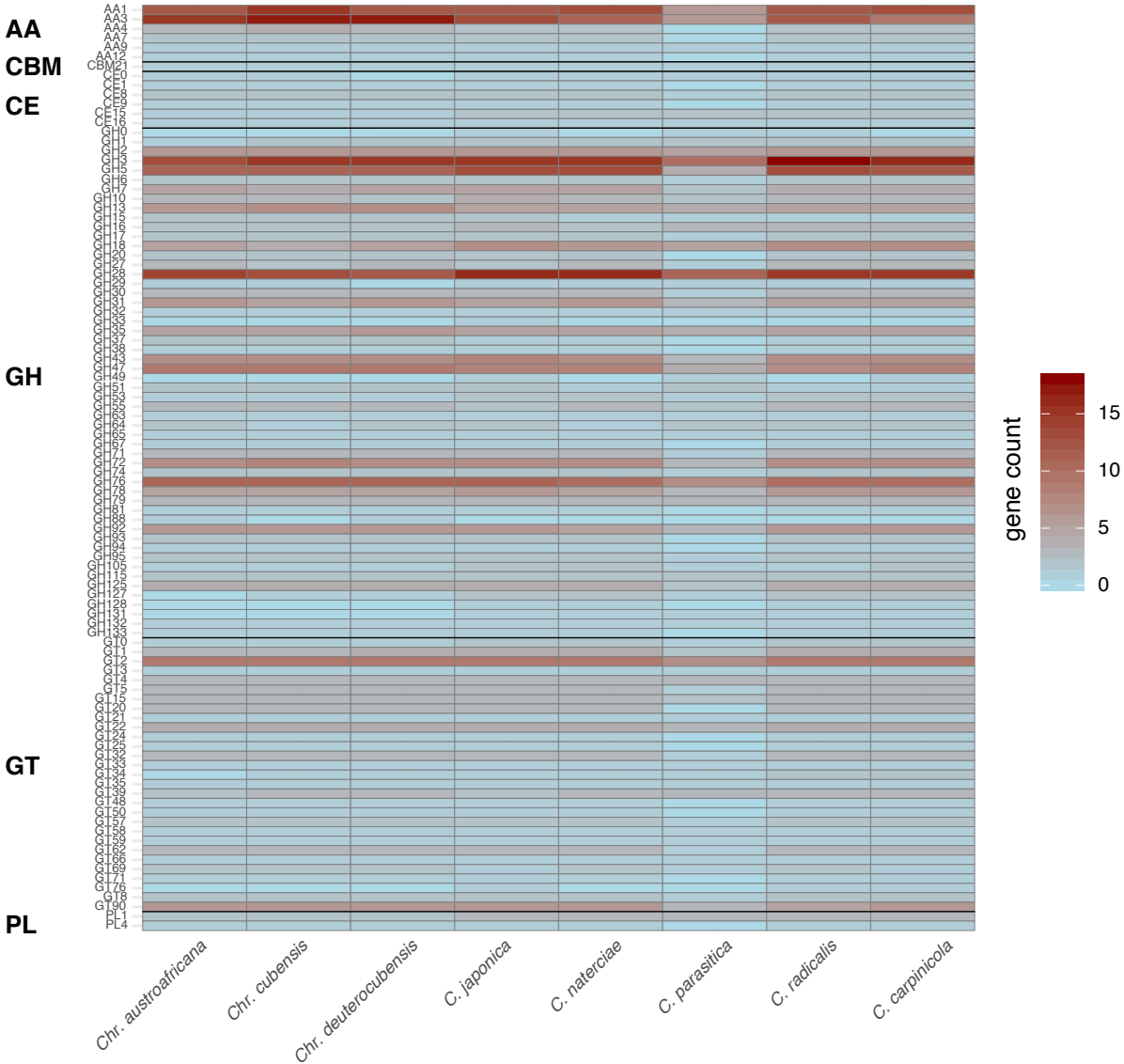
AA
CBM
CE
GH
GT
PL
AA1
AA3
AA4
AA7
AA9
AA12
CBM21
CE0
CE1
CE8
CE9
CE15
CE16
GH0
GH1
GH9
GH13
GH5
GH6
GH7
GH10
GH13
GH15
GH16
GH17
GH18
GH20
GH27
GH28
GH29
GH30
GH31
GH32
GH35
GH36
GH37
GH38
GH43
GH47
GH49
GH51
GH53
GH55
GH63
GH64
GH65
GH67
GH71
GH72
GH74
GH76
GH78
GH79
GH81
GH88
GH92
GH93
GH94
GH95
GH105
GH115
GH125
GH127
GH128
GH131
GH132
GH133
GT0
GT1
GT2
GT3
GT4
GT5
GT15
GT20
GT21
GT22
GT24
GT25
GT32
GT33
GT34
GT35
GT39
GT48
GT50
GT57
GT58
GT59
GT62
GT66
GT69
GT71
GT76
GT8
GT90
PL1
PL4
Chr. austroafricana
Chr. cubensis
Chr. deuterocubensis
C. japonica
C. naterciae
C. parasitica
C. radicalis
C. carpinicola
gene count
15
10
5
0