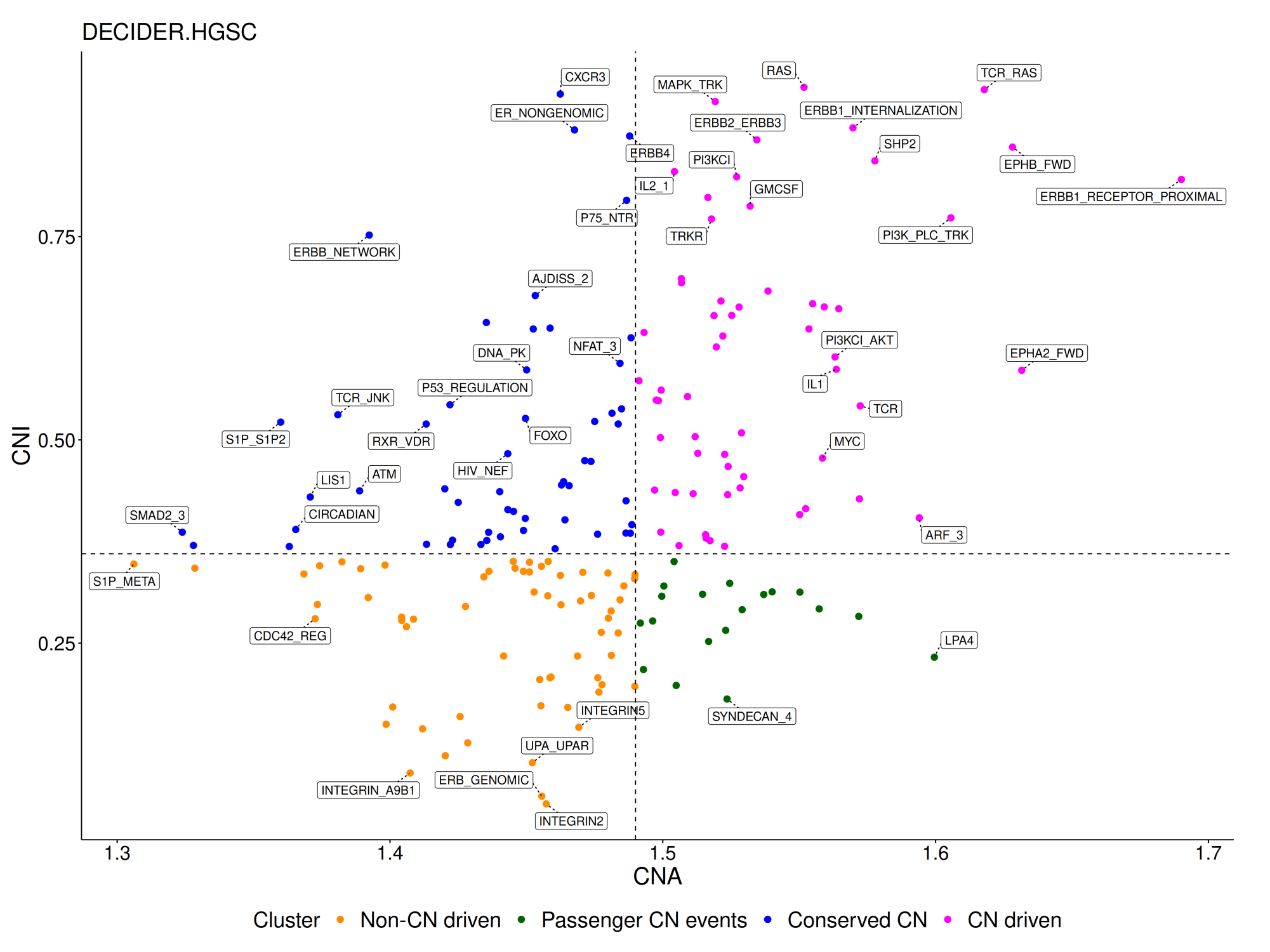

DECIDER.HGSC
CXCR3
MAPK_TRK
RAS
TCR_RAS
ER_NONGENOMIC
ERBB2_ERBB3
ERBB1_INTERNALIZATION
ERBB4
SHP2
PI3KCI
EPHB_FWD
IL2_1
GMCSF
ERBB1_RECEPTOR_PROXIMAL
P75_NTR
TRKR
PI3K_PLC_TRK
ERBB_NETWORK
AJDISS_2
NFAT_3
PI3KCI_AKT
EPHA2_FWD
DNA_PK
IL1
P53_REGULATION
TCR_JNK
TCR
S1P_S1P2
RXR_VDR
FOXO
MYC
HIV_NEF
LIS1
ATM
SMAD2_3
CIRCADIAN
ARF_3
S1P_META
CDC42_REG
LPA4
INTEGRIN5
SYNDECAN_4
UPA_UPAR
ERB_GENOMIC
INTEGRIN_A9B1
INTEGRIN2
0.75
0.50
0.25
CNI
1.3
1.4
1.5
1.6
1.7
CNA
Cluster
Non-CN driven
Passenger CN events
Conserved CN
CN driven