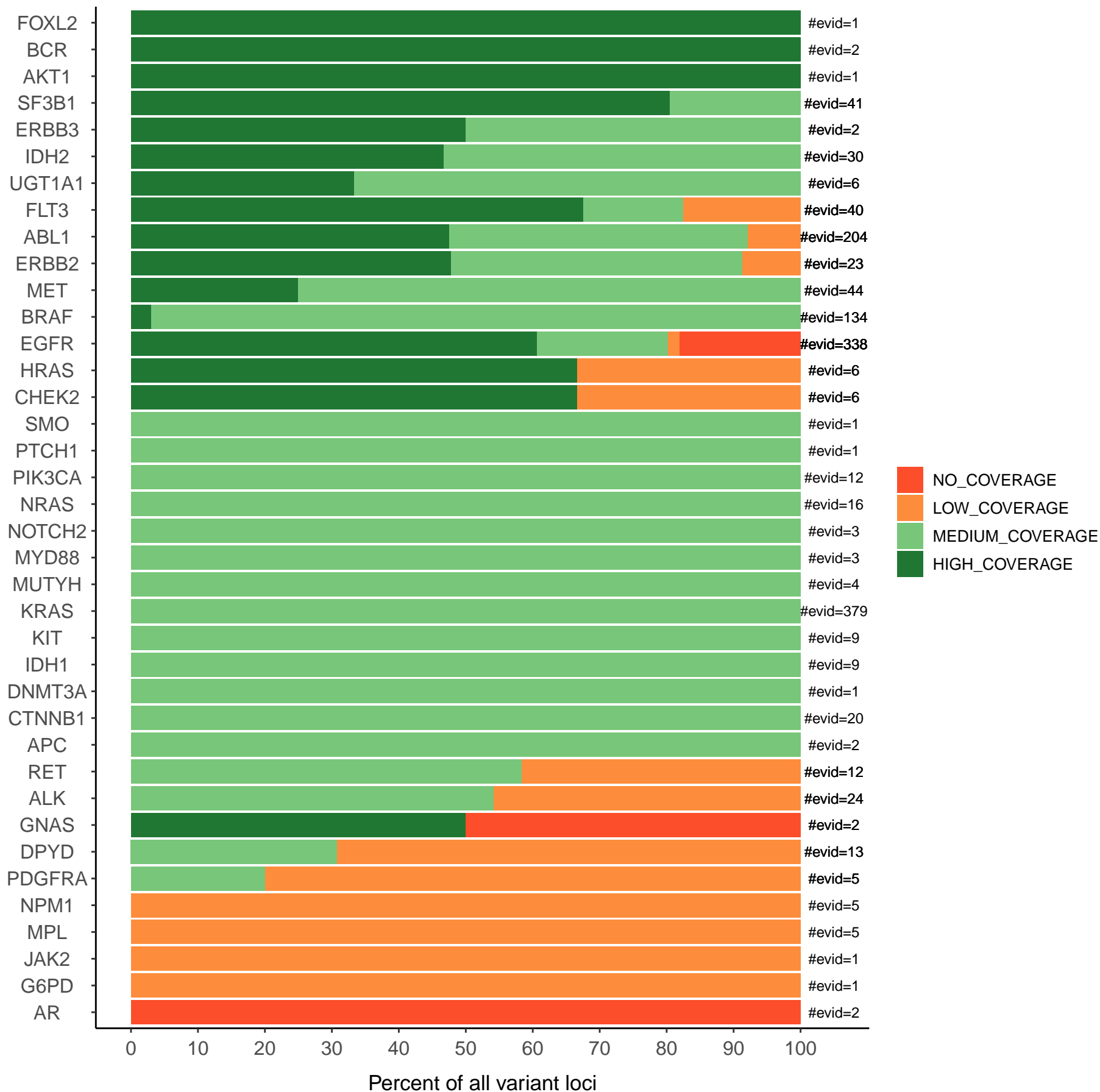
