**Supplementary 2\_ Links to repositories**

1. Link to Figshare repository:

<https://figshare.com/projects/Comparing_feature_selection_and_machine_learning_approaches_for_predicting_CYP2D6_methylation_from_genetic_variations/178203>

1. Scripts for genotype imputation

1 <https://bitbucket.org/sics-sb/gusto-imputation.git>

2 <https://github.com/natacha-beck/imputePrepSanger>