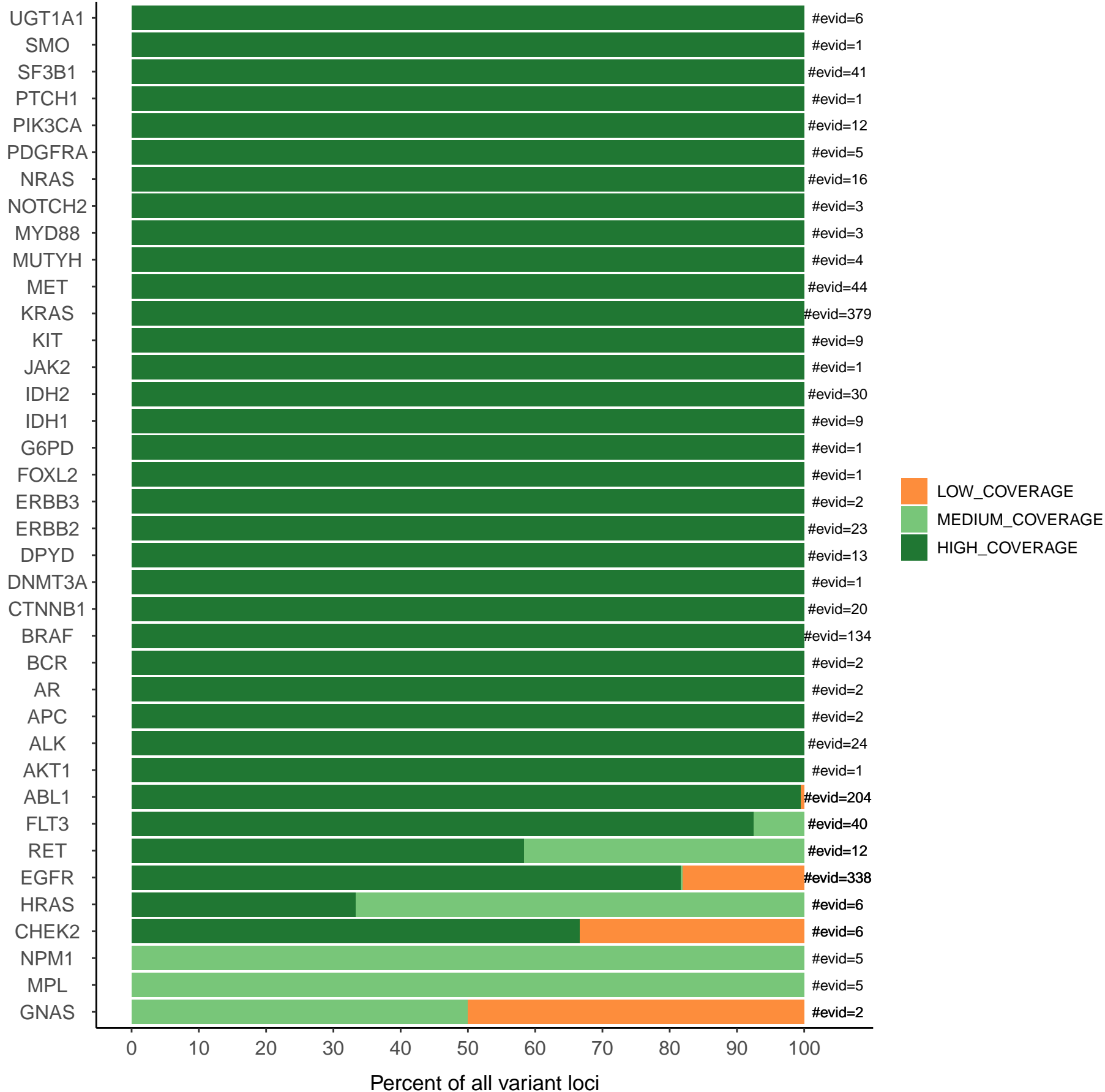
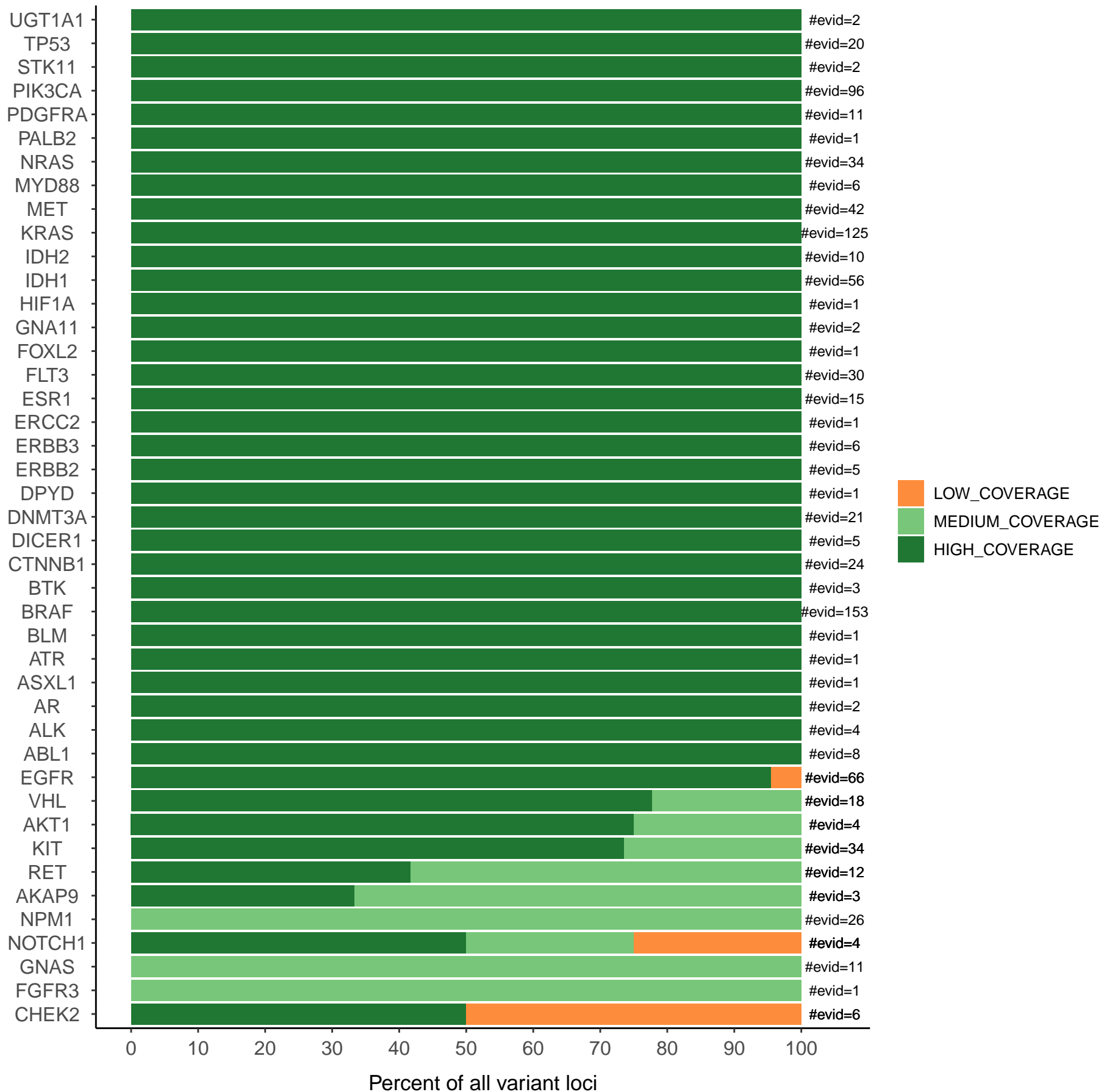


