



# Author Correction: Quantitative fragmentomics allow affinity mapping of interactomes

Correction to: *Nature Communications*  
<https://doi.org/10.1038/s41467-022-33018-0>,  
 published online 17 September 2022

<https://doi.org/10.1038/s41467-022-35177-6>

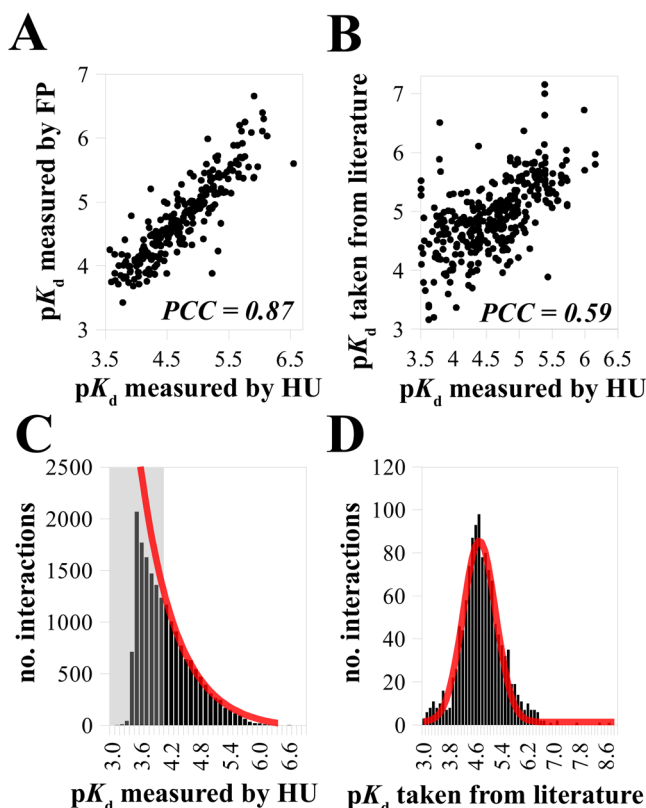
Published online: 07 December 2022



Gergo Gogl , Boglarka Zambo , Camille Kostmann, Alexandra Cousido-Siah, Bastien Morlet , Fabien Durbesson, Luc Negroni , Pascal Eberling, Pau Jané, Yves Nominé , Andras Zeke , Søren Østergaard , Élodie Monsellier, Renaud Vincentelli & Gilles Travé

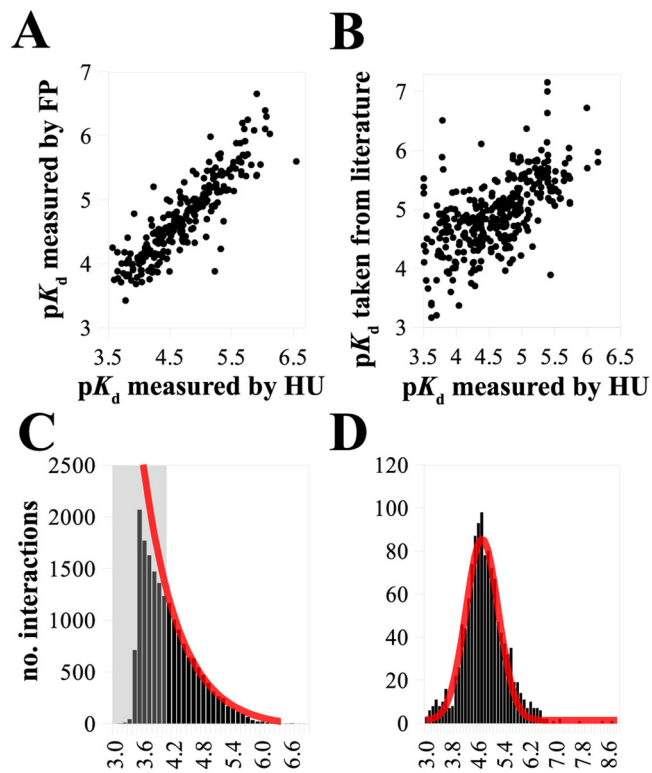
The original version of this article contained errors in Fig. 2, Fig. 4 and Fig. 5.

