

**Hyun Woo “John” Kim**  
**Curriculum Vitae**  
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**Education**

2017 – Present      Ph.D. training in Biochemistry and Molecular Biology, University of Georgia  
2013 – 2016        B.S. in Biochemistry and Molecular Biology, *cum laude*, University of Georgia  
2010 – 2012        B.A. in Music Education and Trombone Performance, Columbus State University  
(did not finish)

**Research Experience**

2017 – Present      Graduate research assistant under Dr. Christopher M. West  
2016 – 2017        Lab technician under Dr. Christopher M. West  
2015 – 2016        Undergraduate research assistant under Dr. Zachary A. Wood

**Publications**

West C.M. & **Kim H.W.** (2019) Nucleocytoplasmic O-Glycosylation in Protists. *Curr. Opin. Struc. Biol.* **56**:204-212  
Kadirvelraj R., Yang J.Y., **Kim H.W.**, Sanders, J.S., Moremen, K.W., Wood, Z.A. (2019) Human poly-N-acetyl-lactosamine synthase structure demonstrates a modular assembly of catalytic subsites for GT-A glycosyltransferases. PNAS. (submitted)  
Rahman K., Mandalasi M., Zhao P., Sheik M.O., Taujale R., **Kim H.W.**, van der Wel H., Matta K., Kannan N., Glushka J.N., West C.M. (2017) Characterization of a cytoplasmic glucosyltransferase that extends the core trisaccharide of the *Toxoplasma* Skp1 E3 ubiquitin ligase subunit. *J.Biol.Chem.* **292**:18644-18659.

**Manuscripts submitted or in preparation**

Mandalasi, M., **Kim, H.W.**, Rahman, K., Zhao, P., Daniel, N., Sheikh, M.O., van der Wel, H., Thieker, D., Ichikawa, T.H., Glushka, J.N., Wells, L., Wood, Z.A., West, C.M. (2019) A glycogenin homolog controls *Toxoplasma gondii* growth via glycosylation of an E3 ubiquitin ligase. (manuscript in preparation)  
Shrestha S., Katiyar S., Sans-Rodriguez C.E., Kemppinen N.R., **Kim H.W.**, Kadirvelraj R., Panagos C., Keyhaninejad N., Chopra P., Byrne D.P., Boons G.J., Knapp E.V., Evers P.A., Edison A.S., Wood Z.A., Kannan N. (2019) Identification of a novel redox-active switch in Fructosamine-3-Kinases expands the regulatory repertoire of the protein kinase super-family. *Sci. Signal.* (submitted)  
Kadirvelraj R., Yang J.Y., **Kim H.W.**, Sanders, J.S., Moremen, K.W., Wood, Z.A. (2019) Human poly-N-acetyl-lactosamine synthase structure demonstrates a modular assembly of catalytic subsites for GT-A glycosyltransferases. PNAS. (submitted)

**Awards and Presentations**

2019    The Protein Society Annual Symposium, Seattle (planned)  
      -“*Structural Insights into the Evolution of the CFAZy GT8 Glycosyltransferase Glycogenin*”  
      -“*Cis-acting Glycan Drives Protein-Protein Interactions of Skp1 in Dictyostelium and Toxoplasma*”  
2018    Biochemistry departmental retreat – poster presentation  
      -“*How does glycosylation modulate Skp1 organization?*”  
2018    Society for Glycobiology - travel grant  
2018    Society for Glycobiology annual conference, New Orleans – poster presentation  
      -“*How does glycosylation modulate Skp1 organization?*”  
2018    Biophysics workshop at NIH, Bethesda – poster presentation  
      -“*Glycosylation modulates Skp1 self-association and its interaction with F-box proteins*”  
2016    UGA Biochemistry Undergraduate Symposium – 1<sup>st</sup> place poster presentation  
      -“*The Crystal Structure and Analysis of Human B3GNT2: a Major Poly lactosamine Synthase*”

## **Memberships**

2019- The Protein Society  
2018- Society for Glycobiology  
2018- American Association for the Advancement of Science

## **Ongoing Projects**

**Kim H.W.**, Eletsky A. van Der Wel H., Prestegard J.H., West C.M. Solution structure of Skp1 reveals dimer interface that occupies F-box binding site.

**Kim H.W.**, Mandalasi M., Wood Z.A., West C.M. *Toxoplasma* Skp1 glycosyltransferase Gat1 exhibits non-catalytic function through protein-protein interaction.

**Kim H.W.**, van Der Wel H., Wood Z.A., West C.M. Skp1-GlcNAc transferase is distantly related to mucin-type GalNAc transferase from GT27 family.

## **PDB depositions**

*Skp1  $\alpha$ -D-galactosyltransferase (Gat1)*

6MW5 - Pt heavy atom derivative

6MW8 - bound Mn<sup>2+</sup> ion and UDP

*$\beta$ 1,3-N-acetylglucosaminyltransferase 2 (B3GNT2)*

6OLB - Selenomethionine derivative

6OLC - bound Mg<sup>2+</sup> ion and UDP

6OLH - bound Mg<sup>2+</sup> ion, UDP, and acceptor (LNnT)

*T-synthase chaperone (Cosmc)*

6OA2 - Cs heavy atom derivative

6OA4 – native protein

6OA5 – truncated protein

## **Scholastic performance**

Undergraduate GPA (UGA) - 3.62

Graduate GPA (UGA) - 3.89

GRE scores:

Verbal reasoning 151 (52%)

Quantitative reasoning 156 (62%)

Analytical writing 2.5 (8%)

## **Graduate courses taken**

*Fall 2017*

8000 – Research Techniques in ILS

8010 – Professional Development for ILS Students

8020 – Critical Reading of Primary Science Literature

8550 – Responsible Conduct in Research

*Spring 2018*

8060 – Student seminar

8070 – Research Discussion

8080 – Journal Club

9000 – Doctoral Research

8113 – Advanced Genetics Cell Biology BCMB

8114 – Advanced Genetics Cell Biology BCMB

8330 – Molecular Modeling

*Summer 2018*

8070 – Research Discussion

8080 – Journal Club

9000 – Doctoral Research

*Fall 2018*

8060 – Student seminar

8070 – Research Discussion

8080 – Journal Club

9000 – Doctoral Research

8213 – Advanced Genetic Cell Biology BCMB

8214 – Advanced Genetic Cell Biology BCMB

*Spring 2019*

8060 – Student seminar

8070 – Research Discussion

8080 – Journal Club

9000 – Doctoral Research

8110 – Protein Structure/Function

8990 – Grant Writing

*Summer 2019*

8070 – Research Discussion

8080 – Journal Club

9000 – Doctoral Research