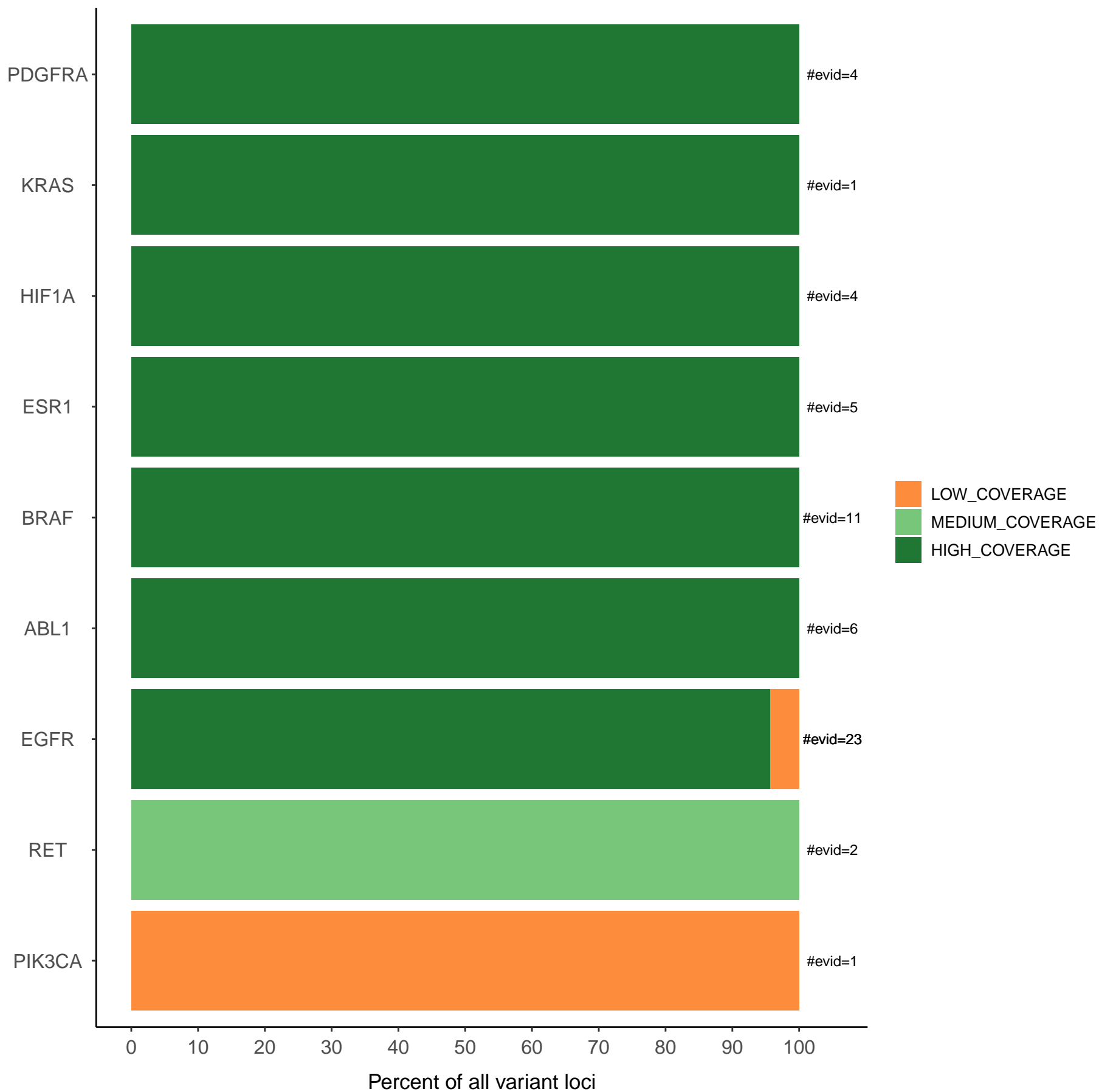
Clinical significant evidences (level A)



