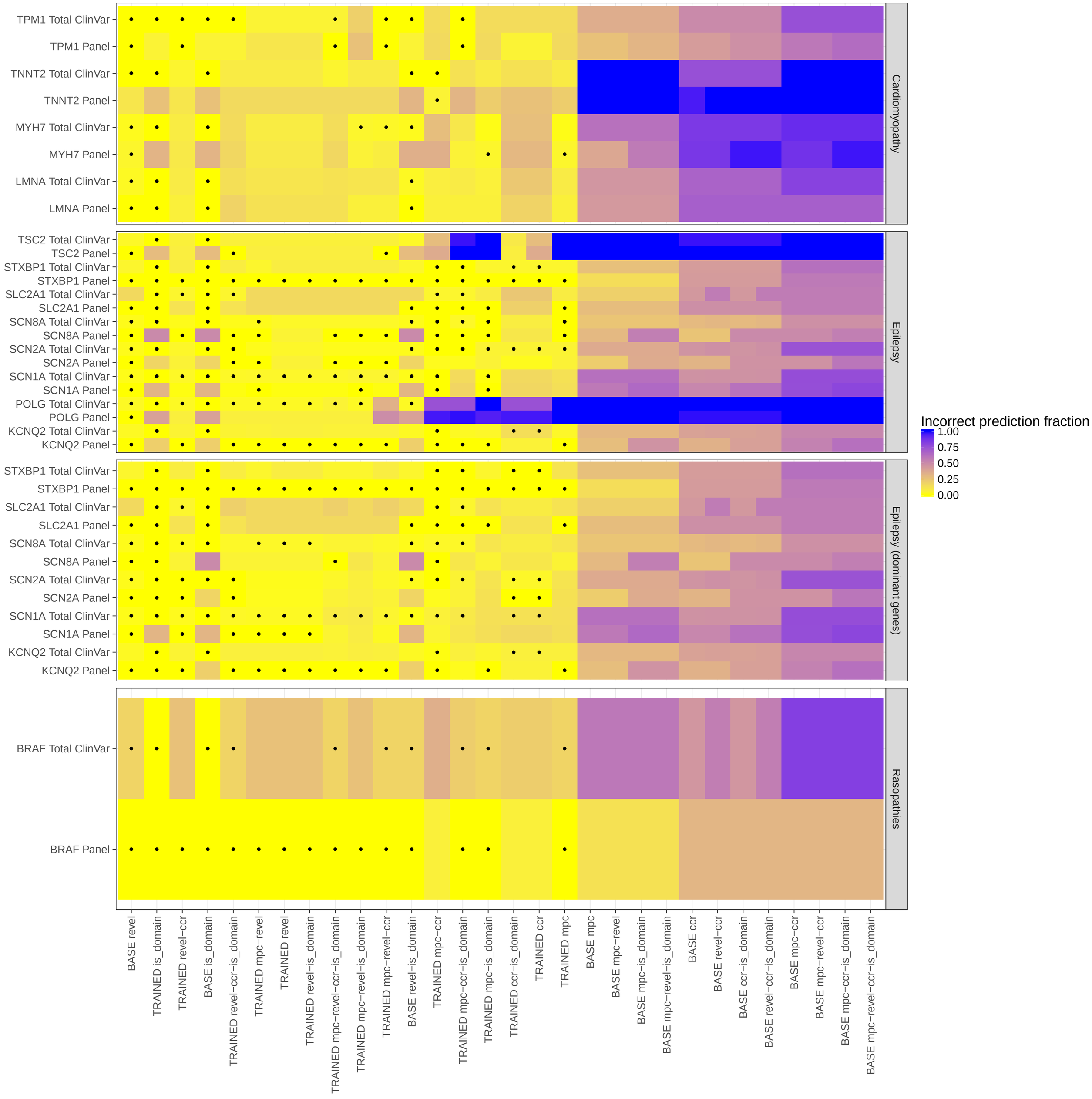
TPM1 Total ClinVar
TPM1 Panel
TNNT2 Total ClinVar
TNNT2 Panel
MYH7 Total ClinVar
MYH7 Panel
LMNA Total ClinVar
LMNA Panel
Cardiomyopathy
TSC2 Total ClinVar
TSC2 Panel
STXBP1 Total ClinVar
STXBP1 Panel
SLC2A1 Total ClinVar
SLC2A1 Panel
SCN8A Total ClinVar
SCN8A Panel
SCN2A Total ClinVar
SCN2A Panel
SCN1A Total ClinVar
SCN1A Panel
POLG Total ClinVar
POLG Panel
KCNQ2 Total ClinVar
KCNQ2 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
KCNQ2 Total ClinVar
KCNQ2 Panel
Epilepsy (dominant genes)
BRAF Total ClinVar
BRAF Panel
Rasopathies
BASE revel
TRAINED is_domain
TRAINED revel−ccr
BASE is_domain
TRAINED revel−ccr−is_domain
TRAINED mpc−revel
TRAINED revel
TRAINED revel−is_domain
TRAINED mpc−revel−ccr−is_domain
TRAINED mpc−revel−is_domain
TRAINED mpc−revel−ccr
BASE revel−is_domain
TRAINED mpc−ccr
TRAINED mpc−ccr−is_domain
TRAINED mpc−is_domain
TRAINED ccr−is_domain
TRAINED ccr
TRAINED mpc
BASE mpc
BASE mpc−revel
BASE mpc−is_domain
BASE mpc−revel−is_domain
BASE ccr
BASE revel−ccr
BASE ccr−is_domain
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