

**ERRATUM**

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**BEAMS (BEAdS Modelling System): a set of computer programs  
for the generation, the visualization and the computation of the  
hydrodynamic and conformational properties of bead models of proteins**

Eur Biophys J (1997) 25: 373–384

The above article unfortunately contained three errors:

1. The e-mail address on the title page should read  
rocco@vega.cba.unige.it
2. The Internet site in the Acknowledgements should be  
www2.cba.unige.it/Macro/Hydro
3. The correct Equation (25) should read

$$R_E = \left[ \frac{3M[\eta]}{10\pi N_A} \right]^{\frac{1}{3}}$$

We apologize for these mistakes.