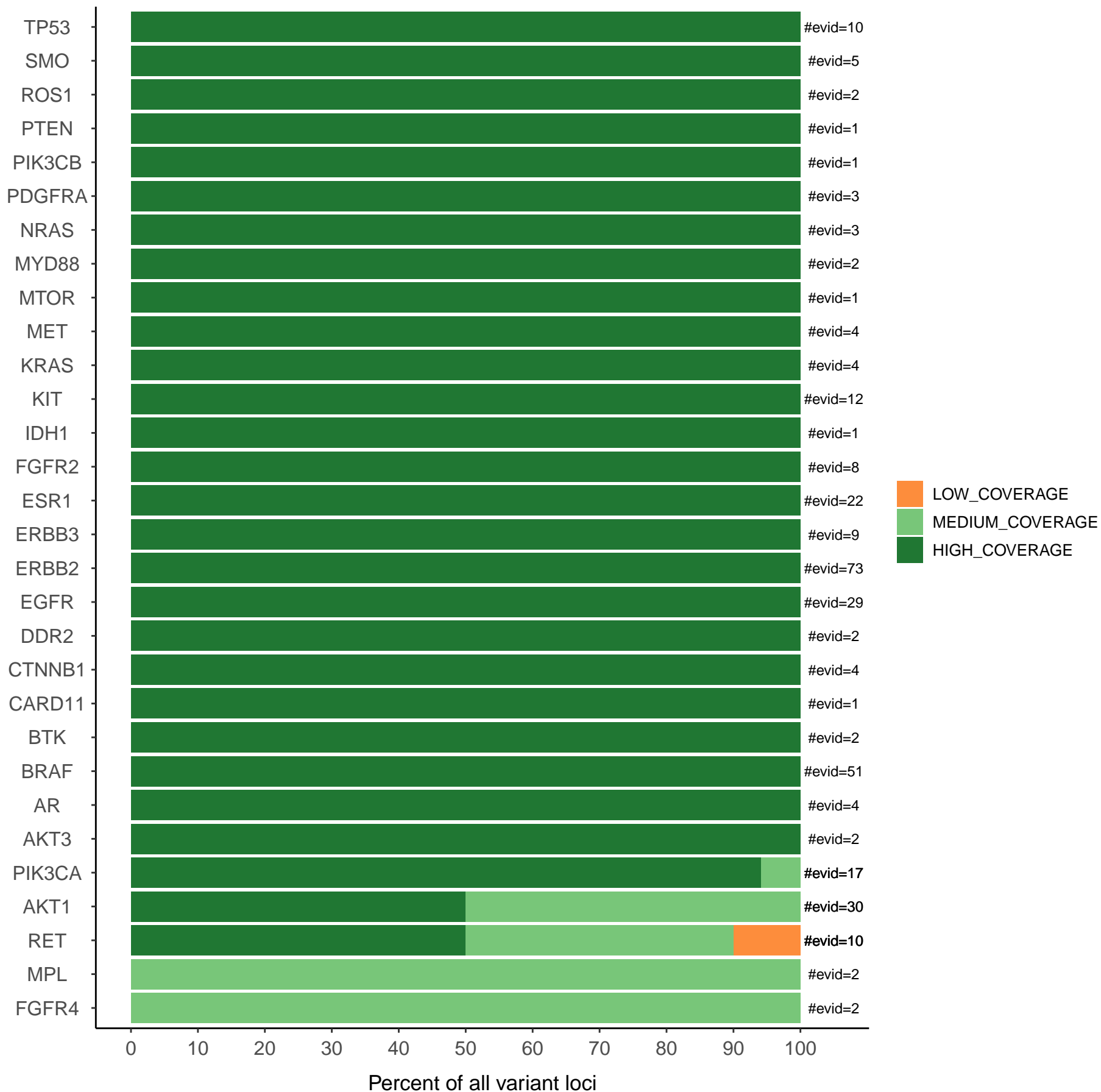
Clinical significant evidences (level B)