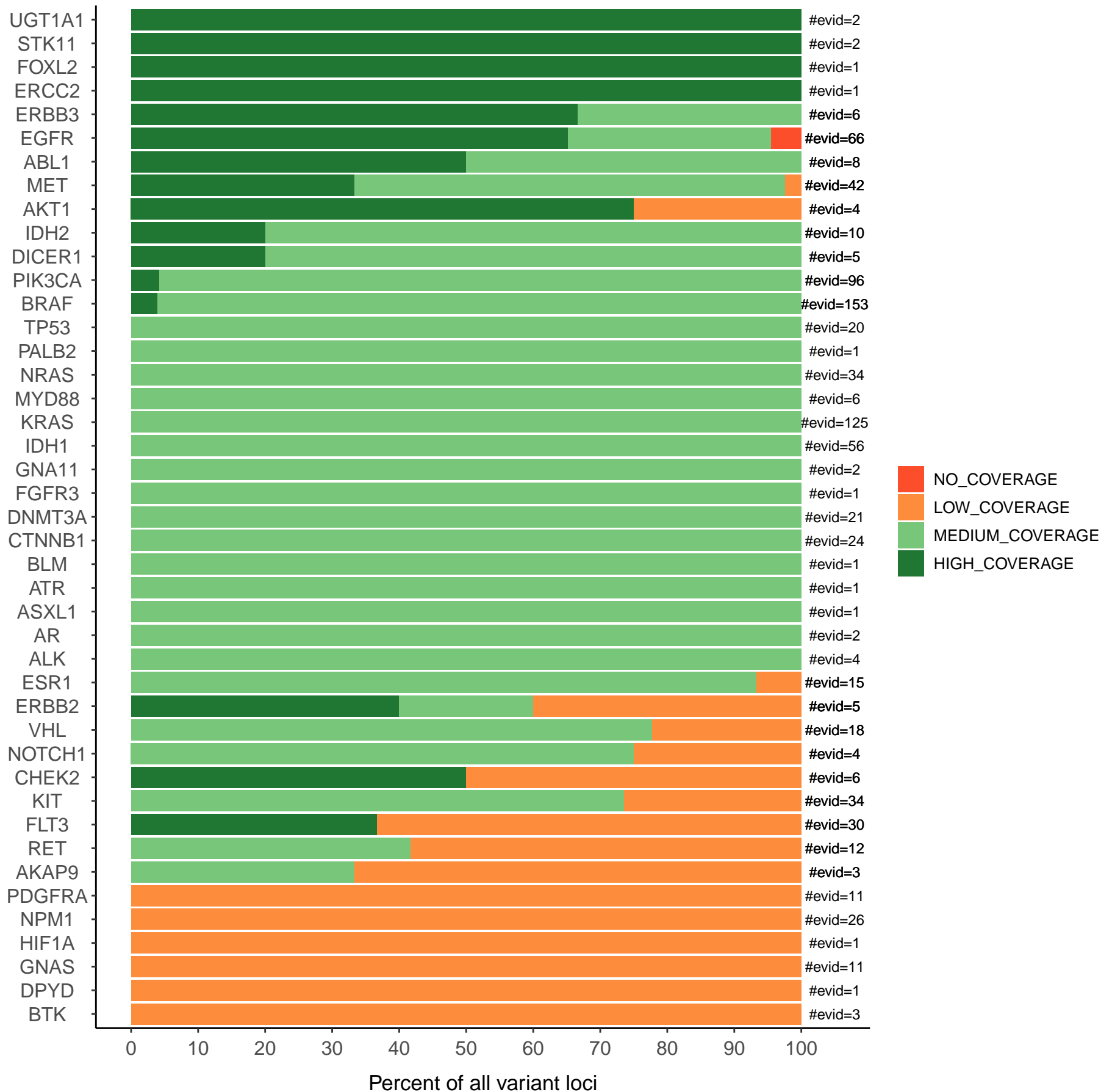


