

Publications

87 research publications, 18 reviews and editorials, 2 book chapters
(citations in WebOfScience Core Collection: 5521; H-index: 39)

Research publications

- 87) Pinto A, El Ali Z, Moniot S, Tamborini L, **Steegborn C**, Foresti R, De Micheli C (2018) Effects of 3-Bromo-4,5-dihydroisoxazole Derivatives on Nrf2 Activation and Heme Oxygenase-1 Expression. *ChemistryOpen* 7, 858-864.
- 86) You W, **Steegborn C** (2018) Structural Basis of Sirtuin 6 Inhibition by the Hydroxamate Trichostatin A: Implications for Protein Deacetylase Drug Development. *J. Med. Chem.*, in press. doi: 10.1021/acs.jmedchem.8b01455
- 85) Iachettini S, Trisciuglio D, Rotili D, Lucidi A, Salvati E, Zizza P, Di Leo L, Del Bufalo D, Ciriolo MR, Leonetti C, **Steegborn C**, Mai A, Rizzo A, Biroccio A (2018) Pharmacological activation of SIRT6 triggers lethal autophagy in human cancer cells. *Cell Death Dis.* 9, 996.
- 84) D. Kalbas, S. Liebscher, T. Nowak, M. Meleshin, M. Pannek, C. Popp, Z. Alhalabi, F. Bordusa, W. Sippl, **C. Steegborn**, M. Schutkowski (2018) Potent and Selective Inhibitors of Human Sirtuin 5. *J. Med. Chem.* 61, 2460-2471.
- 83) M. Pannek, Z. Simic, M. Fuszard, M. Meleshin, D. Rotili, A. Mai, M. Schutkowski, **C. Steegborn** (2017) Crystal structures of the mitochondrial deacetylase Sirtuin 4 reveal isoform-specific acyl recognition and regulation features. *Nat. Commun.* 8:1513.
- 82) Rajabi N, Auth M, Troelsen KR, Pannek M, Bhatt D, Fontenas M, Hirshey MD, **Steegborn C**, Madsen AS, Olsen CA (2017) Mechanism-based Inhibitors of the Human Sirtuin 5 Deacetylase: Structure-Activity Relationship, Biostructural, and Kinetic Insight. *Angew. Chem. Int. Ed. Engl.* 56, 14836-14841.
- 81) Quinti L, Dayalan Naidu S, Träger U, Chen X, Kegel-Gleason K, Llères D, Connolly C, Chopra V, Low C, Moniot S, Sapp E, Tousley AR, Vodicka P, Van Kanegan MJ, Kaltenbach LS, Crawford LA, Fuszard M, Higgins M, Miller JRC, Farmer RE, Potluri V, Samajdar S, Meisel L, Zhang N, Snyder A, Stein R, Hersch SM, Ellerby LM, Weerapana E, Schwarzschild MA, **Steegborn C**, Leavitt BR, Degterev A, Tabrizi SJ, Lo DC, DiFiglia M, Thompson LM, Dinkova-Kostova AT, Kazantsev AG (2017) A KEAP1-modifying small molecule reveals muted NRF2 signaling responses in neural stem cells from Huntington disease patients. *Proc. Natl Acad. Sci. USA* 114, E4676-E4685.
- 80) C.G.F. Graf, C. Schulz, M. Schmälzlein, C. Heinlein, M. Mönnich, L. Perkams, M. Püttner, I. Boos, M. Hessefort, J.N. Lombana Sanchez, M. Weyand, **C. Steegborn**, B. Breiden, K. Ross, G. Schwarzmann, K. Sandhoff, C. Unverzagt (2017) Synthetic Glycoforms reveal Carbohydrate-Dependent Bioactivity of Human Saposin D. *Angew. Chem. Int. Ed. Engl.* 56, 5252-5257.
- 79) Li J, Bonkowski MS, Moniot S, Zhang D, Hubbard BP, Ling AJ, Rajman LA, Qin B, Lou Z, Gorbunova V, Aravind L, **Steegborn C**, Sinclair DA (2017) A conserved NAD⁺-binding pocket that regulates protein-protein interactions during aging. *Science* 355, 1312-1317.
- 78) de Oliveira RM, Vicente Miranda H, Francelle L, Pinho R, Szegö ÉM, Martinho R, Munari F, Lázaro DF, Moniot S, Guerreiro P, Fonseca L, Marijanovic Z, Antas P, Gerhardt E, Enguita FJ, Fauvet B, Penque D, Pais TF, Tong Q, Becker S, Kübler S, Lashuel HA, **Steegborn C**, Zweckstetter M, Outeiro TF (2017) The mechanism of sirtuin 2-mediated exacerbation of alpha-synuclein toxicity in models of Parkinson disease. *PLoS Biol.* 2017 15:e2000374.
- 77) Moniot S, Forgione M, Lucidi A, Hailu GS, Nebbioso A, Carafa V, Baratta F, Altucci L, Giacché N, Passeri D, Pellicciari R, Mai A, **Steegborn C**, Rotili D (2017) Development

