We will also study how biological metal ions Cu(II) and Zn(II) contribute to these phenomena. Finally, we will search for molecules that could stabilize α -helices in A β peptides, thus reducing the risk of their aggregation and gain of toxicity. Such molecules may become AD drug candidates.

To achieve our goals we will chemically synthesize A β 1-40, A β 1-42 and A β 4-42 peptides and optimize the experimental procedures of their handling in order to assure good reproducibility of experiments. Our main studies will be performed using state of the art techniques of biomembrane research, including measurements of ionic currents flowing through microscale phospholipid bilayers and determining bilayer fluidity using spin labels detected by EPR. Finally, we will employ computer simulations and experiments to screen for molecules able to enhance α -helical A β .

7. **References related to conducted /planned research (maximum 3):**

- E. Stefaniak, E. Atrian-Blasco, W. Goch, L. Sabater, C. Hureau, W. Bal, The aggregation pattern of A β 1-40 is altered by the presence of N-truncated A β 4-40 and/or Cu(II) ions in a similar way via ionic interactions. *Chem. Eur. J.* 27, 2798-2809, 2021.
- M. Mital, K. Szutkowski, K. Bossak-Ahmad, P. Skrobecki, S. C. Drew, J. Poznański, I. Zhukov, T. Frączyk, W. Bal, The Palladium(II) Complex of A β ₄₋₁₆ as Suitable Model for Structural Studies of Biorelevant Copper(II) Complexes of N-Truncated Beta-Amyloids, *Int. J. Mol. Sci.* 21, art. no. 9200, 2020, doi:10.3390/ijms21239200
- N. E. Wezynfeld, E. Stefaniak, K. Stachucy, A. Drozd, D. Płonka, S. C. Drew, A. Krężel, W. Bal, Resistance of Cu(A β 4-16) to copper capture by metallothionein 3 supports a function of A β 4-42 peptide as a synaptic Cu^{II} scavenger, *Angew. Chem. Int. Ed.* 55, 8235–8238, 2016.

8. **Scholarship amount (net):** please contact the project supervisor.