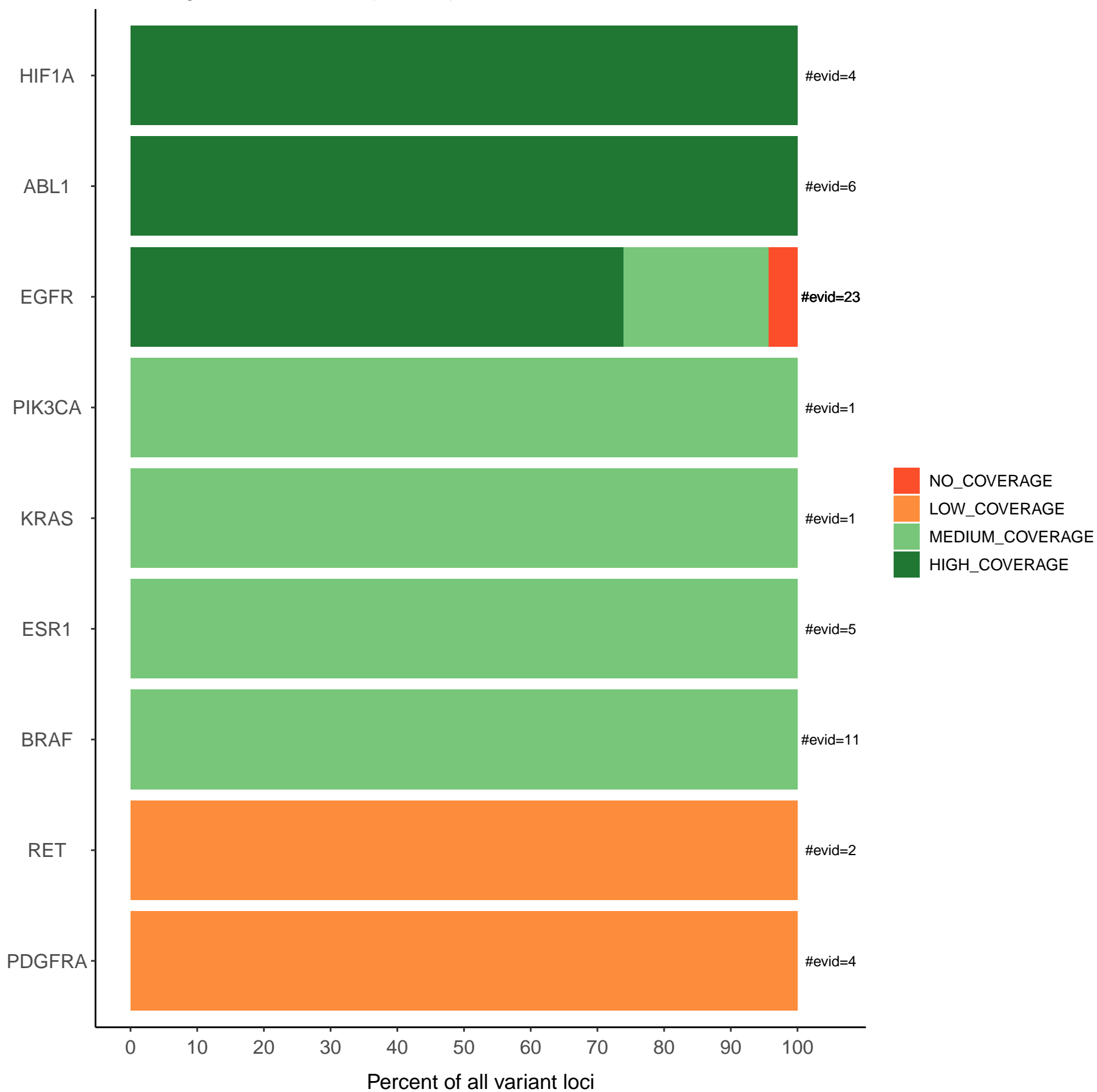
Clinical significant evidences (level A)