of 1,2,4-Oxadiazoles as Potent and Selective Inhibitors of the Human Deacetylase Sirtuin 2: Structure-Activity Relationship, X-ray Crystal Structure, and Anticancer Activity. *J. Med. Chem.* 60, 2344-2360

76) Sundriyal S, Moniot S, Mahmud Z, Yao S, Di Fruscia P, Reynolds CR, Dexter DT, Sternberg MJ, Lam EW, **Steegborn C**, Fuchter MJ (2017) Thienopyrimidinone Based Sirtuin-2 (SIRT2)-Selective Inhibitors Bind in the Ligand Induced Selectivity Pocket. *J. Med. Chem.* 60, 1928-1945.

75) You W, Rotili D, Li TM, Kambach C, Meleshin M, Schutkowski M, Chua KF, Mai A, **Steegborn C**. (2017) Structural Basis of Sirtuin 6 Activation by Synthetic Small Molecules. *Angew. Chem. Int. Ed. Engl.* 56, 1007-1011.

74) L. Ramos-Espiritu, S. Kleinboelting, F.A. Navarrete, A. Alvau, P.E. Visconti, F. Valsecchi, A. Starkov, G. Manfredi, H. Buck, C. Adura, J.H. Zippin, J. van den Heuvel, J.F. Glickman, **C. Steegborn**, L.R. Levin, J. Buck (2016) Discovery of LRE1 as a specific and allosteric inhibitor of soluble adenylyl cyclase. *Nature Chem. Biol.* 12, 838-44.

73) L. Quinti, M. Casale, S. Moniot, T.F. Pais, L.S. Kaltenbach, J. Pallos, R.G. Lim, S.D. Naidu, H. Runne, L. Meisel, N.A. Rauf, D. Leyfer, M.M. Maxwell, E. Saiah, J.E. Landers, R. Luthi-Carter, R. Abagyan, A.T. Dinkova-Kostova, **C. Steegborn**, J.L. Marsh, D.C. Lo, L.M. Thompson, A.G. Kazantsev (2016) SIRT2- and NRF2-targeting thiazole-containing compound with therapeutic activity in Huntington's disease models. *Cell Chem. Biol.* 23, 849-61.

72) Kleinboelting S, Ramos-Espiritu L, Buck H, Colis L, van den Heuvel J, Glickman JF, Levin LR, Buck J, **Steegborn C**. (2016) Bithionol Potently Inhibits Human Soluble Adenylyl Cyclase through Binding to the Allosteric Activator Site. *J. Biol. Chem.* 291, 9776-9784.

71) Schuster S, Roessler C, Meleshin M, Zimmermann P, Simic Z, Kambach C, Schiene-Fischer C, **Steegborn C**, Hottiger MO, Schutkowski M. (2016) A continuous sirtuin activity assay without any coupling to enzymatic or chemical reactions. *Sci. Rep.* 6:22643.

70) Valente S, Mellini P, Spallotta F, Carafa V, Nebbioso A, Polletta L, Carnevale I, Saladini S, Trisciuglio D, Gabellini C, Tardugno M, Zwergel C, Cencioni C, Atlante S, Moniot S, **Steegborn C**, Budriesi R, Tafani M, Del Bufalo D, Altucci L, Gaetano C, Mai A. (2016) 1,4-Dihydropyridines Active on the SIRT1/AMPK Pathway Ameliorate Skin Repair and Mitochondrial Function and Exhibit Inhibition of Proliferation in Cancer Cells. *J. Med. Chem.* 59, 1471-1491.

69) U. Schweizer & **C. Steegborn** (2015) Thyroid hormones—From Crystal Packing to Activity to Reactivity. *Angew. Chem. Int. Ed. Engl.* 54, 12856-8.

68) Z. Simic, M. Weiwig, A. Schierhorn, **C. Steegborn**, M. Schutkowski (2015) The ε-amino group of protein lysine residues is highly susceptible to nonenzymatic acylation by several physiological acyl-CoA thioesters. *ChemBioChem*, 16, 2337-47.

