

February 18<sup>th</sup> 2017

Jill Trehella, Ph.D.  
Chair, wwPDB SAS VTF

Gaetano Montelione, Ph.D. and Micheal Nilges, Ph.D.  
Co-Chairs, wwPDB NMR VTF

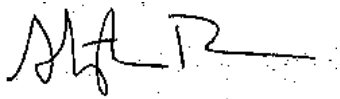
E-mails: [jill.trehella@sydney.edu.au](mailto:jill.trehella@sydney.edu.au), [guy@cabm.rutgers.edu](mailto:guy@cabm.rutgers.edu), [michael.nilges@pasteur.fr](mailto:michael.nilges@pasteur.fr)

Dear Jill, Guy, and Michael:

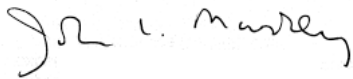
We write to request formally that you three and select members of the wwPDB Small-Angle Scattering Validation Task Force and the wwPDB NMR Validation Task Force meet as soon as is practicable to review the state-of-the-art and develop and promulgate recommendations regarding validation of integrative/hybrid methods NMR-SAS refined atomic coordinate structures. Your guidance is urgently required to ensure that integrative/hybrid NMR-SAS structures entering the Protein Data Bank archive are subject to rigorous community agreed validation procedures.

We are most grateful for your efforts on behalf of the Protein Data Bank archive and the wwPDB partnership, and stand ready to provide any help that we can in terms of meeting logistics, etc.

Yours faithfully,



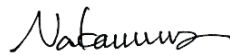
Stephen K. Burley, D.Phil., M.D.  
Distinguished Professor  
Rutgers, The State University of New Jersey  
Director, RCSB Protein Data Bank



John L. Markley, Ph.D.  
Steenbock Professor of Biomolecular  
Structure  
University of Wisconsin-Madison  
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Savmeer Velankar, Ph.D.  
Team Leader, EMBL-European  
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Haruki Nakamura, Ph.D.  
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