

Marija Buljan, PhD

Group Leader: Multi-omics for innovations in healthcare materials

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Education

2006 – 2010 University of Cambridge, UK, PhD thesis at the Wellcome Trust Sanger Institute
2001 – 2006 University of Zagreb, HR, Diploma engineer in Molecular Biology
1997 – 2001 Mathematical Gymnasium, Split, HR

Professional experience

Since 2020 Group Leader at Empa, Materials Science and Technology, CH
2015 – 2019 SNF SystemsX Fellow, ETH Zurich, CH
2013 – 2015 CellNetworks fellow, DKFZ Heidelberg, DE
2011 – 2013 Postdoctoral fellow, MRC LMB, UK
2014 – 2019 Expert adviser for European Commission, Brussels, BE

Invited talks

2019 Swiss Proteomics Meeting, Montreux, CH
2018 European Association for Cancer Research, Bergamo, IT
2018 From Functional Genomics to Systems Biology, EMBL
2018 Personalized Health conference, Zurich, CH
2017 Seattle Institute for Systems Biology, Seattle, USA
2016 European Bioinformatics Institute, Hinxton, UK
2016 Center for Molecular Medicine, Cologne, DE
2016 Anita Borg Birthday Anniversary, Google, Zurich, CH
2016 American Society for Mass Spectrometry, San Antonio, USA
2014 European Bioinformatics Institute, Hinxton, UK
2013 Society for Bioinformatics in Northern Europe, Torun, PL
2013 DyProt meeting, Dresden, DE
2012 Gordon Research Conference, Vermont, USA
2012 Copenhagen Bioscience Conference, Copenhagen, DK
2012 Universite Laval, Montreal, CA
2010 Functional Genomics and Disease, Dresden, DE

Awards

2009 Poster award at IMSB/ECCB conference on Computational Biology, SE
2005 Outstanding student award, Faculty of science, University of Zagreb
2001 Outstanding student award, Mathematical Gymnasium, Split, HR

Reviewing

I reviewed manuscripts for Cell, Cancer Cell, Cell Systems, Cell Reports, Nature Genetics, PLoS Comp Biol, Bioinformatics, Nature Communications and Molecular BioSystems

Publications (Google Scholar: Marija Buljan, Empa)

T Guo, A Luna, VN Rajapakse, CC Koh, Z Wu, W Liu, Y Sun, H Gao, MP Menden, C Xu, L Calzone, L Martignetti, C Auwerx, **M Buljan**, A Banaei-Esfahani, A Ori, M Iskar, L Gillet, R Bi, J Zhang, H Zhang, C Yu, Q Zhong, S Varma, U Schmitt, P Qiu, Q Zhang, Y Zhu, PJ Wild, M J Garnett, P Bork, M Beck, K Liu, J Saez-Rodriguez, F Elloumi, WC Reinhold, C Sander, Y Pommier, Aebersold R: Comparative analysis of mRNA and protein degradation in prostate tissues indicates high stability of proteins. *iScience* (2019)

Shao W, Guo T, Toussaint NC, Xue P, Wagner U, Li L, Charmpi K, Zhu Y, Wu J, **Buljan M**, Sun R, Rutishauser D, Hermanns T, Fankhauser CD, Poyet C, Ljubicic J, Rupp N, Rüschoff JH, Zhong Q, Beyer A, Ji J, Collins BC, Liu Y, Rättsch G, Wild PJ, R Aebersold R: Comparative analysis of mRNA and protein degradation in prostate tissues indicates high stability of proteins. *Nature communications* (2019)

Buljan M, Blattmann P, Aebersold R, M Boutros M: Systematic characterization of pan-cancer mutation clusters. *Molecular systems biology* (2018)

Glusman G, Rose PW, Prlić A, Dougherty J, Duarte JM, Hoffman AS, Barton GJ, Bendixen E, Bergquist T, Bock C, Brunk E, **Buljan M**, Burley SK, Cai B, Carter H, Gao JJ, Godzik A, Heuer M, Hicks M, Hrabe T, Karchin R, Koehler Lemman J, Lane L, Masica DL, Mooney SD, Moulton J, Omenn GS, Pearl F, Pejaver V, Reynolds SM, Rokem A, Schwede T, Song S, Tilgner H, Valasatava Y, Zhang Y, Deutsch EW: Mapping genetic variations to three-dimensional protein structures to enhance variant interpretation: a proposed framework. *Genome medicine* (2017)

