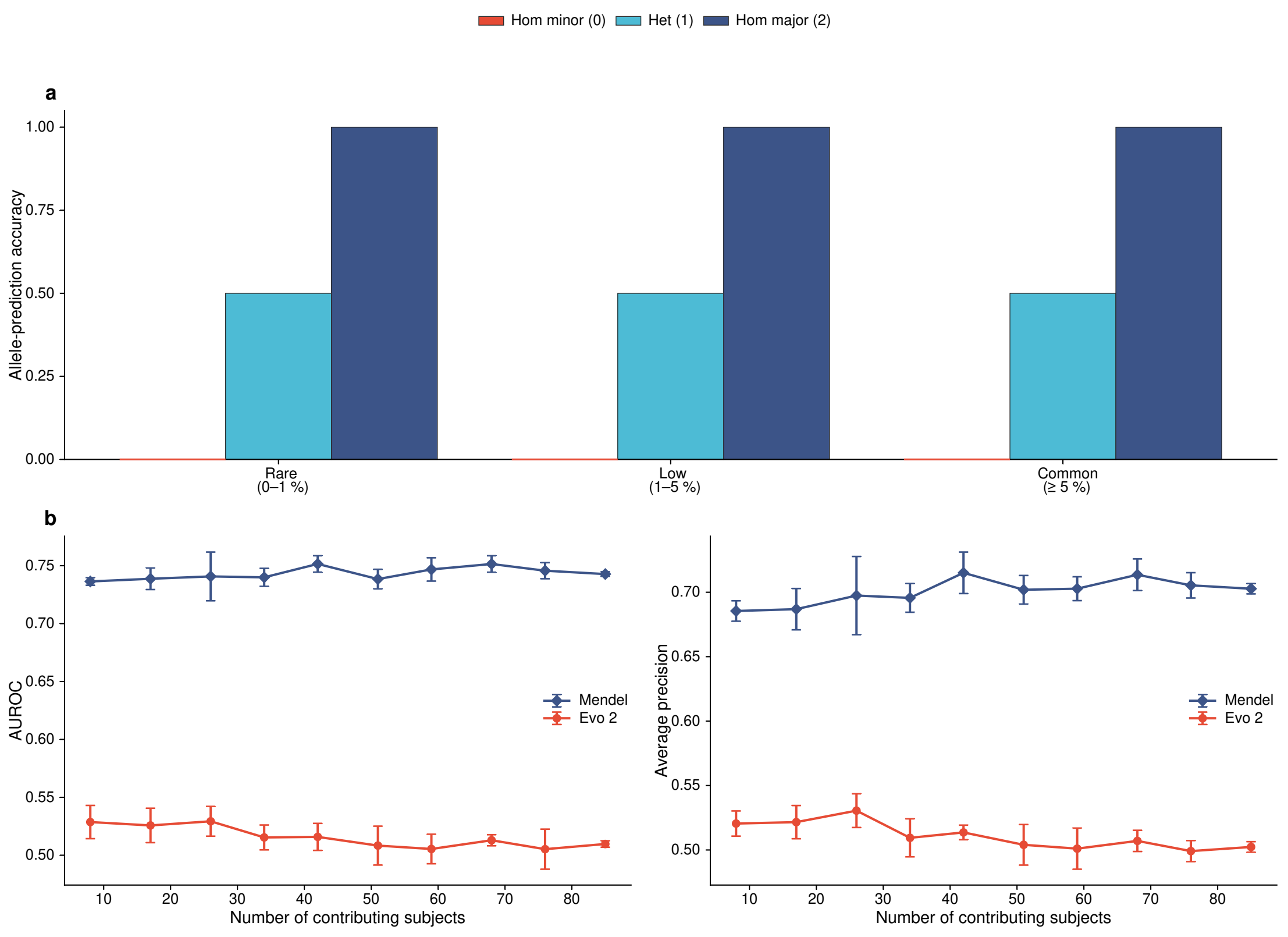
Hom minor (0)
Het (1)
Hom major (2)
a
Allele-prediction accuracy
1.00
0.75
0.50
0.25
0.00
Rare
(0–1 %)
Low
(1–5 %)
Common
(≥ 5 %)
b
AUROC
0.75
0.70
0.65
0.60
0.55
0.50
Mendel
Evo 2
10
20
30
40
50
60
70
80
Number of contributing subjects
Average precision
0.70
0.65
0.60
0.55
0.50
Mendel
Evo 2
10
20
30
40
50
60
70
80
Number of contributing subjects