In the original version of Fig. 2, PCC values were missing from panels a, b and axes labels were missing from panels c, d. The correct version of Fig. 2 is:



# Corrections & amendments

which replaces the previous incorrect version:

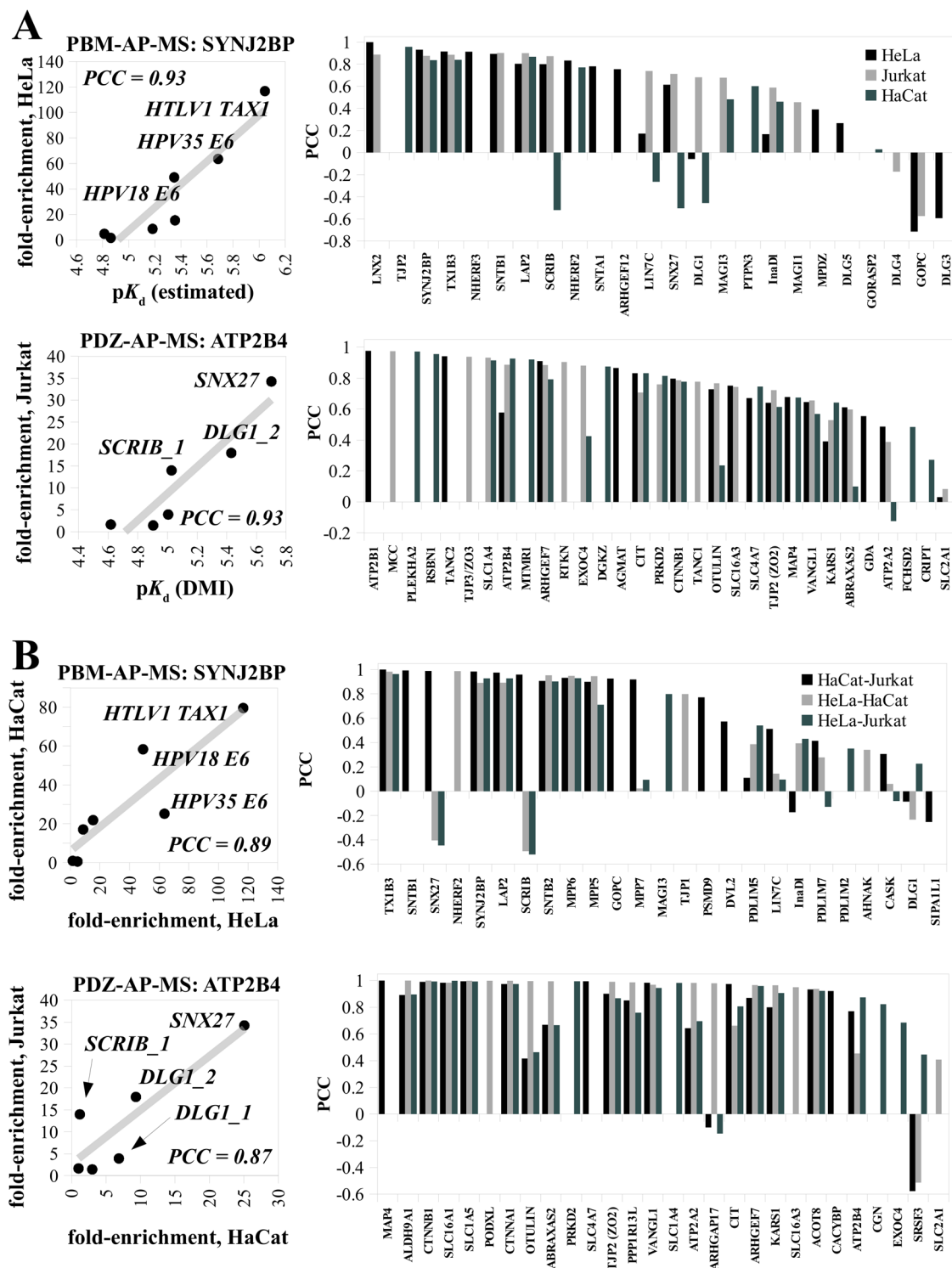






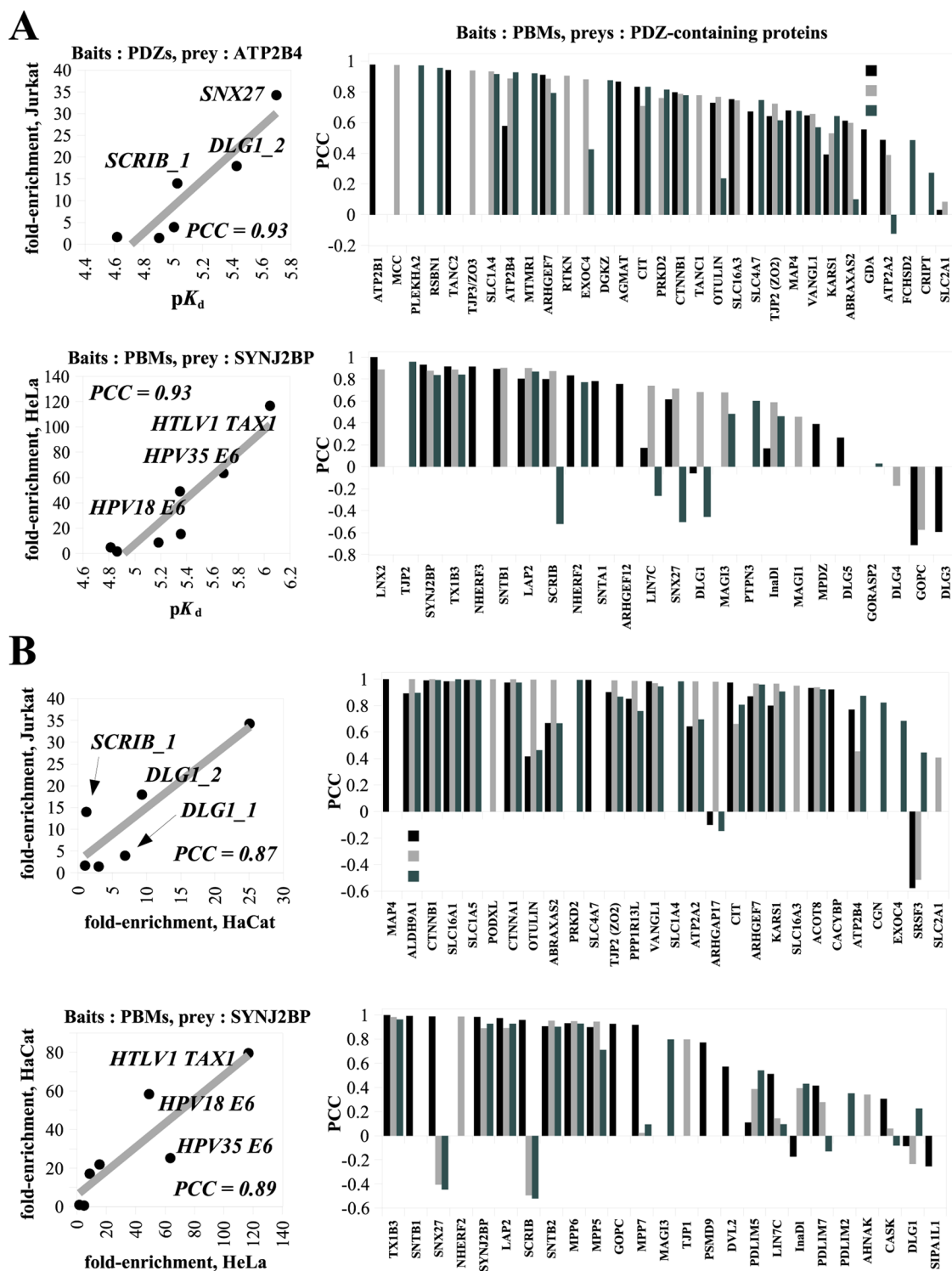
# Corrections & amendments

In the original version of Fig. 5, several legends as well as plot and axes labels were missing or incomplete and the order of panels did not match the descriptions in the figure caption. The correct version of Fig. 5 is:



# Corrections & amendments

which replaces the previous incorrect version:



This has been corrected in both the PDF and HTML versions of the Article.

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# Corrections & amendments

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