Liu Y, Borel C, Li L, Müller T, Williams EG, Germain PL, **Buljan M**, Sajic T, Boersema PJ, Shao W, Faini M, Testa G, Beyer A, Antonarakis SE, Aebersold R: Systematic proteome and proteostasis profiling in human Trisomy 21 fibroblast cells. *Nature communications* (2017)

Latysheva NS, Oates ME, Maddox L, Flock F, Gough J, **Buljan M**, Weatheritt RJ, Babu MM: Molecular principles of gene fusion mediated rewiring of protein interaction networks in cancer. *Molecular cell* (2016)

R Van Der Lee, **M Buljan**, B Lang, RJ Weatheritt, GW Daughdrill, AK Dunker, M Fuxreiter, J Gough, J Gsponer, DT Jones, PM Kim, RW Kriwacki, CJ Oldfield, RV Pappu, P Tompa, VN Uversky, Peter E Wright, M Madan Babu, Classification of intrinsically disordered regions and proteins, *Chemical reviews* (2014)

DM Mitrea, CR Grace, **M Buljan**, MK Yun, NJ Pytel, J Satumba, A Nourse, CG Park, MM Babu, SW White, RW Kriwacki, Structural polymorphism in the N-terminal oligomerization domain of NPM1, *Proceedings of the National Academy of Sciences* (2014)

JK White, AK Gerdin, NA Karp, E Ryder, **M Buljan**, JN Bussell, J Salisbury, S Clare, NJ Ingham, C Podrini, R Houghton, J Estabel, JR Bottomley, DG Melvin, D Sunter, NC Adams, L Baker, C Barnes, R Beveridge, E Cambridge, D Carragher, P Chana, K Clarke, Y Hooks, N Igosheva, C Ismail, H Jackson, L Kane, R Lacey, DT Lafont, M Lucas, S Maguire, K McGill, RE McIntyre, S Messenger, L Mottram, L Mulderrig, S Pearson, HJ Protheroe, LA Roberson, G Salisbury, M Sanderson, D Sanger, C Shannon, PC Thompson, E Tuck, VE Vancollie, L Brackenbury, W Bushell, R Cook, P Dalvi, D Gleeson, B Habib, M Hardy, K Liakath-Ali, E Miklejewska, S Price, D Sethi, E Trenchard, D von Schiller, S Vyas, AP West, J Woodward, E Wynn, A Evans, D Gannon, M Griffiths, S Holroyd, V Iyer, C Kipp, M Lewis, W Li, D Oakley, D Richardson, D Smedley, C Agu, J Bryant, L Delaney, NI Gueorguieva, H Tharagonnet, AJ Townsend, D Biggs, E Brown, A Collinson, CE Dumeau, E Grau, S Harrison, James Harrison, Catherine E Ingle, Helen Kundi, Alla Madich, Danielle Mayhew, T Metcalf, S Newman, J Pass, L Pearson, H Reynolds, C Sinclair, H Wardle-Jones, M Woods, L Alexander, T Brown, F Flack, C Frost, N Griggs, S Hrniciarova, A Kirton, J McDermott, C Rogerson, G White, P Zielezinski, T DiTommaso, A Edwards, E Heath, MA Mahajan, B Yalcin, D Tannahill, DW Logan, DG MacArthur, J Flint, VB Mahajan, SH Tsang, I Smyth, FM Watt, WC Skarnes, G Dougan, DJ Adams, R Ramirez-Solis, A Bradley, KP Steel, Genome-wide generation and systematic phenotyping of knockout mice reveals new roles for many genes, *Cell* (2013)

M Buljan, G Chalancon, AK Dunker, A Bateman, S Balaji, M Fuxreiter, MM Babu, Alternative splicing of intrinsically disordered regions and rewiring of protein interactions, *Current opinion in structural biology* (2013)

M Buljan, G Chalancon, S Eustermann, GP Wagner, M Fuxreiter, A Bateman, MM Babu, Tissue-specific splicing of disordered segments that embed binding motifs rewires protein interaction networks, *Molecular cell* (2012)

M Buljan, A Frankish, A Bateman, Quantifying the mechanisms of domain gain in animal proteins, *Genome biology* (2010)

M Buljan, A Bateman, The evolution of protein domain families, *Biochemical Society Transactions* (2009)