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Hakim, S. S.; Olsson, M. H. M.; Sorensen, H. O.; Bovet, Nicolas Emile; Bohr, J.; Feidenhans'l, Robert Krarup; Stipp, Susan Louise Svane

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Publisher Correction: Interactions of the Calcite \{10.4\} Surface with Organic Compounds: Structure and Behaviour at Mineral–Organic Interfaces

S. S. Hakim\textsuperscript{1,} M. H. M. Olsson\textsuperscript{1,} H. O. Sørensen\textsuperscript{1,} N. Bovet\textsuperscript{1,} J. Bohr\textsuperscript{2,} R. Feidenhans'li\textsuperscript{3} \& S. L. S. Stipp\textsuperscript{1}

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This Article contained an error in the legend of Table 1 where,

“...For each interactors, the interacting TANC protein, the detection method and the binding region (experimentally validated) are here listed. Y2H: Yeast two hybrid; Co-IP: Co-immunoprecipitation; SPR: Surface plasmon resonance HTS: High-Throughput System; AC: Affinity Capture; PL: Proximity Label; MS: Mass spectrometry; CLIP: Cross-Linking ImmunoPrecipitation SF-TAP/MS: systematic tandem affinity purifications coupled to mass spectrometry; LIG\_PDZ\_Class\_1: PDZ-binding motif; LIG\_EVH1\_1: Proline-rich motif binding to signal transduction class I EVH1 domains; DEG\_SCF\_TRCP1: SCF-betaTrCP1 complex target site; MOD\_LATS\_1: phosphorylation motif recognised by the LATS kinases; DOC\_PP1\_RVXF\_1: PP1 docking motif; LIG\_14-3-3\_2: phospho-motif mediating the interaction with 14-3-3 proteins; LIG\_Actin\_WH2\_2: Actin-binding motif; TRG\_NES\_CRM1\_1: Nuclear Export Signal.”

now reads:

“The tabulated parameters are obtained from fitting the reflectivity data with box models. For models with more than one layer, the layer closest to the calcite surface is indexed 1 (e.g. Methanol-1). The density for calcite (2.71 g/cm\textsuperscript{3}) and helium (0.03 g/cm\textsuperscript{3}) were kept fixed during the fitting procedure, and their thicknesses were infinite. Error estimates were obtained for every parameter by varying the parameter until the \(\chi^2\) based R-value is changed by 5\% [70, 71]. The method does not reveal correlations between the parameters. On the right side, theoretical bulk density and the molecular length of the molecules are given for comparison.”

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\textsuperscript{1}Nano-Science Center, Department of Chemistry, University of Copenhagen, Universitetsparken 5, 2100, Copenhagen, Denmark. \textsuperscript{2}DTU Nanotech, Department of Micro- and Nanotechnology, Technical University of Denmark, Ørsted Plads, 2800 Kgs. Lyngby, Denmark. \textsuperscript{3}Niels Bohr Institute, University of Copenhagen, Blegdamsvej 17, 2100, Copenhagen, Denmark. Correspondence and requests for materials should be addressed to S.S.H. (email: hakim@nano.ku.dk) or H.O.S. (email: osholm@nano.ku.dk)