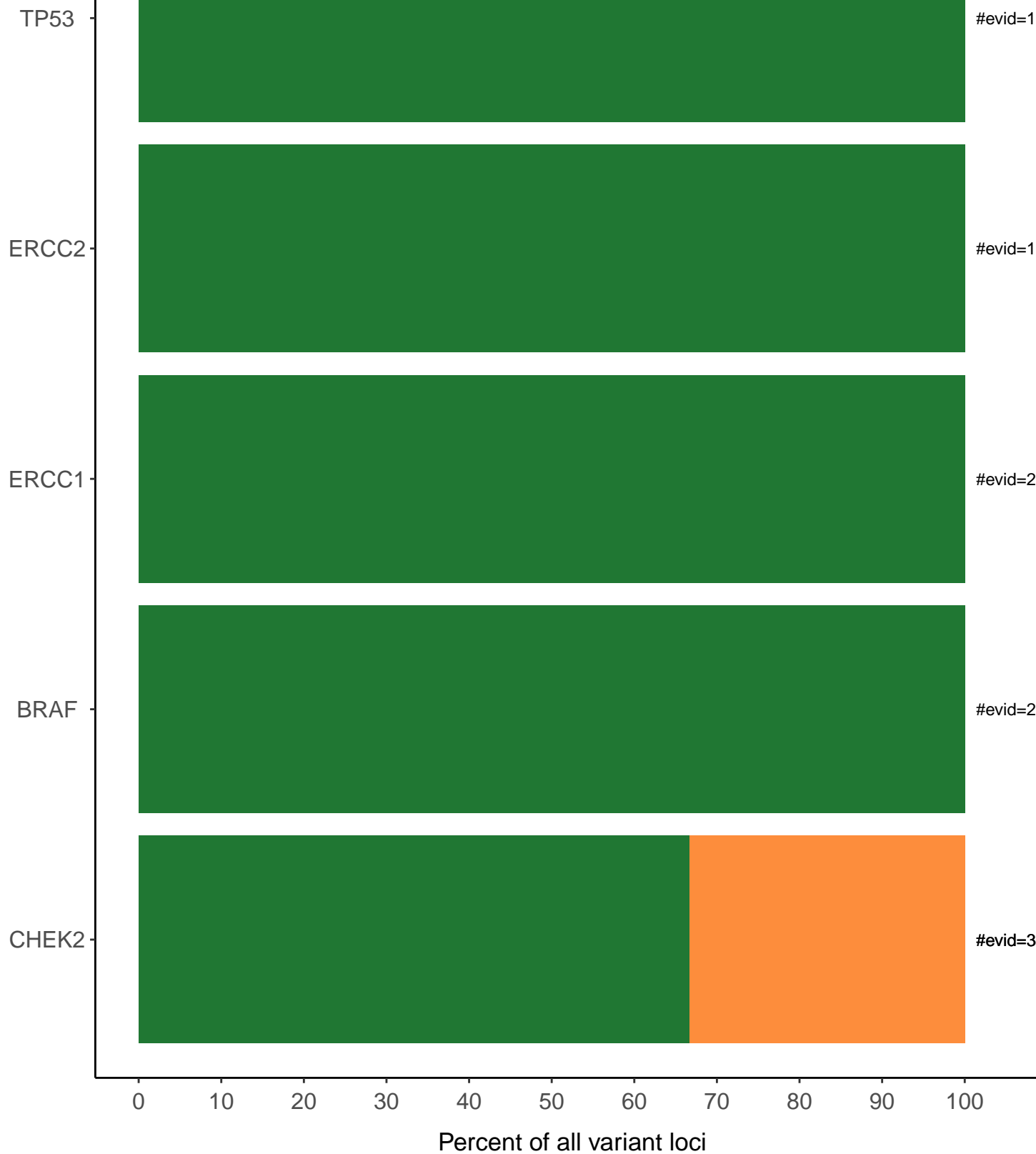
Clinical significant evidences (level B1)



