

## **SlgN affiliated publications by Wei WU**

*Last updated: 30 November 2022*

### **2022**

van Bergen W, Hevler JF, Wu W, Baggelaar MP, Heck AJR. SITE-SPECIFIC ACTIVITY-BASED PROTEIN PROFILING USING PHOSPHONATE HANDLES. *Mol Cell Proteomics*. 2022 Nov 23:100455.

Beumer J, Bauzá-Martinez J, Veth TS, Geurts V, Boot C, Gilliam-Vigh H, Poulsen SS, Knop FK, Wu W, Clevers H. Mapping prohormone processing by proteases in human enteroendocrine cells using genetically engineered organoid models. *Proc Natl Acad Sci U S A*. 2022 Nov 16;119(46):e2212057119.

He GW, Lin L, DeMartino J, Zheng X, Staliarova N, Dayton T, Begthel H, van de Wetering WJ, Bodewes E, van Zon J, Tans S, Lopez-Iglesias C, Peters PJ, Wu W, Kotlarz D, Klein C, Margaritis T, Holstege F, Clevers H. Optimized human intestinal organoid model reveals interleukin-22-dependency of paneth cell formation. *Cell Stem Cell*. 2022 Sep 1;29(9):1333-1345.e6.

Bauzá-Martinez J, Armony G, Pronker MF, Wu W. Characterization of protein complexes in extracellular vesicles by intact extracellular vesicle crosslinking mass spectrometry (iEVXL). *J Extracell Vesicles*. 2022 Jul;11(8):e12245.