

Rare diseases but common problems: Shwachman Diamond Syndrome from genetics to therapy through the combination of nanoscale techniques

Dritan Siliqi¹

Giuseppe Mangiatordi¹, Nuria Sánchez-Puig², Abril Gijsbers³, Pietro Delre¹ and Michele Saviano¹

¹ Institute of Crystallography , National Council for Research, Via G. Amendola 122/O, Bari, Italy

² Instituto de Química, Universidad Nacional Autónoma de México, Ciudad Universitaria, Ciudad de México, Mexico

³ The Maastricht Multimodal Molecular Imaging Institute (M4I), Division of Nanoscopy, Maastricht University, Maastricht, The Netherlands

dritan.siliqi@ic.cnr.it

Shwachman-Diamond Syndrome (SDS) is a ribosomopathy with a wide spectrum of clinical presentations [1] associated with the loss of function of Shwachman-Bodian-Diamond Syndrome (SBDS) protein [2] and as we described for the first time [3], the Elongation Factor-Like 1 (EFL1). Together, these proteins remove the anti-association factor eIF6 from the surface of the pre-60S ribosomal subunit to promote the formation of mature ribosomes. Due to the lack of knowledge of the molecular mechanisms responsible for SDS pathogenesis, current therapy is nonspecific and focuses only at alleviating the symptoms. For that reason, we studied [4] the interaction mechanism of the proteins in solution and demonstrated that binding SBDS*EFL1 consists of two independent and cooperative events, with domains 2–3 of SBDS directing the initial interaction with EFL1, followed by docking of domain 1. In solution, both proteins exhibited large flexibility and consisted of an ensemble of conformations, as demonstrated by Small Angle X-ray Scattering (SAXS) experiments [4, 5]. SAXS is a powerful technique for structural investigation of macromolecules in solution as for nanoparticles in solution or in solid state. Building on the recent observation that EFL1 single-point mutations clinically manifest as SDS-like phenotype, we carried out comparative Molecular Dynamics (MD) simulations on three mutants, T127A, M882K and R1095Q and wild type EFL1 [6] combining with SAXS experiments. This study supports the notion that EFL1 function is governed by an allosteric mechanism involving the concerted action of GTPase domains and can help point towards new approaches to SDS treatment.

References

- [1] Y. Dror, J. Donadieu, J. Kogelmeier, J. Dodge, S. Toiviainen-Salo, O. Makitie, E. Kerr, C. Zeidler, A. Shimamura, N. Shah, M. Cipolli, T. Kuijpers, P. Durie, J. Rommens, L. Siderius, J.M. Liu (2011). Draft consensus guidelines for diagnosis and treatment of Shwachman-Diamond syndrome, *Ann N Y Acad Sci* 1242, 40-55.
- [2] Boocock, G. R. B., Morrison, J. A., Popovic, M., Richards, N., Ellis, L., Durie, P. R., & Rommens, J. M. (2003). Mutations in SBDS are associated with Shwachman-Diamond Syndrome. *Nature Genetics*, 33(1), 97–101
- [3] Stepensky, P., Chacón-Flores, M., Kim, K. H., Abuzaitoun, O., Bautista-Santos, A., Simanovsky, N., Siliqi, D., Altamura, D., Méndez-Godoy, A., NaserEddin, N., Dor, T., Charrow, J., Sánchez-Puig, N., Elpele. O. (2017). Mutations in EFL1, an SBDS partner, are associated with infantile pancytopenia, exocrine pancreatic insufficiency and skeletal anomalies in a Shwachman- Diamond like syndrome. *J Med Genet*, 0: 1-9.
- [4] Gijsbers A, Montagut D, Méndez-Godoy A, Altamura D, Saviano M, Siliqi D*, Sánchez-Puig N*(2018). Interaction of the GTPase Elongation Factor Like-1 with the Shwachman-Diamond Syndrome Protein and Its Missense Mutations. *Int J Mol Sci* 19(12):4012.
- [5] Siliqi, D.; Foadi, J.; Mazzorana, M.; Altamura, D.; Méndez-Godoy, A.; Sánchez-Puig, N. (2018). Conformational Flexibility of Proteins Involved in Ribosome Biogenesis: Investigations via Small Angle X-ray Scattering (SAXS). *Crystals*, 8, 109.
- [6] P. Delre, D. Alberga, A. Gijsbers, N. Sánchez-Puig, O. Nicolotti, M. Saviano, D. Siliqi* & G. Felice Mangiatordi* (2019) Exploring the role of elongation Factor-Like 1 (EFL1) in Shwachman-Diamond syndrome through molecular dynamics, *Journal of Biomolecular Structure and Dynamics*, DOI: 10.1080/07391102.2019.1704883.