67) C. Roessler, C. Tueting, M. Meleshin, D. Rauh, **C. Steegborn**, M. Schutkowski (2015) A novel continuous assay for the deacetylase Sirtuin 5 and other deacetylases. *J. Med. Chem.* 58, 7217-23.

66) A. Banerjee, R.S. Adolph, J. Gopalakrishnapai, S. Kleinboelting, C. Emmerich, **C. Steegborn**, S.S. Visweswariah (2015) A Universal Stress Protein (USP) in Mycobacteria binds cAMP. *J. Biol. Chem.* 290, 12731-43.

65) L. Polletta, E. Vernucci, I. Carnevale, T. Arcangeli, D. Rotili, S. Palmerio, **C. Steegborn**, T. Nowak, M. Schutkowski, L. Pellegrini, L. Sansone, L. Villanova, A. Runci, B. Pucci, E. Morgante, M. Fini, A. Mai, M. A. Russo, M. Tafani (2015). SIRT5 regulation of ammonia-induced autophagy and mitophagy. *Autophagy* 11, 253-70.

- 64) T. Rumpf, M. Schiedel, B. Karaman, C. Roessler, B.J. North, A. Lehotzky, J. Oláh, K.I. Ladwein, K. Schmidkunz, M. Gajer, M. Pannek, **C. Steegborn**, D.A. Sinclair, S. Gerhardt, J. Ovádi, M. Schutkowski, W. Sippl, O. Einsle, M. Jung (2015). Selective Sirt2 inhibition by ligand-induced rearrangement of the active site. *Nat. Commun.* 6:6263
- 63) X. Chen, P. Wales, L. Quinti, F. Zuo, S. Moniot, F. Herisson, N.A. Rauf, H. Wang, R.B. Silverman, C. Ayata, M.M. Maxwell, **C. Steegborn**, M.A. Schwarzschild, T.F. Outeiro, A.G. Kazantsev (2015). The Sirtuin-2 Inhibitor AK7 Is Neuroprotective in Models of Parkinson's Disease but Not Amyotrophic Lateral Sclerosis and Cerebral Ischemia. *PLoS One* 10, e0116919
- 62) P. Di Fruscia, E. Zacharioudakis, C. Liu, S. Moniot, S. Laohasinnarong, M. Khongkow, I.F. Harrison, K. Koltsida, C.R. Reynolds, K. Schmidkunz, M. Jung, K.L. Chapman, **C. Steegborn**, D.T. Dexter, M.J. Sternberg, E.W. Lam, M.J. Fuchter (2014) The Discovery of a Highly Selective 5,6,7,8-Tetrahydrobenzo[4,5]thieno[2,3-d]pyrimidin-4(3H)-one SIRT2 Inhibitor that is Neuroprotective in an in vitro Parkinson's Disease Model. *ChemMedChem* 10, 69-82.
- 61) B. Fränzel, F. Fischer, **C. Steegborn**, D.A. Wolters (2014) Proteinase K improves quantitative acylation studies. *Proteomics* 15, 44-7.
- 60) Roessler C, Nowak T, Pannek M, Gertz M, Nguyen GT, Scharfe M, Born I, Sippl W, **Steegborn C**, Schutkowski M. (2014) Chemical Probing of the Human Sirtuin 5 Active Site Reveals Its Substrate Acyl Specificity and Peptide-Based Inhibitors. *Angew. Chem. Int. Ed.* 53, 10728-32.
- 59) S. Kleinboelting, J. van den Heuvel, **C. Steegborn** (2014) Structural analysis of human soluble adenylyl cyclase and crystal structures of its nucleotide complexes - implications for cyclase catalysis and evolution. *FEBS J.* 281, 4151-64.
- 58) U. Schweizer, C. Schlicker, D. Braun, J. Köhrle, **C. Steegborn** (2014) Crystal structure of mammalian selenocysteine-dependent iodothyronine deiodinase suggests a peroxiredoxin-like catalytic mechanism. *Proc. Natl Acad. Sci. USA* 111, 10526-31.
- 57) M.H. Suhre, T. Scheibel, **C. Steegborn**, M. Gertz (2014) Crystallization and preliminary X-ray diffraction analysis of proximal thread matrix protein 1 (PTMP1) from *Mytilus galloprovincialis*. *Acta Crystallogr. Sect. F* 70, 769-72.
- 56) S. Kleinboelting, J. van den Heuvel, C. Kambach, M. Weyand, M. Leipelt, **C. Steegborn** (2014) Expression, purification, crystallization and preliminary X-ray diffraction analysis of a mammalian type 10 adenylyl cyclase. *Acta Crystallogr. Sect. F* 70, 467-9.
- 55) M.H. Suhre, M. Gertz, **C. Steegborn**, T. Scheibel (2014) Structural and functional features of a collagen-binding matrix protein from the mussel byssus. *Nat. Commun.* 5:3392.
- 54) S. Kleinboelting, A. Diaz, S. Moniot, J. van den Heuvel, M. Weyand, L.R. Levin, J. Buck, **C. Steegborn** (2014) Crystal structures of human soluble adenylyl cyclase reveal mechanisms of catalysis and of its activation through bicarbonate. *Proc. Natl Acad. Sci. USA* 111, 3727-32.
- 53) G.T.T. Nguyen, S. Schaefer, M. Gertz, M. Weyand, **C. Steegborn** (2013) Crystal structures of Sirt3 complexes with the resveratrol derivative 5-(2-(4-bromophenyl)vinyl)-1,3-benzenediol reveal binding sites and inhibition mechanism. *Chem. Biol.* 20:1375-85.
- 52) D. Rauh, F. Fischer, M. Gertz, M. Lakshminarasimhan, T. Bergbrede, F. Aladini, C. Kambach, C.F.W. Becker, J. Zerweck, M. Schutkowski, **C. Steegborn** (2013). An acetylome peptide microarray reveals specificities and deacetylation substrates for all human Sirtuin isoforms. *Nat. Commun.* 4:2327.