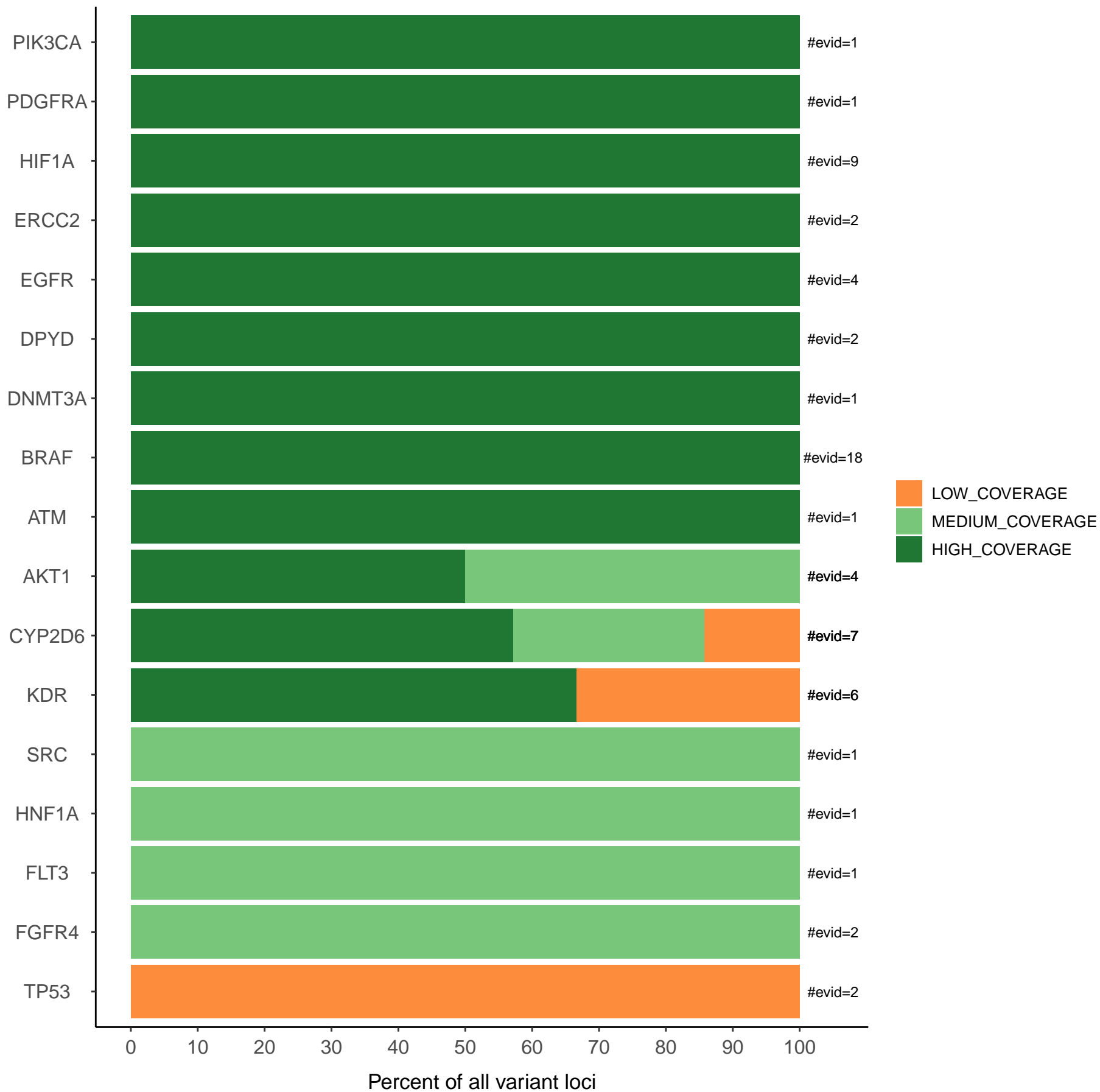
Clinical significant evidences (level B2)