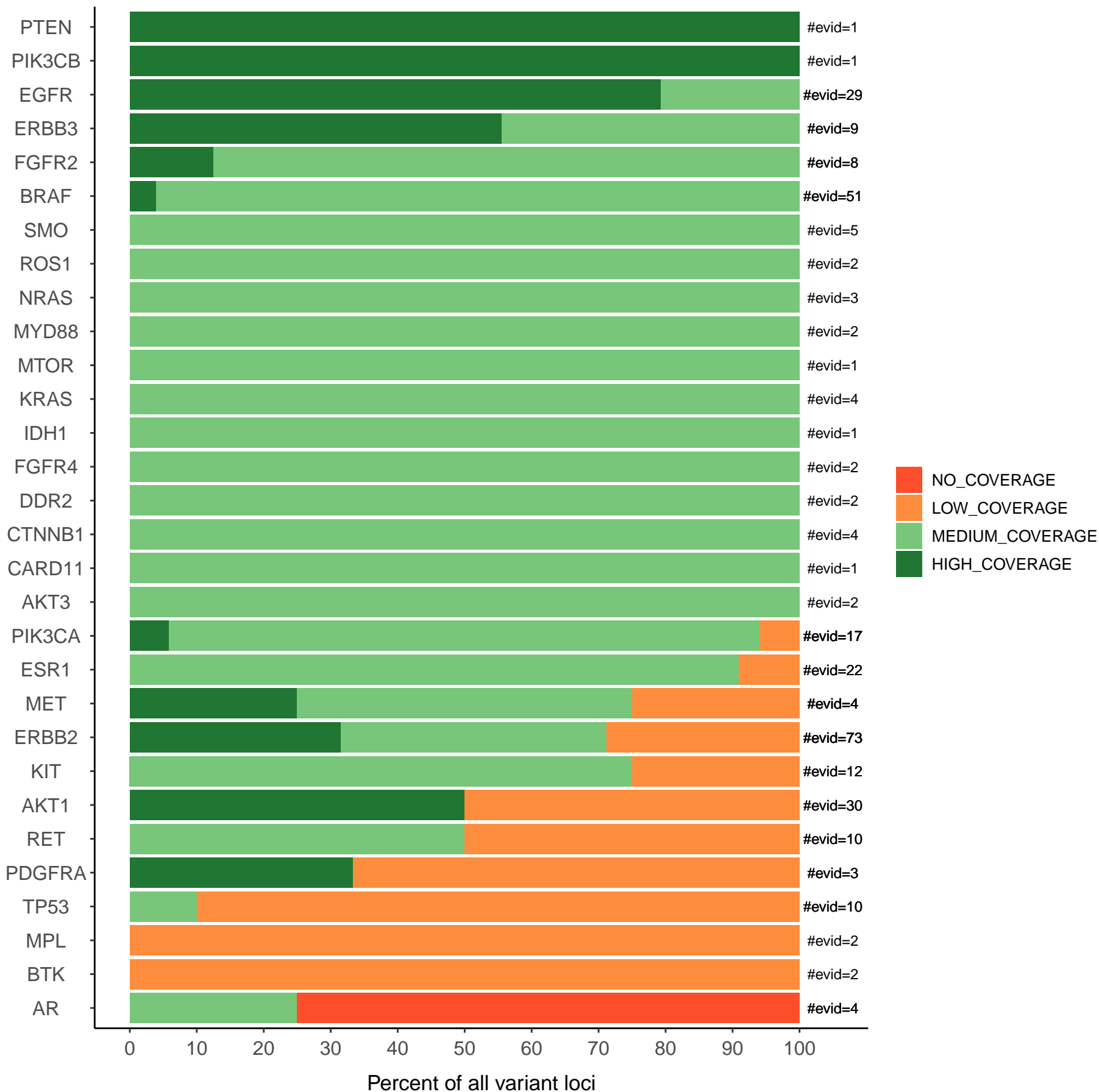
Clinical significant evidences (level B)



