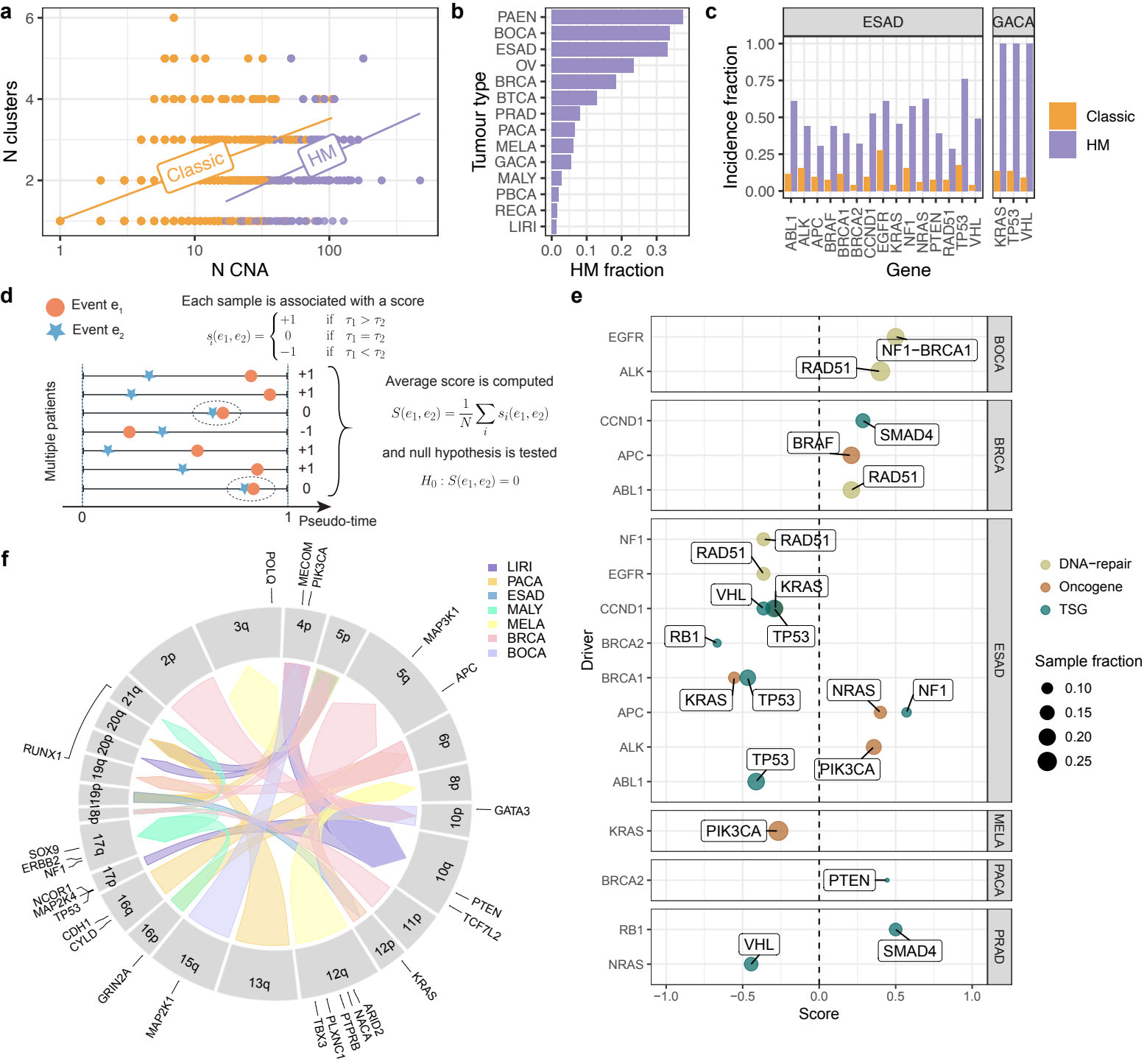
a
N clusters
N CNA
Classic
HM
b
Tumour type
HM fraction
PAEN
BOCA
ESAD
OV
BRCA
BTCA
PRAD
PACA
MELA
GACA
MALY
PBCA
RECA
LIRI
c
ESAD
GACA
Incidence fraction
Gene
Classic
HM
d
Event e1
Event e2
Each sample is associated with a score
Multiple patients
Pseudo-time
Average score is computed
and null hypothesis is tested
e
Driver
Score
DNA−repair
Oncogene
TSG
Sample fraction
f
LIRI
PACA
ESAD
MALY
MELA
BRCA
BOCA