- 51) G.T.T. Nguyen, S. Schaefer, M. Gertz, M. Weyand, **C. Steegborn** (2013) Structures of human sirtuin 3 complexes with ADP-ribose and with carba-NAD⁺ and SRT1720: binding details and inhibition mechanism. *Acta Cryst. D* 69, 1423-32.
- 50) M. Gertz, F. Fischer, G.T.T. Nguyen, M. Lakshminarasimhan, M. Schutkowski, M. Weyand, **C. Steegborn** (2013) Ex-527 inhibits Sirtuins by exploiting their unique NAD⁺-dependent deacetylation mechanism. *Proc. Natl Acad. Sci. USA* 110:E2772-81.
- 49) G. Laurent, N.J. German, A.K. Saha, V.C. de Boer, M. Davies, T.R. Koves, N. Dephoure, F. Fischer, G. Boanca, B. Vaiteesvaran, S.B. Lovitch, A.H. Sharpe, I.J. Kurland, **C. Steegborn**, S.P. Gygi, D.M. Muoio, N.B. Ruderman, M.C. Haigis (2013) SIRT4 coordinates the balance between lipid synthesis and catabolism by repressing malonyl CoA decarboxylase. *Mol. Cell* 50, 686-98.
- 48) M. Lakshminarasimhan, U. Curth, S. Moniot, S. Mosalaganti, S. Raunser, **C. Steegborn** (2013) Molecular architecture of the human protein deacetylase Sirt1 and its regulation by AROS and resveratrol. *Biosci. Rep.* 33:e00037.
- 47) M. Lakshminarasimhan, D. Rauh, M. Schutkowski, **C. Steegborn** (2013) Sirt1 activation by resveratrol is substrate sequence-selective. *Aging* 5, 151-4.
- 46) S. Moniot, M. Schutkowski, **C. Steegborn** (2013) Crystal structure analysis of human Sirt2 and its ADP-ribose complex. *J. Struct. Biol.* 182, 136-43.
- 45) B. Suenkel, F. Fischer, **C. Steegborn** (2013) Inhibition of the human deacylase Sirtuin 5 by the indole GW5074. *Bioorg. Med. Chem. Lett.* 23, 143-6.
- 44) Gertz M, Nguyen GT, Fischer F, Suenkel B, Schlicker C, Fränzel B, Tomaschewski J, Aladini F, Becker C, Wolters D, **Steegborn C.** (2012) A molecular mechanism for direct sirtuin activation by resveratrol. *PLoS One* 7:e49761.
- 43) Fischer F, Gertz M, Suenkel B, Lakshminarasimhan M, Schutkowski M, **Steegborn C.** (2012) Sirt5 deacylation activities show differential sensitivities to nicotinamide inhibition. *PLoS One* 7:e45098.
- 42) S. Middelhaufe, M. Leipelt, L.R. Levin, J. Buck, **C. Steegborn** (2012) Identification of a haem domain in human soluble adenylate cyclase. *Biosci. Rep.* 32, 491-9.
- 41) H. Topal, N.B. Fulcher, J. Bitterman, E. Salazar, J. Buck, L.R Levin, M.J Cann, MC Wolfgang, **C. Steegborn** (2012). Crystal Structure and Regulation Mechanisms of the CyaB Adenylyl Cyclase from the Human Pathogen Pseudomonas aeruginosa. *J. Mol. Biol.* 416, 271-86.
- 40) C. Schlicker, G. Boanca, M. Lakshminarasimhan, **C. Steegborn** (2011). Structure-based development of novel sirtuin inhibitors. *Aging* 3, 852-72.
- 39) R. Acin-Perez, M. Russwurm, K. Günnewig, M. Gertz, G. Zoidl, L. Ramos, J. Buck, L.R. Levin, J. Rassow, G. Manfredi, **C. Steegborn** (2011). A Phosphodiesterase 2A Isoform Localized to Mitochondria Regulates Respiration. *J. Biol. Chem.* 286, 30423-32.
- 38) M. Russwurm, C. Schlicker, M. Weyand, D. Koesling, **C. Steegborn** (2011). Crystal structure of the GAF-B domain from human phosphodiesterase 5. *Proteins* 79, 1682-1687.
- 37) A. Hall, L. De Sordi, D.M. MacCallum, H. Topal, R. Eaton, J.W. Bloor, G.K. Robinson, L.R. Levin, J. Buck, Y. Wang, N.A.R. Gow, **C. Steegborn**, F.A. Mühlischlegel (2010). CO₂ Acts as a Signalling Molecule in Populations of the Fungal Pathogen Candida albicans. *PLoS Pathogens* 6:e1001193.
- 36) A. Innocenti, R.A. Hall, C. Schlicker, A. Scozzafava, **C. Steegborn**, F.A. Mühlischlegel, C.T. Supuran (2009) Carbonic anhydrase inhibitors. Inhibition and homology modeling studies of the fungal beta-carbonic anhydrase from Candida albicans with sulfonamides. *Bioorg. Med. Chem.* 17, 4503-4509.

