Possible interesting articles

## Elisa

Modeling gene-wise dependencies improves the identification of drug response biomarkers in cancer studies.

* <https://europepmc.org/article/MED/28082455>

## Max

Chen et al. - Predicting HLA class II antigen presentation through integrated deep learning (2019)

De Groot et al. - Large-scale image-based profiling of single-cell phenotypes in arrayed CRISPR-Cas9 gene perturbation screens (2018)