Discovery and synthesis of novel inhibitors of the UPF1 helicase required for nonsense-mediated mRNA decay (NMD)

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Nonsense mRNA decay (NMD) recognizes and eliminates mRNA that contain a premature translation termination codon (PTC). The NMD pathway consists of a trimetric complex of hUPF proteins (UPF1, UPF2 and UPF3). In contrast, the ATP-dependent UPF1 helicase deserves special attention, as it performs the main regulatory functions in the NMD machinery.^[1] The activity of the pathway is observed in the pathophysiology of human genetic diseases, including malignancies^[2] For this reason, the UPF1 protein is an important molecular target in anti-cancer therapy. It should be emphasized that the mechanistic aspects of the NMD pathway are not fully understood. The presented approach focuses on the synthesis of potential UPF1 inhibitors small molecule chemicals selected by analyzing the results of a computer simulation. Using the Fischer indolization method, a number of heterocyclic derivatives based on the structural scaffold of 2,3,4,9-tetrahydro-1H-carbazole were obtained. The synthesized chemical compounds are potential UPF1 inhibitors. In vitro evaluation of active derivative ligands will be checked using thermal shift assay (TSA) and cellular thermal shift assay (CETSA).



[1] Kalathiya, U.; Padariya, M.; Pawlicka, K.; Verma, C. S.; Houston, D.; Hupp, T. R.; Alfaro, J. A. Insights into the Effects of Cancer Associated Mutations at the UPF2 and ATP-Binding Sites of NMD Master Regulator: UPF1. Int. J. Mol. Sci., 20, 5644 (**2019**)

[2] Nogueira, G.; Fernandes, R.; García-Moreno, J.F. et al. *Nonsense-mediated RNA decay and its bipolar function in cancer*. Mol Cancer., 20, 72 (**2021**)