- 35) M. Gertz, F. Fischer, M. Leipelt, D. Wolters, **C. Steegborn** (2009) Identification of Peroxiredoxin 1 as a novel interaction partner for the lifespan regulator protein p66^{Shc}. *Aging* 1, 254-265.
- 34) C. Schlicker, R.A. Hall, D. Vullo, S. Middelhaufe, M. Gertz, C.T. Supuran, F.A. Mühlischlegel, **C. Steegborn** (2009) Structure and inhibition of the CO₂-sensing carbonic anhydrase Can2 from the pathogenic fungus *Cryptococcus neoformans*. *J. Mol. Biol.* 385, 1207-1220.
- 33) A. Rauch, M. Leipelt, M. Russwurm, **C. Steegborn** (2008) Crystal structure of the guanylyl cyclase Cya2. *Proc. Natl. Acad. Sci. USA* 105, 15720-15725.
- 32) A. Innocenti, F.A. Mühlischlegel, R.A. Hall, **C. Steegborn**, A. Scozzafava, C.T. Supuran (2008) Carbonic anhydrase inhibitors. Inhibition of the beta-class enzymes from the fungal pathogens *Candida albicans* and *Cryptococcus neoformans* with simple anions. *Bioorg. Med. Chem. Lett.* 18, 5066-5070.
- 31) M. Tammenkoski, K. Koivula, E. Cusanelli, M. Zollo, **C. Steegborn**, A.A. Baykov, R. Lahti (2008) The human metastasis regulator protein h-prune is a short-chain exopolyphosphatase. *Biochemistry* 47, 9707-9713.
- 30) C. Schlicker, M. Gertz, D. Papatheodorou, B. Kachholz, C.F.W. Becker, **C. Steegborn** (2008) Substrates and regulation mechanisms for the human mitochondrial Sirtuins Sirt3 and Sirt5. *J. Mol Biol.* 382, 790-801.
- 29) C. Schlicker, A. Rauch, K.C. Hess, B. Kachholz, L.R. Levin, J. Buck, **C. Steegborn** (2008) Structure-based development of novel adenylyl cyclase inhibitors. *J. Med. Chem.* 51, 4456-4464.
- 28) M. Gertz, F. Fischer, D. Wolters, **C. Steegborn** (2008) Activation of the life-span regulator p66^{Shc} through reversible disulfide bond formation. *Proc. Natl Acad. Sci. USA* 105, 5705-5709.
- 27) L. Garzia, A. D'Angelo, A. Amoresano, S.K. Knauer, C. Cirulli, C. Campanella, R.H. Stauber, **C. Steegborn**, A. Iolascon, M. Zollo (2007) Phosphorylation of nm23H1 by CKI induces complex formation with h-prune and promotes cell motility. *Oncogene* 27, 1853-1864.
- 26) S. Middelhaufe, L. Garzia, U.-M. Ohndorf, B. Kachholz, M. Zollo, **C. Steegborn** (2007) Domain mapping on the human metastasis regulator protein h-Prune reveals a C-terminal dimerization domain. *Biochem. J.* 407, 199-205.
- 25) H.G. Mannherz, E. Ballweber, M. Galla, S. Villard, C. Granier, **C. Steegborn**, A. Schmidtmann, K. Jaquet, B. Pope, A.G.M. Weeds (2007) Mapping the ADF/Cofilin Binding Site on Monomeric Actin by Competitive Cross-linking and Peptide Array: Evidence for a Second Binding Site on Monomeric Actin. *J. Mol. Biol.* 366, 745-755.
- 24) E.G. Mogensen, G. Janbon, J. Chaloupka, **C. Steegborn**, M. Shun Fu, F. Moyrand, T. Klengel, D. S. Pearson, M. A. Geeves, J. Buck, L. R. Levin, and F. A. Mühlischlegel (2006) CO₂ sensing in *Cryptococcus neoformans* involves the carbonic anhydrase Can2. *Eukaryot. Cell* 5, 103-111.
- 23) **C. Steegborn**, T.N. Litvin, K.C. Hess, A.B. Capper, R. Taussig, J. Buck, L.R. Levin, H. Wu (2005) A novel mechanism for adenylyl cyclase inhibition from the crystal structure of its complex with catechol estrogen. *J. Biol. Chem.* 280, 31754 - 31759.
- 22) **C. Steegborn**, T.N. Litvin, L.R. Levin, J. Buck, H. Wu (2005) Bicarbonate activation of adenylyl cyclase via promotion of catalytic active site closure and metal recruitment. *Nature Struct. Mol. Biol.* 12, 32-37.
- 21) H. Ye, T.-S. Chen, X. Xu, M. Pennycooke, H. Wu, **C. Steegborn** (2004) Crystal

