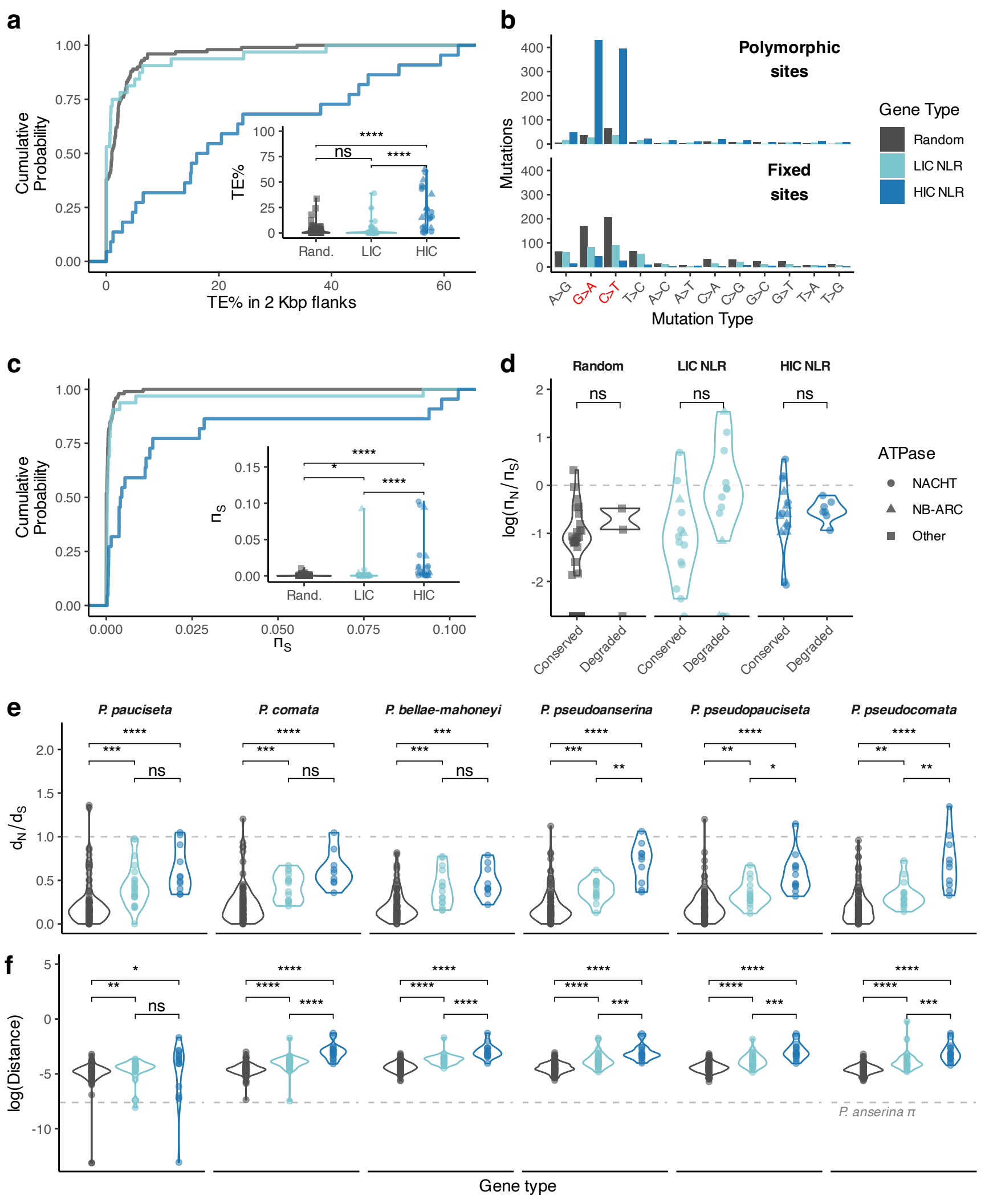
a
Cumulative Probability
TE% in 2 Kbp flanks
TE%
Rand.
LIC
HIC
ns
****
b
Polymorphic sites
Fixed sites
Mutations
Mutation Type
A>G
G>A
C>T
T>C
A>C
A>T
C>A
C>G
G>C
G>T
T>A
T>G
Gene Type
Random
LIC NLR
HIC NLR
c
Cumulative Probability
π_S
Rand.
LIC
HIC
****
*
d
Random
LIC NLR
HIC NLR
ns
$\log(\pi_N/\pi_S)$
Conserved
Degraded
ATPase
NACHT
NB-ARC
Other
e
P. pauciseta
P. comata
P. bellae-mahoneyi
P. pseudoanserina
P. pseudopauciseta
P. pseudocomata
d_N/d_S
****
***
**
*
ns
f
log(Distance)
P. anserina π
Gene type