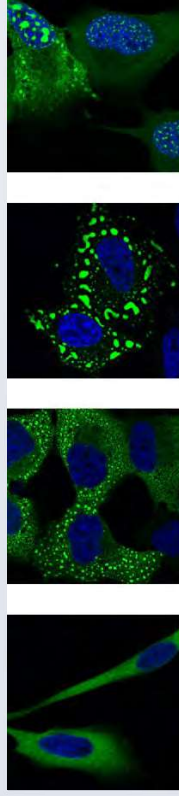




The 4th Cell Stress Society International workshop on

Small Heat Shock Proteins



The beauty and the complexity of the Small Heat Shock Proteins

Program

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4th CSSI workshop on Small Heat Shock Proteins

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Abstract Title:
Frameshift mutations in the Heat Shock Protein B8 share common pathogenic mechanisms and impair proteostasis

Names:

Barbara Tedesco^{*1,2}, Leen Vendredy^{*3}, Elias Adriaenssens³, Maria Cozzi¹, Bob Asselbergh^{4,5}, Valeria Crippa¹, Riccardo Cristofani¹, Paola Rusmini¹, Veronica Ferrari¹, Elena Casarotto¹, Marta Chierichetti¹, Francesco Mina¹, Paola Framaggione¹, Mariarita Galbiati¹, Margherita Piccollella¹, Jonathan Baets^{6,7}, Femke Baekle⁸, Riet De Rycke⁸, Vincent Mouly⁹, Tommaso Laurenzi¹, Ivano Eberini¹, Lan Weiss¹⁰, Virginia Kimonis¹⁰, Vincent Timmerman³, Angelo Poletti¹

Affiliations:

¹Dipartimento di Scienze Farmacologiche e Biomolecolari, Dipartimento di eccellenza 2018-2022, Università degli Studi di Milano, Milan, 20133, Italy; ²Unit of Medical Genetics and Neurogenetics, Fondazione IRCCS Istituto Neurologico Carlo Besta, Milan 20133, Italy; ³Peripheral Neuropathy Research Group, Department of Biomedical Sciences and Institute Born Bunge, University of Antwerp, Antwerpen, Belgium; ⁴Neuromics Support Facility, VIB Center for Molecular Neurology, VIB, Antwerp, Belgium; ⁵Neuromics Support Facility, Department of Biomedical Sciences, University of Antwerp, Antwerp, Belgium; ⁶Laboratory of Neuromuscular Pathology, Institute Born Bunge, and Translational Neurosciences, Faculty of Medicine, University of Antwerp, Antwerp, 2610, Belgium; ⁷Neuromuscular Reference Centre, Department of Neurology, Antwerp University Hospital, Antwerp, 2610, Belgium; ⁸Department of Biomedical Molecular Biology, and VIB Center for Inflammation Research, and VIB Bioimaging Core, Ghent University, 9000 Ghent, Belgium; ⁹Sorbonne Université, Inserm, Institut de Myologie, Centre de Recherche en Myologie, Paris, France; ¹⁰Dept. of Pediatrics, University of California, Irvine, United States.

Abstract:

The Heat Shock Protein B8 (HSPB8) is highly expressed in muscles where it favours the Chaperone Assisted Selective Autophagy (CASA) of damaged structural proteins. In neurons, HSPB8 enhances the clearance of misfolded substrates, such as those associated with motoneuron diseases. This is achieved by the interaction of HSPB8 with the co-chaperone BAG3, forming the CASA complex together with the HSP70 and the E3-ubiquitin ligase CHIP. When misfolded proteins are recognized by the chaperones of the CASA complex, they can be refolded or routed to aggregates for autophagy-mediated degradation. Frameshift mutations in *HSPB8* gene have been reported in neuromuscular diseases. These HSPB8 frameshift mutations cause the elongation of the HSPB8 protein product at the carboxy-terminus and a variable modification of the carboxy-terminal domain. Here, we show that the expression of the HSPB8 frameshift mutants associates with insolubility and aggregation propensity. The HSPB8 frameshift mutants retain the ability to take part in the CASA complex, determining the sequestration of the HSPB8 wildtype, BAG3, HSP70 and CHIP. As a result, misfolded and ubiquitinated substrates are entrapped in HSPB8-mutants aggregates together with autophagy receptors and CASA members. Notably, HSPB8 mutants aggregation is not driven by the CASA members, although BAG3 depletion affect aggregate number and size. Autophagy receptors do not cause HSPB8 mutants aggregation as well. Instead, we found that the elongated carboxy-terminus of HSPB8 mutants possesses intrinsic properties to aggregation. In summary, here we describe a gain of toxic function mechanism through which different HSPB8 frameshift mutations may cause neuromuscular diseases.