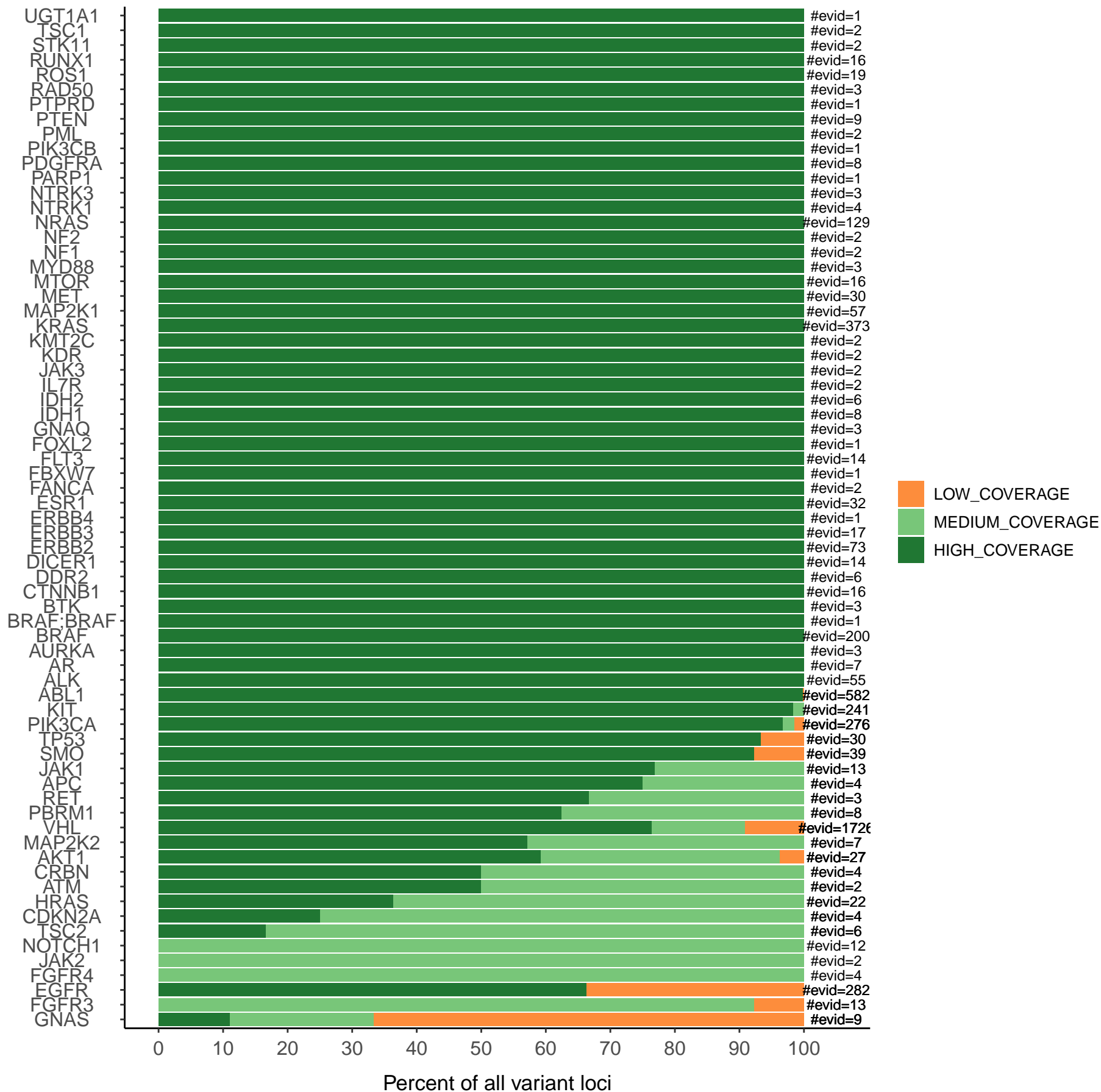
Clinical significant evidences (level B3)