Clinical significant evidences (level B1)



Clinical significant evidences (level B2)



Clinical significant evidences (level B3)

ERCC2

#evid=1

ERCC1

#evid=2

CHEK2

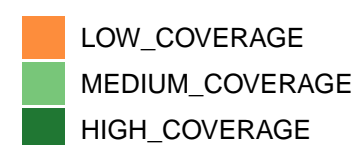
#evid=3

BRAF

#evid=2

TP53

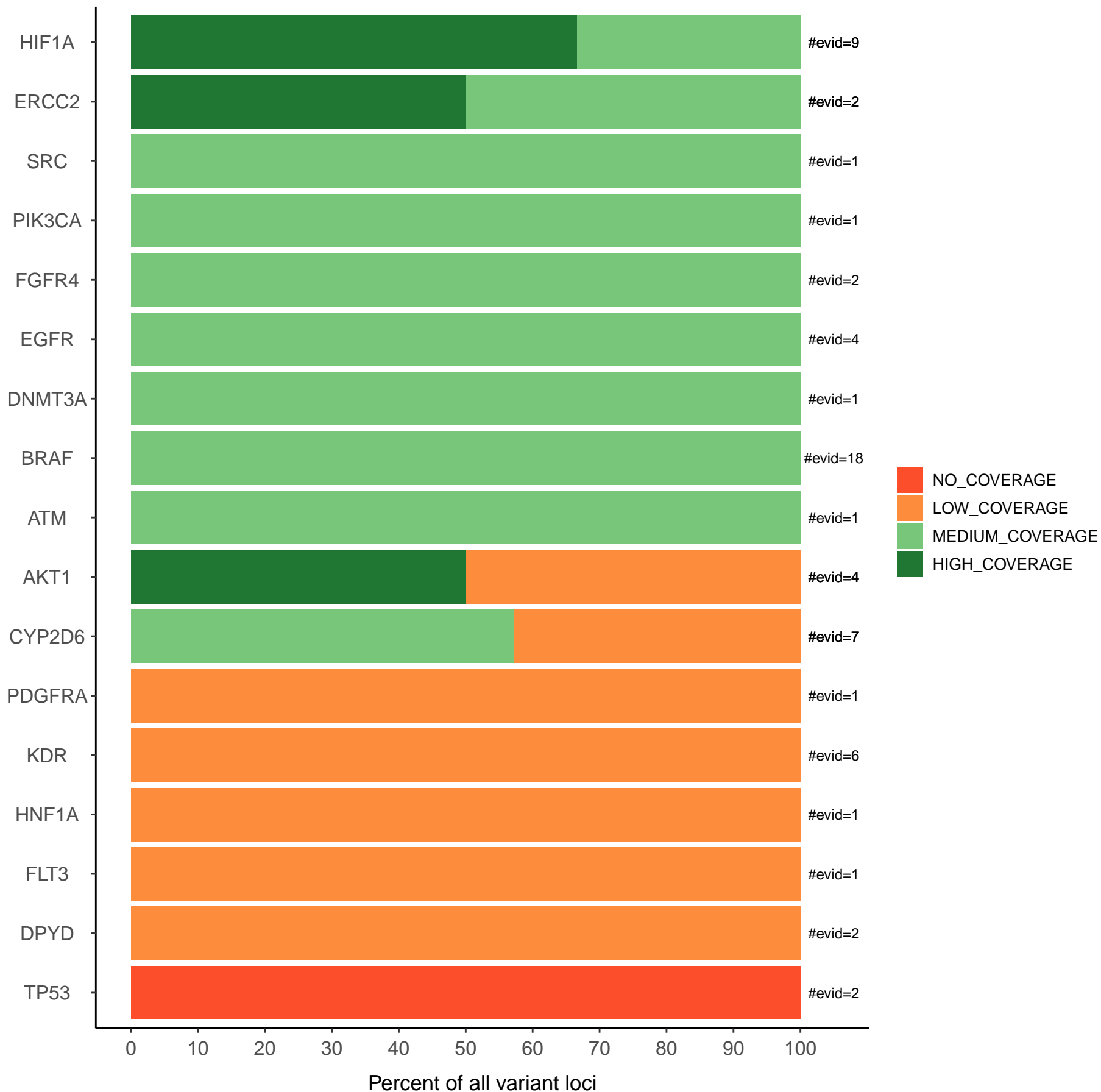
#evid=1



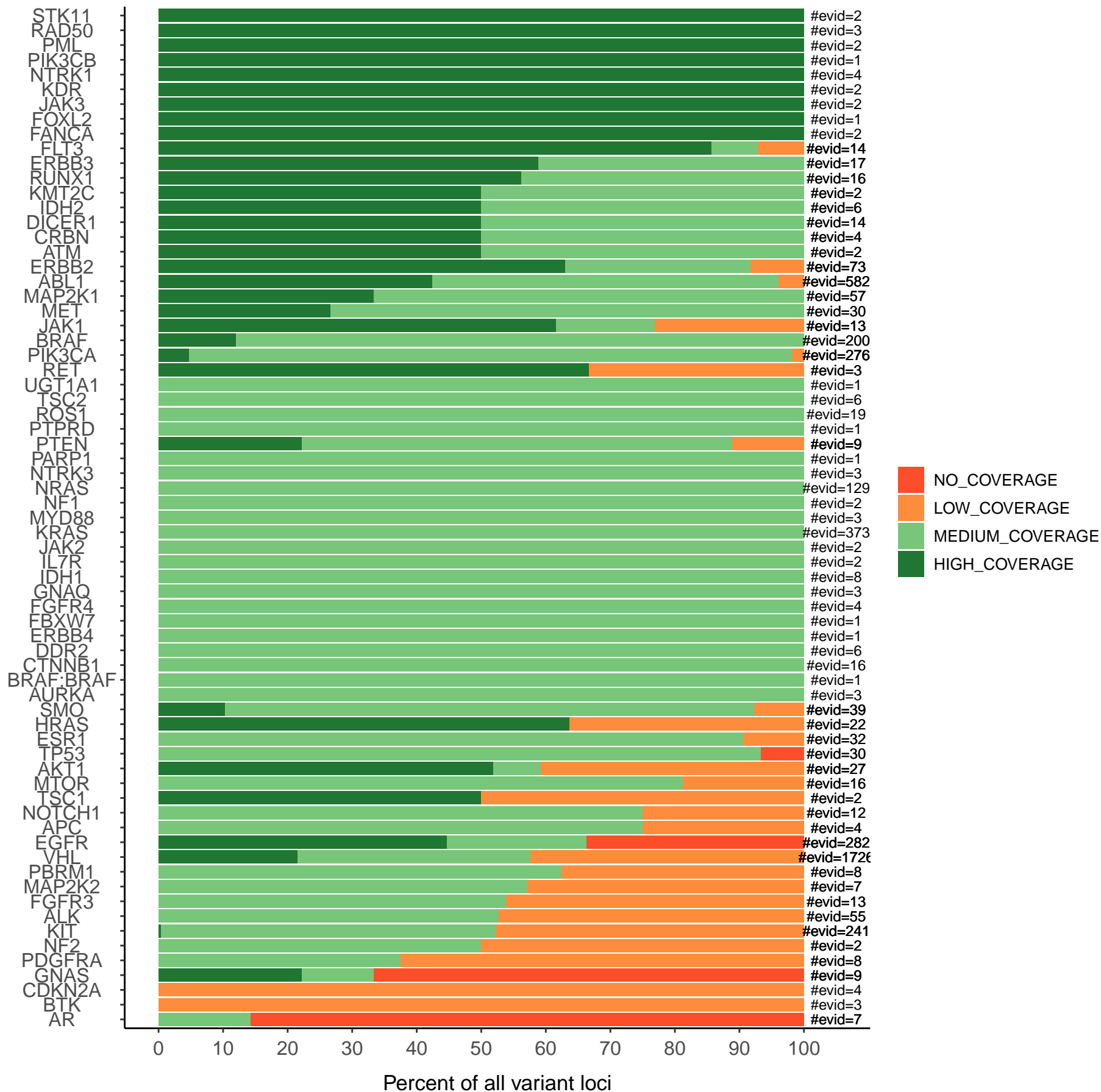
0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

