Correction to ‘The basic keratin 10-binding domain of the virulence-associated pneumococcal serine-rich protein PsrP adopts a novel MSCRAMM fold’

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The residue numbers in the associated PDB files 3ZGH and 3ZGI were corrected during the PDB deposition process to have the same numbers as given in the associated Uniprot entry Q97P71. Unfortunately, we missed to update the residue numbers in the figures and text of the manuscript. The numbers in the manuscript should have an offset of −1. However, the first and last residue numbers given for the protein constructs BR187–385 and STII-BR187–385 remain valid.