

# Brock Schuman

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6517523/publications.pdf>

Version: 2024-02-01

9  
papers

109  
citations

1684188  
5  
h-index

1588992  
8  
g-index

10  
all docs

10  
docs citations

10  
times ranked

155  
citing authors

| # | ARTICLE   | IF  | CITATIONS |
|---|---|-----|-----------|
| 1 | Conserved residues Arg188 and Asp302 are critical for active site organization and catalysis in human ABO(H) blood group A and B glycosyltransferases. <i>Glycobiology</i> , 2018, 28, 624-636.   | 2.5 | 9         |
| 2 | Glycosyltransfer in mutants of putative catalytic residue Glu303 of the human ABO(H) A and B blood group glycosyltransferases GTA and GTB proceeds through a labile active site. <i>Glycobiology</i> , 2017, 27, 370-380.                                       | 2.5 | 5         |
| 3 | High-resolution crystal structures and STD NMR mapping of human ABO(H) blood group glycosyltransferases in complex with trisaccharide reaction products suggest a molecular basis for product release. <i>Glycobiology</i> , 2017, 27, 966-977.                 | 2.5 | 3         |
| 4 | pH-induced conformational changes in human ABO(H) blood group glycosyltransferases confirm the importance of electrostatic interactions in the formation of the semi-closed state. <i>Glycobiology</i> , 2014, 24, 237-246.                                     | 2.5 | 5         |
| 5 | Geometric Attributes of Retaining Glycosyltransferase Enzymes Favor an Orthogonal Mechanism. <i>PLoS ONE</i> , 2013, 8, e71077.   | 2.5 | 27        |
| 6 | Sequence-dependent effects of cryoprotectants on the active sites of the human ABO(H) blood group A and B glycosyltransferases. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2012, 68, 268-276.  | 2.5 | 10        |
| 7 | Cysteine-to-Serine Mutants Dramatically Reorder the Active Site of Human ABO(H) Blood Group B Glycosyltransferase without Affecting Activity: Structural Insights into Cooperative Substrate Binding. <i>Journal of Molecular Biology</i> , 2010, 402, 399-411. | 4.2 | 11        |
| 8 | The effect of heavy atoms on the conformation of the active-site polypeptide loop in human ABO(H) blood-group glycosyltransferase B. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2007, 63, 860-865.                                   | 2.5 | 12        |
| 9 | Glycosyltransferase Structure and Function. , 2006, , 217-257.  |     | 27        |