nature COMMUNICATIONS

DOI: 10.1038/ncomms7209

Erratum: Hydrogen bond rotations as a uniform structural tool for analyzing protein architecture

Robert C. Penner, Ebbe S. Andersen, Jens L. Jensen, Adriana K. Kantcheva, Maike Bublitz, Poul Nissen, Anton M.H. Rasmussen, Katrine L. Svane, Bjørk Hammer, Reza Rezazadegan, Niels Chr Nielsen, Jakob T. Nielsen & Jørgen E. Andersen

Nature Communications 5:5803 doi: 10.1038/ncomms6803 (2015); Published 17 Dec 2014; Updated 3 Feb 2015

The authors Robert C. Penner and Ebbe S. Andersen were incorrectly omitted from the list of corresponding authors in this Article. The correct information for correspondence is: 'Correspondence and requests for materials should be addressed to R.C.P. (email: rpenner@qgm.au.dk) or to E.S.A. (email: esa@inano.au.dk) or to J.E.A. (email: andersen@qgm.au.dk)'. The Article also contains errors in the labelling of Figs 1 and 2. In Fig. 1a, the labels RP_i, RP_j, $(RP_i)^{-1}$ and $R^H_{i,j}$ should read $\mathcal{R}_{P_i}$, $\mathcal{R}_{P_j}$, $(\mathcal{R}_{P_i})^{-1}$ and $\mathcal{R}^H_{i,j}$, respectively. In Fig. 2, labels R^B_{i-5}, R^B_{i-4}, R^B_{i-3}, R^B_{i-2}, R^B_{i-1}, R^B_i, R^B_{i+1}, R^B_{i+2}, R^B_{i+3}, R^B_{i+4}, R^B_{i+5} and R^B_{i+6} should read $\mathcal{R}^B_{i-5}$, $\mathcal{R}^B_{i-4}$, $\mathcal{R}^B_{i-3}$, $\mathcal{R}^B_{i-2}$, $\mathcal{R}^B_{i-1}$, $\mathcal{R}^B_i$, $\mathcal{R}^B_{i+1}$, $\mathcal{R}^B_{i+2}$, $\mathcal{R}^B_{i+3}$, $\mathcal{R}^B_{i+4}$, $\mathcal{R}^B_{i+5}$ and $\mathcal{R}^B_{i+6}$, respectively. The correct versions of these figures follow.

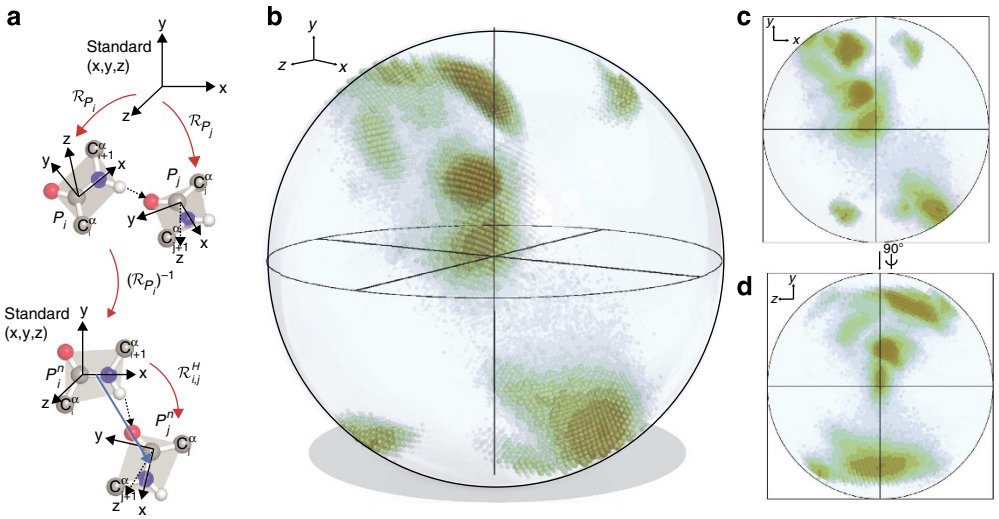

Figure 1

NATURE COMMUNICATIONS | DOI: 10.1038/ncomms7209

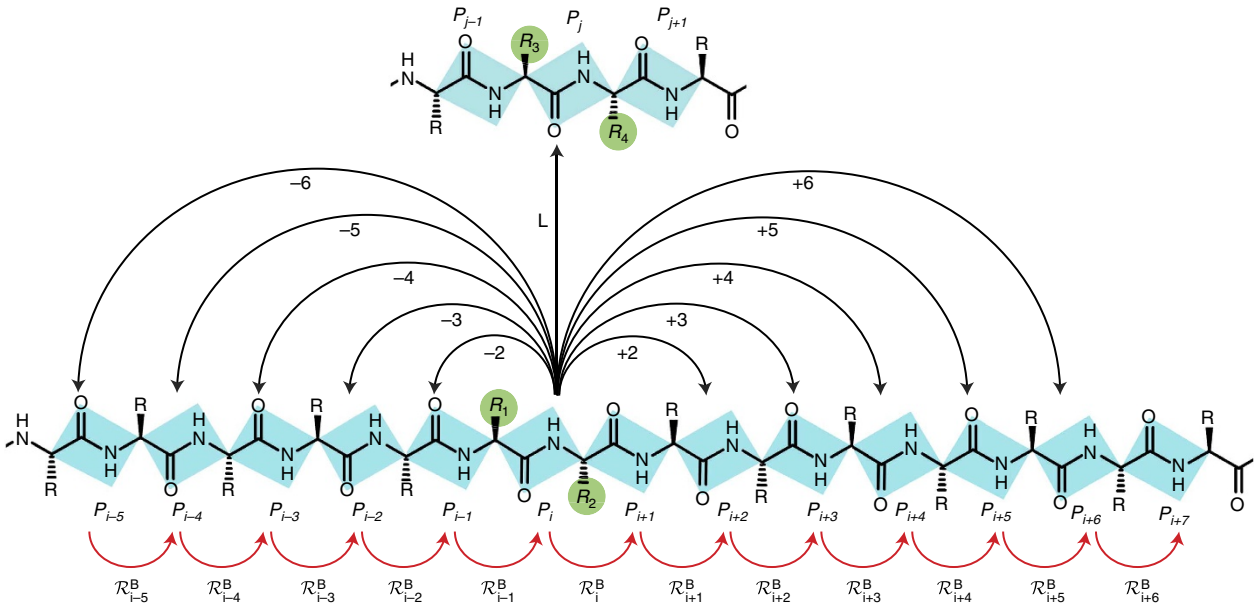

Figure 2