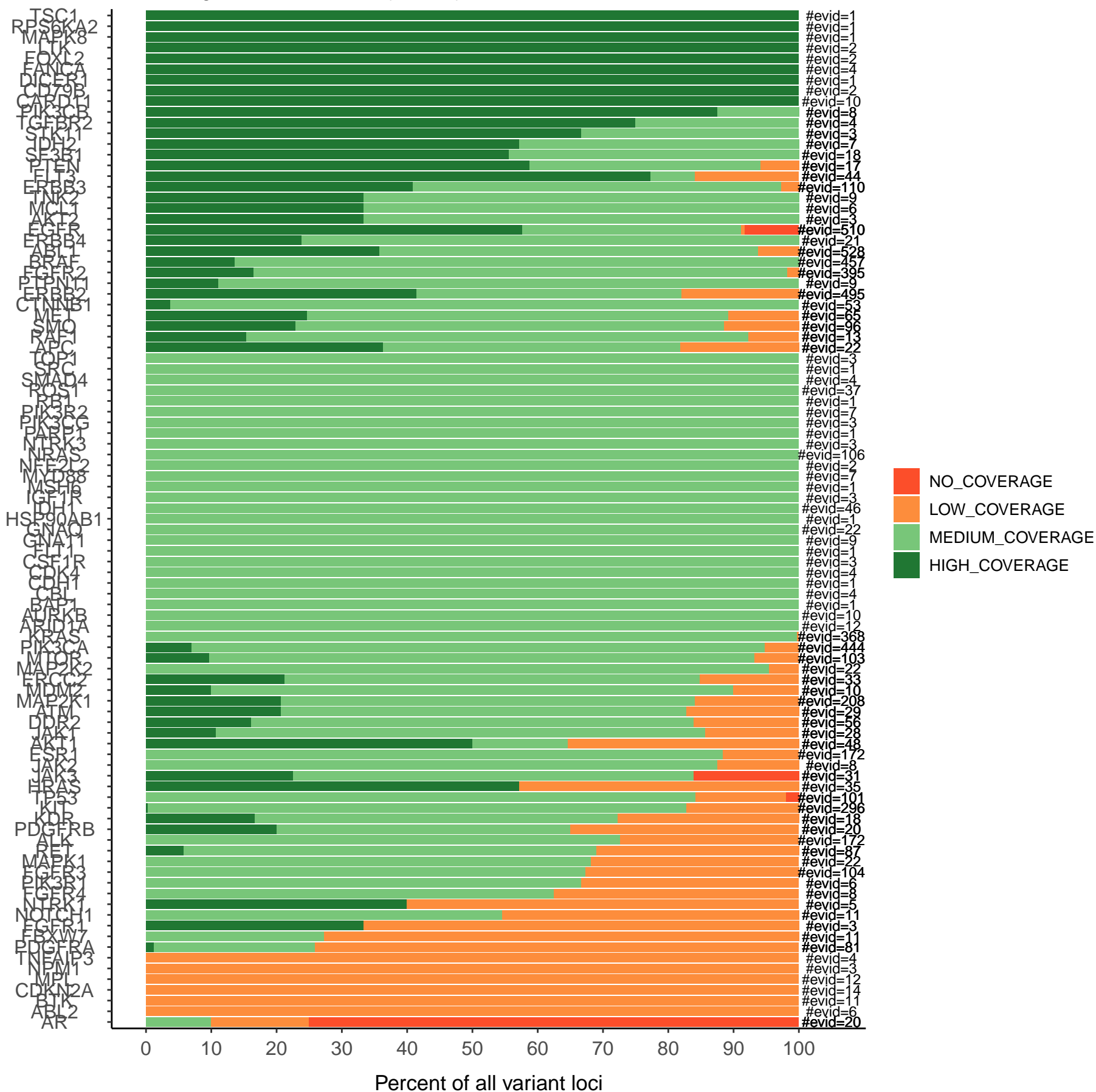
Clinical significant evidences (level B4)



Clinical significant evidences (level C)



Clinical significant evidences (level D)



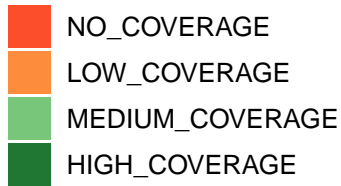
Clinical significant evidences (level A), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=1408



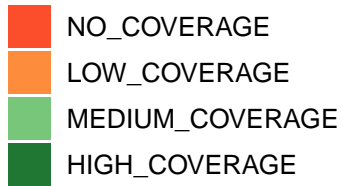
Clinical significant evidences (level B), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=875



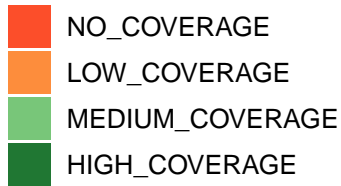
Clinical significant evidences (level B1), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=57



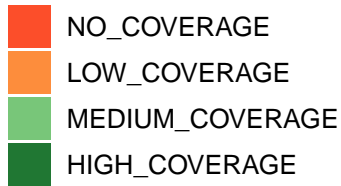
Clinical significant evidences (level B2), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=317



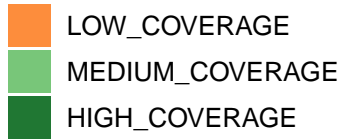
Clinical significant evidences (level B3), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=9



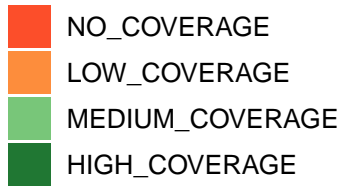
Clinical significant evidences (level B4), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=63



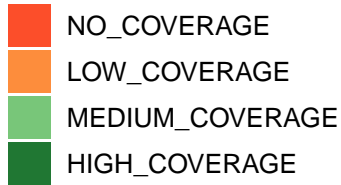
Clinical significant evidences (level C), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=4471



Clinical significant evidences (level D), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=5820

