



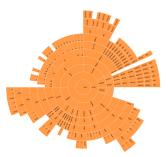


October 2014

Call 4 Students!

The SEMS group at the **Department of Systems Biology and Bioinformatics** aims at improving the management of computational models. To boost our studies we seek motivated students to extend our team.

Right now we need a tool to **visualise hierarchical contents** (such as XML trees). There is already software available, e.g. TreeViz¹. While it gives you an idea of what we are looking for, the tool lacks some essential features necessary to meet our requirements. We want you to either extend the tool or to develop an alternative which fits our needs.



A suitable candidate is **highly motivated** to join our junior research group. You need to have at least some basic programming skills, preferably Java and Javascript. We would be happy to integrate you in our team.

Are you the right person for the job?

Do not hesitate to apply for a **HiWi position**! Simply email us your CV and tell us a bit about yourself. We will then invite you for a short interview to align our visions. This work could also be embedded in a **Bachelor or Master thesis**. In case there are open questions feel free to drop us a line!



Contact

martin.scharm@uni-rostock.de
https://sems.uni-rostock.de/jobs

¹http://hypertree.sourceforge.net/