

Publication List – Thomas Monecke

* shared first authorships; # corresponding authorship

16. Zinggrebe, J., Moepps, B., **Monecke, T.**, Gierschik, P., Schlichtig, F., Barth, T.F.E., Strauß, G., Boldrin, E., Posovszky, C., Schulz, A., Beringer, O., Rieser, E., Jacobsen, E.M., Lorenz, M.R., Schwarz, K., Pannicke, U., Walczak, H., Niessing, D., Schuetz, C., Fischer-Posovszky, P., Debatin, K.M. (2022) Compound heterozygous variants in OTULIN are associated with fulminant atypical late-onset ORAS. *EMBO Mol Med*, 14(3), e14901.
15. Behrens, G., Edelmann, S.L., Raj, T., Kronbeck, N., **Monecke, T.**, Davydova, E., Wong, E.H., Kifinger, L., Giesert, F., Kirmaier, M.E., Hohn, C., de Jonge, L.S., Pisfil, M.G., Fu, M., Theurich, S., Feske, S., Kawakami, N., Wurst, W., Niessing, D., Heissmeyer, V. (2021) Disrupting Roquin-1 interaction with Regnase-1 induces autoimmunity and enhances antitumor responses. *Nat Immunol*, 22(12), 1563-1576.
14. Netz, E., Dijkstra, T.M.H., Sachsenberg, T., Zimmermann, L., Walzer, M., **Monecke, T.**, Ficner, R., Dybkov, O., Urlaub, H. and Kohlbacher, O. (2020) OpenPepXL: An open-source tool for sensitive identification of cross-linked peptides in XL-MS. *Mol Cell Proteomics*, 19(12), 2157-2168.
13. **Monecke, T.**, Dickmanns, A., Weiss, M.S., Port, S.A., Kehlenbach, R.H., Ficner, R. (2015) Combining dehydration, construct optimization and improved data collection to solve the crystal structure of a CRM1-RanGTP-SPN1-Nup214 quaternary nuclear export complex. *Acta Crystallogr F*, 71(12), 1481-7.
12. Port, S.A.* **Monecke, T.***, Dickmanns, A., Spillner, C., Hofele, R., Urlaub, H., Ficner, R., Kehlenbach, R. (2015) Structural and functional characterization of CRM1-Nup214 interactions reveals multiple FG-binding sites involved in nuclear export. *Cell Rep*, 13(4), 690-702.
11. Dickmanns, A., **Monecke, T.**, Ficner, R. (2015) Structural Basis of Targeting the Exportin CRM1 in Cancer. *Cells*, 4(3), 538-68.
10. **Monecke, T.**, Dickmanns, A., Ficner, R. (2014) Allosteric control of the exportin CRM1 unraveled by crystal structure analysis. *FEBS J*, 281(18), 4179-94.
9. Liu, Y., Neumann, P., Kuhle, B., **Monecke, T.**, Schell, S., Chari, A., Ficner, R. (2014) Translation initiation factor eIF3b contains a nine-bladed β -propeller and interacts with the 40S ribosomal subunit. *Structure*, 22(6), 923-930.
8. **Monecke, T.#**, Buschmann, J., Neumann, P., Wahle, E., Ficner, R. (2014) Crystal Structures of the Novel Cytosolic 5'-Nucleotidase IIIB Explain Its Preference for m⁷GMP. *PLoS ONE*, 9(3), e90915.
7. Dölker, N., Blanchet, C.E., Voß, B., Haselbach, D., Kappel, C., **Monecke, T.**, Svergun, D., Stark, H., Ficner, R., Zachariae, U., Grubmüller, H., Dickmanns, A. (2013) Structural Determinants and Mechanism of Mammalian CRM1 Allostery. *Structure*, 21(8), 1350-1360.
6. **Monecke, T.**, Haselbach, D., Voß, B., Russek, A., Neumann, P., Thomson, E., Hurt, E., Zachariae, U., Stark, H., Grubmüller, H., Dickmanns, A., Ficner, R. (2013) Structural basis for cooperativity of CRM1 export complex formation. *Proc Natl Acad Sci USA*, 110(3), 960-965.
5. Güttler, T., Madl, T., Neumann, P., Deichsel, D., Corsini, L., **Monecke, T.**, Ficner, R., Sattler, M., Görlich, D. (2010) NES consensus redefined by structures of PKI-type and Rev-type nuclear export signals bound to CRM1. *Nat Struct & Mol Biol*, 17(11), 1367-1376.

4. **Monecke, T.***, Güttler, T.*, Neumann, P., Dickmanns, A., Görlich, D., Ficner, R. **(2009)** Crystal structure of the nuclear export receptor CRM1 in complex with Snurportin1 and RanGTP. *Science*, 324(5930), 1087-1091.
3. **Monecke, T.**, Dickmanns, A., Ficner, R. **(2009)** Structural basis for m⁷G-cap hypermethylation of small nuclear, small nucleolar and telomerase RNA by the dimethyltransferase TGS1. *Nucleic Acids Res*, 37(12), 3865-3877.
2. **Monecke, T.**, Dickmanns, A., Strasser, A., Ficner, R. **(2009)** Structure analysis of the conserved methyltransferase domain of human trimethylguanosine synthase TGS1. *Acta Crystallogr D*, 65(4), 332-8.
1. **Monecke, T.***, Schell, S.*, Dickmanns, A., Ficner, R. **(2008)** Crystal structure of the RRM domain of poly(A)-specific ribonuclease reveals a novel m⁷G-cap-binding mode. *J Mol Biol*, 382(4), 827-834.