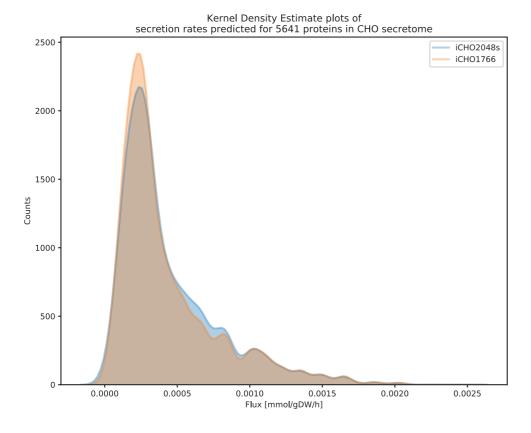
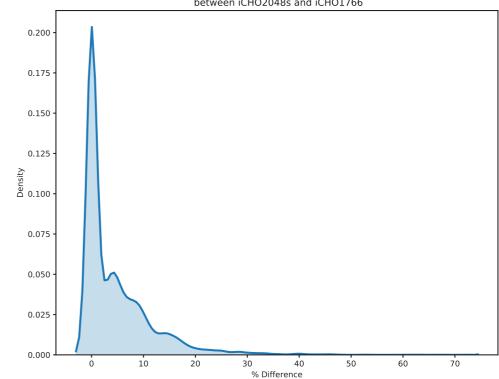
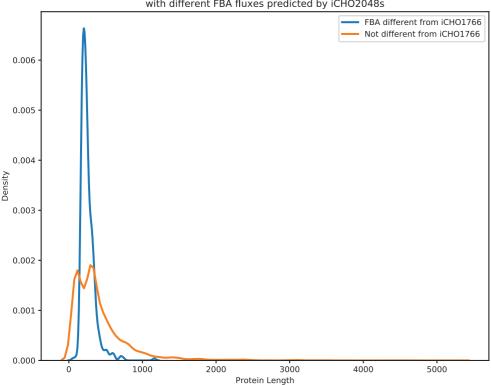
A

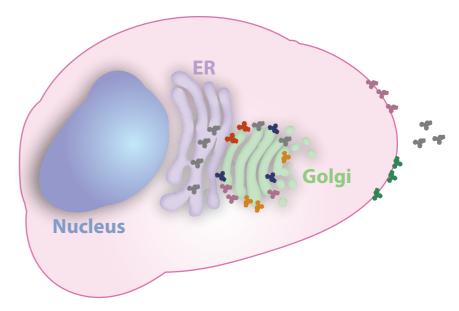






Distribution of Protein length across proteins with different FBA fluxes predicted by iCHO2048s





- Protein with no modifications
- O-linked glycosylation only
- GPI-anchor only
- Transmembrane domain only
- Disulfide bonds and Transmembrane domain
- O-linked glycosylation and Transmembrane domain