

Merck Research Symposium Poster Session

Due to work being proprietary, I cannot share the contents of the poster I created.

I was invited and created a poster for the annual Merck Research Symposium. This was related to my main project of the Automation of the Liquid Chromatography-Mass Spectrometry (LC-MS) Workflow Using a Run Sequence Wizard. For those that are curious about the project, information can be found [here](#). The poster received quite a lot of attention. I was only one of three bioinformaticians out of nearly 200 undergraduate, graduate and post-doc scientists. The other two bioinformaticians were doing their graduate studies, meaning that **I was the only undergraduate bioinformatician invited to the symposium.**

My presentation was viewed by approximately 200 people of which half were scientists, directors and vice presidents of the research divisions. The other half included my peers that were presenting in the symposium.

I took questions from around 50 people in varying group sizes about what bioinformatics is, why my project was important and what my future was. I was then invited to give a talk to Merck Research Laboratories (MRL), Pharmacokinetics, Pharmacodynamics and Drug Metabolism (PPDM), Tissue Distribution and Metabolite Identification (TD/ID) group.

Information about the talk given can be found [here](#).