BIOINF 2019/20, winter semester, 1st exam

* Synonymous mutations
* Protein structure
* Dotplot
* Distance calculation using algorithm
* How can we see that a phylogenetic study is robust
* How can we predict a protein structure
* Darwin theory on a molecular sequence level
* EST’s – what it is and its uses (?)
* Shotgun in metagenomics
* Applications of bioinformatics