

Containerization of microbial proteomics data processing pipelines

Timothy Bergström

March 11, 2019

tib@kth.se

950502-1312

KTH: Royal institute of technology

Abstract

Hello there

Sammanfattning

Hej på dig

Contents

1	Introduction	1
2	Background	2

Nomenclature

GUI Graphical User Interface

HPC High Performance Computing

MS Mass Spectrometry

1 Introduction

2 Background