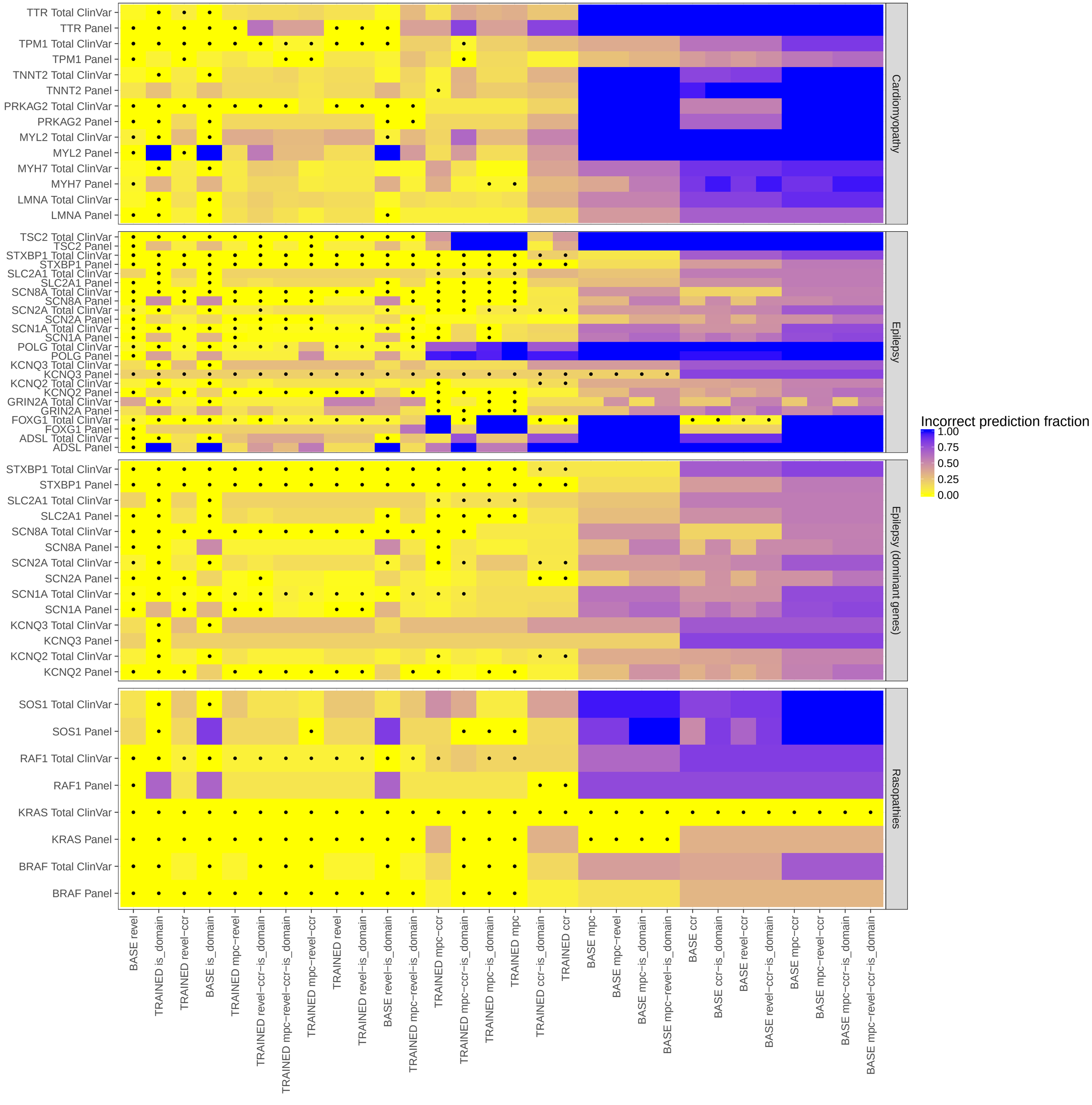
TTR Total ClinVar
TTR Panel
TPM1 Total ClinVar
TPM1 Panel
TNNT2 Total ClinVar
TNNT2 Panel
PRKAG2 Total ClinVar
PRKAG2 Panel
MYL2 Total ClinVar
MYL2 Panel
MYH7 Total ClinVar
MYH7 Panel
LMNA Total ClinVar
LMNA Panel
Cardiomyopathy
TSC2 Total ClinVar
TSC2 Panel
STXBP1 Total ClinVar
STXBP1 Panel
SLC2A1 Total ClinVar
SLC2A1 Panel
SCN8A Total ClinVar
SCN8A Panel
SCN2A Total ClinVar
SCN2A Panel
SCN1A Total ClinVar
SCN1A Panel
POLG Total ClinVar
POLG Panel
KCNQ3 Total ClinVar
KCNQ3 Panel
KCNQ2 Total ClinVar
KCNQ2 Panel
GRIN2A Total ClinVar
GRIN2A Panel
FOXG1 Total ClinVar
FOXG1 Panel
ADSL Total ClinVar
ADSL Panel
Epilepsy
STXBP1 Total ClinVar
STXBP1 Panel
SLC2A1 Total ClinVar
SLC2A1 Panel
SCN8A Total ClinVar
SCN8A Panel
SCN2A Total ClinVar
SCN2A Panel
SCN1A Total ClinVar
SCN1A Panel
KCNQ3 Total ClinVar
KCNQ3 Panel
KCNQ2 Total ClinVar
KCNQ2 Panel
Epilepsy (dominant genes)
SOS1 Total ClinVar
SOS1 Panel
RAF1 Total ClinVar
RAF1 Panel
KRAS Total ClinVar
KRAS Panel
BRAF Total ClinVar
BRAF Panel
Rasopathies
BASE revel
TRAINED is_domain
TRAINED revel−ccr
BASE is_domain
TRAINED mpc−revel
TRAINED revel−ccr−is_domain
TRAINED mpc−revel−ccr−is_domain
TRAINED mpc−revel−ccr
TRAINED revel
TRAINED revel−is_domain
BASE revel−is_domain
TRAINED mpc−revel−is_domain
TRAINED mpc−ccr
TRAINED mpc−ccr−is_domain
TRAINED mpc−is_domain
TRAINED mpc
TRAINED ccr−is_domain
TRAINED ccr
BASE mpc
BASE mpc−revel
BASE mpc−is_domain
BASE mpc−revel−is_domain
BASE ccr
BASE ccr−is_domain
BASE revel−ccr
BASE revel−ccr−is_domain
BASE mpc−ccr
BASE mpc−revel−ccr
BASE mpc−ccr−is_domain
BASE mpc−revel−ccr−is_domain
Incorrect prediction fraction
1.00
0.75
0.50
0.25
0.00