- structure of the putative adapter protein MTH1859. *J. Struct. Biol.* 148, 251-256.
- 20) G. Xu, R.L. Rich, **C. Steegborn**, T. Min, Y. Huang, D.G. Myszka, H. Wu (2003) Mutational analyses of the p35/caspase Interaction. A bowstring kinetic model of caspase inhibition by p35. *J. Biol. Chem.* 278, 5455-61.
- 19) M.A. Augustin, A.S. Reichert, H. Betat, R. Huber, M. Moerl, **C. Steegborn** (2003) Crystal Structure of the human CCA-adding enzyme: Insights into template-independent polymerization. *J. Mol. Biol.* 328, 985-994.
- 18) A. Messerschmidt, M. Wörbs, **C. Steegborn**, M.C. Wahl, R. Huber, B. Laber, T. Clausen (2003) What determines enzymatic specificity in the cys-met-metabolism PLP-dependent enzyme family: Crystal structure of cystathionine γ -lyase from yeast and intrafamilial structural comparison. *Biol. Chem.* 384, 373-386.
- 17) S.P. Stabler, **C. Steegborn**, M.C. Wahl, J. Oliveriusova, J.P. Kraus, R.H. Allen, C. Wagner, S.H. Mudd (2002) Elevated plasma total homocysteine in severe MAT I/III deficiency. *Metabolism* 51, 981-988.
- 16) W. Rossoll, A.-K. Kroening, U.-M. Ohndorf, **C. Steegborn**, S. Jablonka, M. Sendtner (2002) Specific interaction of Smn, the spinal muscular atrophy determining gene product, with hnRNP-R and gry-rbp/hnRNP-Q: a role for Smn in RNA processing in motor axons? *Hum. Mol. Genet.* 11, 93-105.
- 15) **C. Steegborn**, O. Danot, R. Huber, T. Clausen (2001) Crystal structure of transcription factor MalT domain III: A novel helix repeat fold implicated in regulated oligomerisation. *Structure* 9, 1051-1060.
- 14) **C. Steegborn**, B. Laber, A. Messerschmidt, R. Huber, T. Clausen (2001) Crystal structures of cystathionine γ -synthase inhibitor complexes rationalize the increased affinity of a novel inhibitor. *J. Mol. Biol.* 311, 789-801.
- 13) U.-M. Ohndorf, **C. Steegborn**, R. Knijff, P. Sondermann (2001) Contributions of the individual domains in human La protein to its RNA 3'-end binding activity. *J. Biol. Chem.* 276, 27188-27196.
- 12) A. Dorowski, A. Hofmann, **C. Steegborn**, M. Bociu, R. Huber (2001) Crystal structure of paprika ferredoxin-NADP⁺ reductase. Implications for the electron transfer pathway. *J. Biol. Chem.* 276, 9253-9263.
- 11) **C. Steegborn**, H. Schneider-Hassloff, M. Zeeb, J. Balbach (2000) Cooperativity of a protein folding reaction probed at multiple chain positions by real-time 2D NMR spectroscopy. *Biochemistry* 39, 7910-7919.
- 10) T. Clausen, J. Kaiser, **C. Steegborn**, R. Huber, D. Kessler (2000) Crystal structure of the cystine C-S lyase from *Synechocystis*: stabilization of cysteine persulfide for FeS cluster biosynthesis. *Proc. Natl Acad. Sci. USA* 97, 3856-3861.
- 9) T. Clausen, A. Schlegel, R. Peist, E. Schneider, **C. Steegborn**, Y.-S. Chang, A. Haase, G.P. Bourenkov, H.D. Bartunik, W. Boos (2000) X-ray structure of MalY from *Escherichia coli*: a pyridoxal 5'-phosphate-dependent enzyme acting as a modulator in *mal* gene expression. *EMBO J.* 19, 831-842.
- 8) V. Schreiber, **C. Steegborn**, T. Clausen, W. Boos, E. Richet (1999) A new mechanism for the control of a prokaryotic transcriptional regulator: antagonistic binding of positive and negative effectors. *Mol. Microbiol.* 35, 765-776.
- 7) T. Clausen, M.C. Wahl, A. Messerschmidt, R. Huber, J. Fuhrmann, B. Laber, W. Streber, **C. Steegborn** (1999) Cloning, purification and characterisation of cystathionine γ -synthase from *Nicotiana tabacum*. *Biol. Chem.* 380, 1237-1242.
- 6) **C. Steegborn**, A. Messerschmidt, B. Laber, W. Streber, R. Huber, T. Clausen (1999) The crystal structure of cystathionine γ -synthase from *Nicotiana tabacum* reveals its substrate and reaction specificity. *J. Mol. Biol.* 290, 983-996.