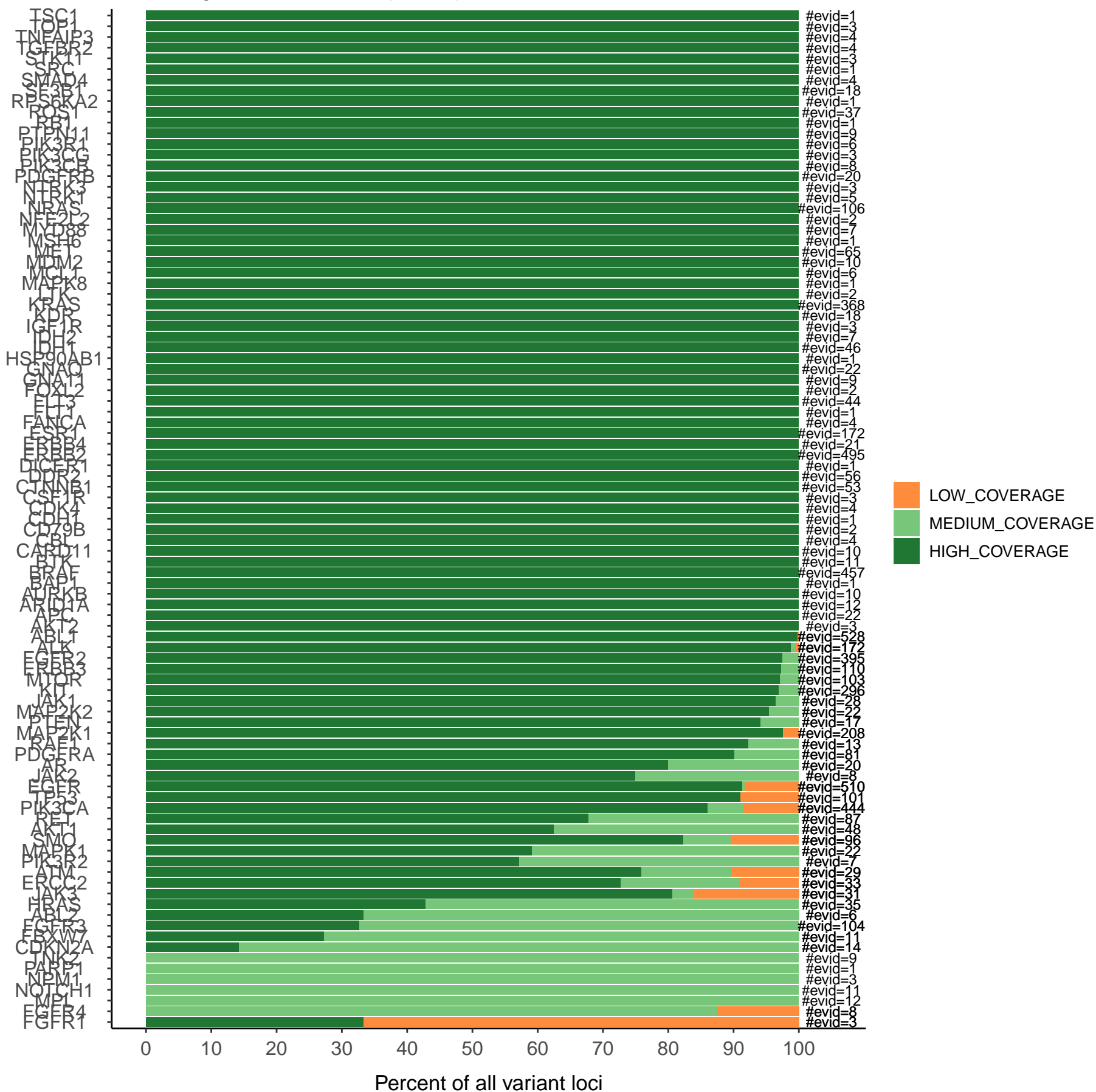
Clinical significant evidences (level B4)



