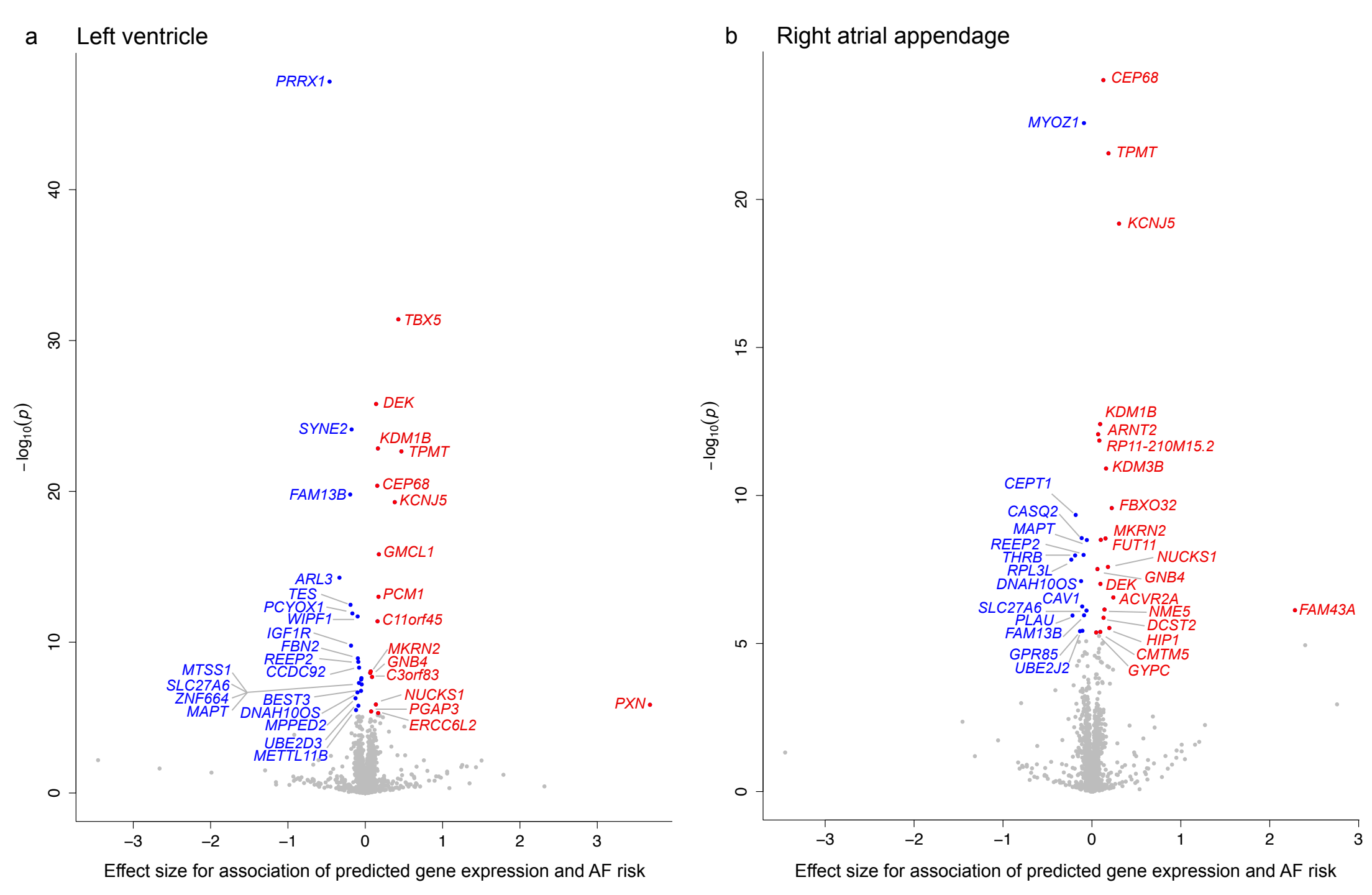
a
Left ventricle
PRRX1
TBX5
DEK
SYNE2
KDM1B
TPMT
CEP68
FAM13B
KCNJ5
GMCL1
ARL3
TES
PCM1
PCYOX1
WIPF1
C11orf45
IGF1R
FBN2
MKRN2
REEP2
GNB4
CCDC92
C3orf83
MTSS1
SLC27A6
NUCKS1
ZNF664
BEST3
PGAP3
MAPT
DNAH10OS
ERCC6L2
MPPED2
UBE2D3
METTL11B
PXN
−log10(p)
Effect size for association of predicted gene expression and AF risk
b
Right atrial appendage
CEP68
MYOZ1
TPMT
KCNJ5
KDM1B
ARNT2
RP11-210M15.2
KDM3B
CEPT1
FBXO32
CASQ2
MAPT
MKRN2
REEP2
FUT11
THRB
NUCKS1
RPL3L
GNB4
DNAH10OS
DEK
CAV1
ACVR2A
SLC27A6
NME5
FAM43A
PLAU
DCST2
FAM13B
HIP1
GPR85
CMTM5
UBE2J2
GYPC
−log10(p)
Effect size for association of predicted gene expression and AF risk