- 5) **C. Steegborn**, T. Clausen, P. Sondermann, U. Jacob, M. Worbs, S. Marinkovic, R. Huber, M.C. Wahl (1999) Kinetics and inhibition of recombinant human cystathionine γ -lyase - Toward the rational control of transsulfuration. *J. Biol. Chem.* 274, 12675-12684.
- 4) J. Balbach, **C. Steegborn**, T. Schindler, F.X. Schmid (1999) A protein folding intermediate of ribonuclease T₁ characterized at high resolution by 1D and 2D real-time NMR spectroscopy. *J. Mol. Biol.* 285, 829-842.
- 3) **C. Steegborn**, P. Skladal (1997) Construction and characterization of the direct piezoelectric immunosensor for atrazine operating in solution. *Biosens. Bioelectron.* 12, 19-27.
- 2) A.U. Metzger, T. Schindler, D. Willbold, M. Kraft, **C. Steegborn**, A. Volkmann, R.W. Frank, P. Roesch (1996) Structural rearrangements on HIV-1 Tat (32-72) TAR complex formation. *FEBS Lett.* 384, 255-259.
- 1) M. Sprinzl, **C. Steegborn**, F. Huebel, S. Steinberg (1996) Compilation of tRNA sequences and sequences of tRNA genes. *Nucleic Acids Res.* 24, 68-72.

