

Dr. Christoph W. Müller



02/04/1960, Herrenberg, Germany

Structural and Computational Biology Unit
European Molecular Biology Laboratory (EMBL)
D-69117 Heidelberg, Germany

Phone: +49(0)6221 387 8320

Fax: +49(0)6221 387 8591

Email: christoph.mueller@embl.de

Joint Head of Unit and Senior Scientist

SCIENTIFIC VITA

1986 Diplom-Chemiker, Albert-Ludwigs-Universität Freiburg, Germany
1991 Ph.D. thesis in the laboratory of Prof. G.E. Schulz, University of Freiburg:
 "Wildtype and mutant structures of adenylate kinase from *Escherichia coli*."
1991 Postdoctoral Fellow at University of Freiburg, Germany,
 Advisor: Prof. G.E. Schulz
1992-1995 Postdoctoral Fellow at Harvard University, Cambridge, U.S.A.
 Advisor: Prof. S.C. Harrison
1995-2000 Group Leader, EMBL Grenoble Outstation
1997 Habilitation at the University Joseph Fourier, Grenoble, France
2000-2007 Senior Scientist and Deputy Head of the Outstation, EMBL Grenoble
Since 2007 Joint Head of the Structural and Computational Biology Unit, EMBL Heidelberg

AWARDS

2003 Steinhofer-Award of the Steinhofer Foundation, Freiburg University, Germany
2005 Member of the European Molecular Biology Organisation (EMBO)

COORDINATING FUNCTIONS

2000-2007 Deputy Head of the EMBL Grenoble Outstation
2002-2007 Member of the PSB Grenoble Steering committee
Since 2007 Joint Head of the Structural and Computational Biology Unit
Since 2007 EMBL representative in the Heidelberg Molecular Life Science (HMLS) council
Since 2007 Member of the EMBO course committee
Since 2008 INSTRUCT - Working group leader - light microscopy
Since 2009 Local coordinator of the research infrastructure project PCUBE

Reviewing activities for the following agencies/institutes: Aeres - France, DFG, Wellcome Trust, MRC, Austrian Science Fund (FWF), HFSP, CNIO Madrid, IMBB Crete, PSI Villingen, Institute Pasteur, Nordic Partnership for Molecular Medicine

FIELDS OF INTEREST

Structural biology, X-ray crystallography, single-particle electron microscopy, transcriptional regulation, molecular machines, RNA polymerases, chromatin remodeling factors, transcription factors, nuclear transport.

CURRENTLY FUNDED PROJECTS

Infrastructure for Protein Production Platforms (PCUBE), Integrated Project 3D-Repertoire, EU Marie-Curie training fellowship, EMBO, INSTRUCT RTD project.

PUBLICATIONS (10 selected recent publications):

Fernández-Tornero, C., Böttcher, B., Rashid, U.J., Steuerwald, U., Flörchinger, B., Devos, D.P., Lindner, D., **Müller, C.W.** (2010). Conformational flexibility of RNA polymerase III during transcriptional elongation. *EMBO J.* 29, 3762-72.

Stirnemann, C.U., Petsalaki, E., Russell, R.B., **Müller, C.W.** (2010). WD40 proteins propel cellular networks. *TIBS* 35, 565-74

Morinière, J., Rousseaux, S., Steuerwald, U., Soler-López, M., Curtet, S., Vitte, A-L., Govin, J., Gaucher, J., Sadoul, K., Hart, D.J., Krijgsveld, J., Khochbin, S., **Müller, C.W.** & Petosa, C. (2009). Cooperative binding of two acetylation marks on a histone tail by a single bromodomain. *Nature* 461, 664-668.

Grimm, C., Matos, R., Ly-Hartig, N., Steuerwald, U., Lindner, D., Rybin, V., Müller, J. and **Müller C.W.** (2009). Molecular recognition of histone lysine methylation by the Polycomb group repressor dSfmbt. *EMBO J.* 28, 1965-1977.

Hamès, C., Ptchelkine, D., Grimm, C., Thevenon, E., Moyroud, E., Gérard, F., Martiel, J.L., Benlloch, R., Parcy, F., **Müller, C.W.** (2008). Structural basis for LEAFY floral switch function and similarity with helix-turn-helix proteins. *EMBO J.* 27, 2628-37.

Fernandez-Tornero, C., Böttcher, B., Riva, M., Carles, C., Steuerwald, U., Ruigrok, R.W., Sentenac, A., **Müller, C.W.**, Schoehn, G. (2007). Insights into transcription initiation and termination from the electron microscopy structure of yeast RNA polymerase III. *Mol Cell.* 25, 813-23.

Mylona, A., Fernandez-Tornero, C., Legrand, P., Haupt, M., Sentenac, A., Acker, J., **Müller, C.W.** (2006). Structure of the $\tau 60/\Delta\tau 91$ subcomplex of yeast transcription factor IIIC: insights into pre-initiation complex assembly. *Mol. Cell* 24, 221-232.

Petosa, C., Patrice M., Baudin, F., Moulin, M. Artero, J.B., **Müller, C.W.** (2006). Structural basis of lytic cycle activation by the Epstein-Barr virus ZEBRA protein. *Mol. Cell* 21, 565-572.

Petosa, C., Schoehn, G., Askjaer, P., Bauer, U., Moulin, M., Steuerwald, U., Soler López, M., Baudin, F., Mattaj, I.W., **Müller, C.W.** (2004). Architecture of CRM1/Exportin1 suggests how cooperativity is achieved during formation of a nuclear export complex. *Mol. Cell* 16, 761-775.

Soler-Lopez, M., Petosa, C., Fukuzawa, M., Ravelli, R., Williams, J.G., **Müller, C.W.** (2004). Structure of an activated *Dictyostelium* STAT in its DNA-unbound form. *Mol. Cell* 13, 791-04.