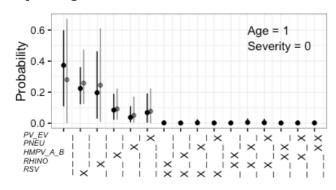
## Compare with nplcm using Kenya site data

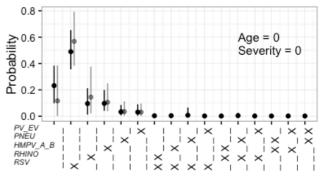
WBPCR is not used. TPR & FPR are not stratified. No regression on FPR.

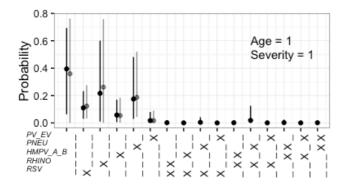
## 1. Multinomial non-regression model with 32 candidate pathogens.

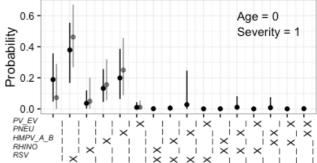
Eti.Name	pEti	Eti.Name	pEti	Eti.Name	pEti	Eti.Name	pEti
RSV	0.3491	PARA_3	0.0330	ADENO	0.0064	ENTRB	0.0017
RHINO	0. 1528	NoA	0.0261	PARA_2	0.0058	CMV	0.0015
HMPV_A_B	0. 1016	PARA_1	0.0261	FUNGI	0.0052	COR	0.0014
PNEU	0.0754	FLU_A	0.0220	MCAT	0.0040	NMEN	0.0013
PV_EV	0.0370	PCP	0.0211	OTHSTR	0.0039	HBOV	0.0011
HINF	0.0368	PARA_4	0.0202	M_PNEU	0.0033	FLU_C	0.0007
TB	0.0331	SAUR	0.0140	NFGNR	0.0032	C_PNEU	0.0006
PARA_3	0.0330	FLU_B	0.0087	SASP	0.0025	BOPE	0.0005

## 2. Top 5 singleton causes and NoA









## 3. Top 5 singleton causes with pairs and NoA