Reviews & Editorials

- 18) Wiggins SV, Steegborn C, Levin LR, Buck J. (2018) Pharmacological modulation of the CO₂/HCO₃-/pH-, calcium-, and ATP-sensing soluble adenylyl cyclase. *Pharmacol. Ther.*, 190, 173-186.
- 17) H. Dai, D.A. Sinclair, J.L. Ellis, **C. Steegborn** (2018) Sirtuin activators and inhibitors: Promises, achievements, and challenges. *Pharmacol. Ther.* 188, 140-154.
- 16) **C. Steegborn** (2017) JSB Paper-of-the Year Award: "Sébastien Moniot et al. (2013) Crystal structure analysis of human Sirt2 and its ADP-ribose complex. *J. Struct. Biol.* 182, 136–43". *J. Struct. Biol.* 198, 1-2.
- 15) U. Schweizer, H. Towell, A. Vit, A. Rodriguez-Ruiz, **C. Steegborn** (2017) Structural aspects of thyroid hormone binding to proteins and competitive interactions with natural and synthetic compounds. *Mol. Cell. Endocrinol.* 55, R37-52.
- 14) M. Gertz, **C. Steegborn** (2016) Using mitochondrial sirtuins as drug targets: disease implications and available compounds. *Cell. Mol. Life Sci.* 73, 2871-9.
- 13) B. Suenkel, **C. Steegborn** (2016) Recombinant preparation, biochemical analysis, and structure determination of Sirtuin family histone/protein deacetylases. *Methods in Enzymology*, Vol. Enzymes of Epigenetics 573, 183-208.
- 12) Klionsky DJ, Abdelmohsen K, et al., **C. Steegborn**, et al. (2016) Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). *Autophagy* 12, 1-222.
- 11) U. Schweizer, **C. Steegborn** (2015) New insights into the structure and mechanism of iodothyronine deiodinases. *J. Mol. Endocrinol.* 55, R37-52.
- 10) **C. Steegborn** (2014) Structure, mechanism, and regulation of soluble adenylyl cyclases – similarities and differences to transmembrane adenylyl cyclases. *Biochim. Biophys. Acta - Molecular Basis of Disease* 1842, 2535-47.
- 9) M. Schutkowski, F. Fischer, C. Roessler, **C. Steegborn** (2014) New assays and approaches for discovery and design of Sirtuin modulators. *Expert Opin. Drug Discov.* 9, 183-99.
- 8) S. Moniot, M. Weyand, **C. Steegborn** (2012). Structures, substrates, and regulators of Mammalian sirtuins - opportunities and challenges for drug development. *Front. Pharmacol.* 3:16.

- 7) M. Lakshminarasimhan, **C. Steegborn** (2011) Emerging mitochondrial signaling mechanisms in physiology, aging processes, and as drug targets. *Exp. Gerontol.* 46, 174-7.
- 6) M. Gertz, **C. Steegborn** (2010) The mitochondrial apoptosis pathway and p66Shc--a regulatory redox enzyme or an adapter protein snuggling around? *Cell Cycle* 9, 4425-6.
- 5) J.A. Baur, D. Chen, E.N. Chini, K. Chua, H.Y. Cohen, R. de Cabo, C. Deng, S. Dimmeler, D. Gius, L.P. Guarante, S.L. Helfand, S. Imai, H. Itoh, T. Kadokawa, D. Koya, C. Leeuwenburgh, M. McBurney, Y. Nabeshima, C. Neri, P. Oberdoerffer, R.G. Pestell, B. Rogina, J. Sadoshima, V. Sartorelli, M. Serrano, D.A. Sinclair, **C. Steegborn**, M. Tatar, H.A. Tissenbaum, Q. Tong, K. Tsubota, A. Vaquero, E. Verdin (2010) Dietary restriction: standing up for sirtuins. *Science* 329, 1012-3.
- 4) M. Gertz M, **C. Steegborn** (2010) The lifespan-regulator p66Shc in mitochondria - redox enzyme or redox sensor? *Antioxid. Redox Signal.* 13, 1417-28.
- 3) M. Gertz, **C. Steegborn** (2010) Function and regulation of the mitochondrial Sirtuin isoform Sirt5 in mammalia. *Biochim. Biophys. Acta* 1804, 1658-65.
- 2) M. Kamenetsky, S. Middelhaufe, E.M. Bank, L.R. Levin, J. Buck, **C. Steegborn** (2006) Molecular details of cAMP generation in mammalian cells: a tale of two systems. *J. Mol. Biol.* 362, 623-639.
- 1) **C. Steegborn**, T. Clausen (2000) Toward specific inhibition of the pyridoxal-5'-phosphate dependent transsulfuration enzymes. *Recent Res. Dev. Biochem.* 2, 191-207.

Book chapters

- S. Moniot, W. You, **C. Steegborn** (2018) Structural and mechanistic insights in Sirtuin catalysis and pharmacological modulation. In “Introductory Review on Sirtuins in Biology and Disease” Eds. L. Guarante, R. Mostoslavsky, A. Kazantsev
- U.-M. Ohndorf, C. Schlicker, **C. Steegborn** (2009) Crystallographic studies on carbonic anhydrases from fungal pathogens for structure-assisted drug development. In “Drug Design of Zinc-Enzyme Inhibitors: Functional, Structural, and Disease Applications” Eds. C.T. Supuran, J.-Y. Winum.

Patent applications **C. Steegborn**, R. Huber, T. Clausen, A. Messerschmidt, *et al.* (1999) Inhibitoren der pflanzlichen Cystathionin gamma-Synthase. *German Patent Application 199 535 19.1*

J. Buck, L.R. Levin, L. Ramos-Espiritu, **C. Steegborn** (2017) Inhibitors of soluble adenylyl cyclase. *US patent application 62/328,806*