

WE ARE LOOKING FOR TWO BRILLIANT PROTEOMICS SCIENTISTS

We developed and launched the Evosep One chromatography system with plans to dramatically expand the possibilities in the field of clinical LC-MS. Those plans are unfolding rather well and we are now seeking new colleagues to achieve even more.

So we wish to expand the team at our Odense office with two Scientists, who can help develop new applications, protocols, and methods.

Responsibilities

- Develop and test new separation methods and workflows for LC-MS.
- Support and conduct in-house research and development projects in general.
- Create and maintain application documentation, protocols and scientific literature.
- Represent Evosep in a few longer-term scientific collaborations.
- Help fight the fires that inevitably happen when you least expect them.

Qualifications

- B.Sc., M.Sc., or Ph.D. in Analytical Sciences, Biochemistry or related fields.
- 3+ years of hands-on experience with LC-MS instrumentation.
- Experience with automation and robotics will be considered beneficial.
- Excellent communication and “people” skills are essential. This goes for both writing and presenting skills, as well as listening and interaction skills in general.
- Moderate travel will be required within Europe, as well as occasional overseas trips.

At Evosep, we are a small but very dedicated team with a long history in the nanoLC and proteomics community. We value an informal and constructive tone both internally and when working with our users. We are not perfect (and neither are you) but we enjoy the quest for high performance (and so should you).

The company is headquartered in Odense, Denmark, but our customers are evenly split between Europe and North America.

If you are interested in learning more or want to submit an application, please contact Ole Vorm at ov@evosep.com or phone +45 26 33 23 23.

Evosep aims to improve quality of life and patient care by radically innovating protein based clinical diagnostics. We will make sample preparation and separation for MS analysis 10 times faster and 100 times more robust in order to enable truly large cohort studies and provide the foundation for precision medicine.