Clinical significant evidences (level C)



Clinical significant evidences (level D)



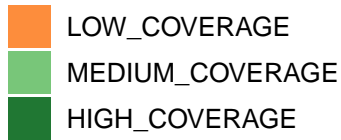
Clinical significant evidences (level A), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=1408



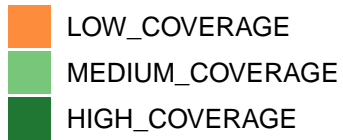
Clinical significant evidences (level B), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=875



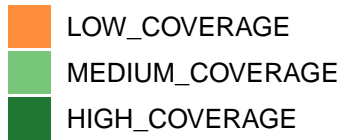
Clinical significant evidences (level B1), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=57



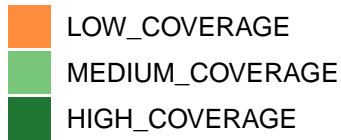
Clinical significant evidences (level B2), all gene summary

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



LOW_COVERAGE

HIGH_COVERAGE

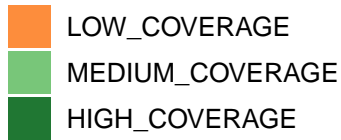
Clinical significant evidences (level B4), all gene summary

CALLABILITY

0 10 20 30 40 50 60 70 80 90 100

Percent of all variant loci

#evid=63



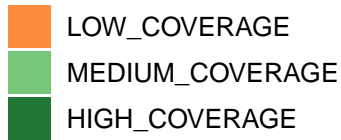
Clinical significant evidences (level C), all gene summary

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

