

Moriuchi, H., Koda, N., Okuda-Ashitaka, E., Daiyasu, H., Ogasawara, K., Toh, H., ... & Watanabe, K. (2008). Molecular characterization of a novel type of prostamide/prostaglandin F synthase, belonging to the thioredoxin-like superfamily. *Journal of biological chemistry*, 283(2), 792-801.

Morris, G. M., Goodsell, D. S., Halliday, R. S., Huey, R., Hart, W. E., Belew, R. K., & Olson, A. J. (1998). Automated docking using a Lamarckian genetic algorithm and an empirical binding free energy function. *Journal of computational chemistry*, 19(14), 1639-1662.

Morris, G. M., Huey, R., Lindstrom, W., Sanner, M. F., Belew, R. K., Goodsell, D. S., & Olson, A. J. (2009). AutoDock4 and AutoDockTools4: Automated docking with selective receptor flexibility. *Journal of computational chemistry*, 30(16), 2785-2791.

Muhi-eldeen Z., Essentials of medicinal chemistry update, 2^{ed} Edition, 2010/2011.

Nelson, R. J. (2005). *An introduction to behavioral endocrinology*. Sinauer Associates.

Ogorochi, T., Ujihara, M., & Narumiya, S. (1987). Purification and Properties of Prostaglandin H-E Isomerase from the Cytosol of Human Brain: Identification as Anionic Forms of Glutathione S-Transferase. *Journal of neurochemistry*, 48(3), 900-909.

RCSB Protein Data Bank. <http://www.pdb.org/>.