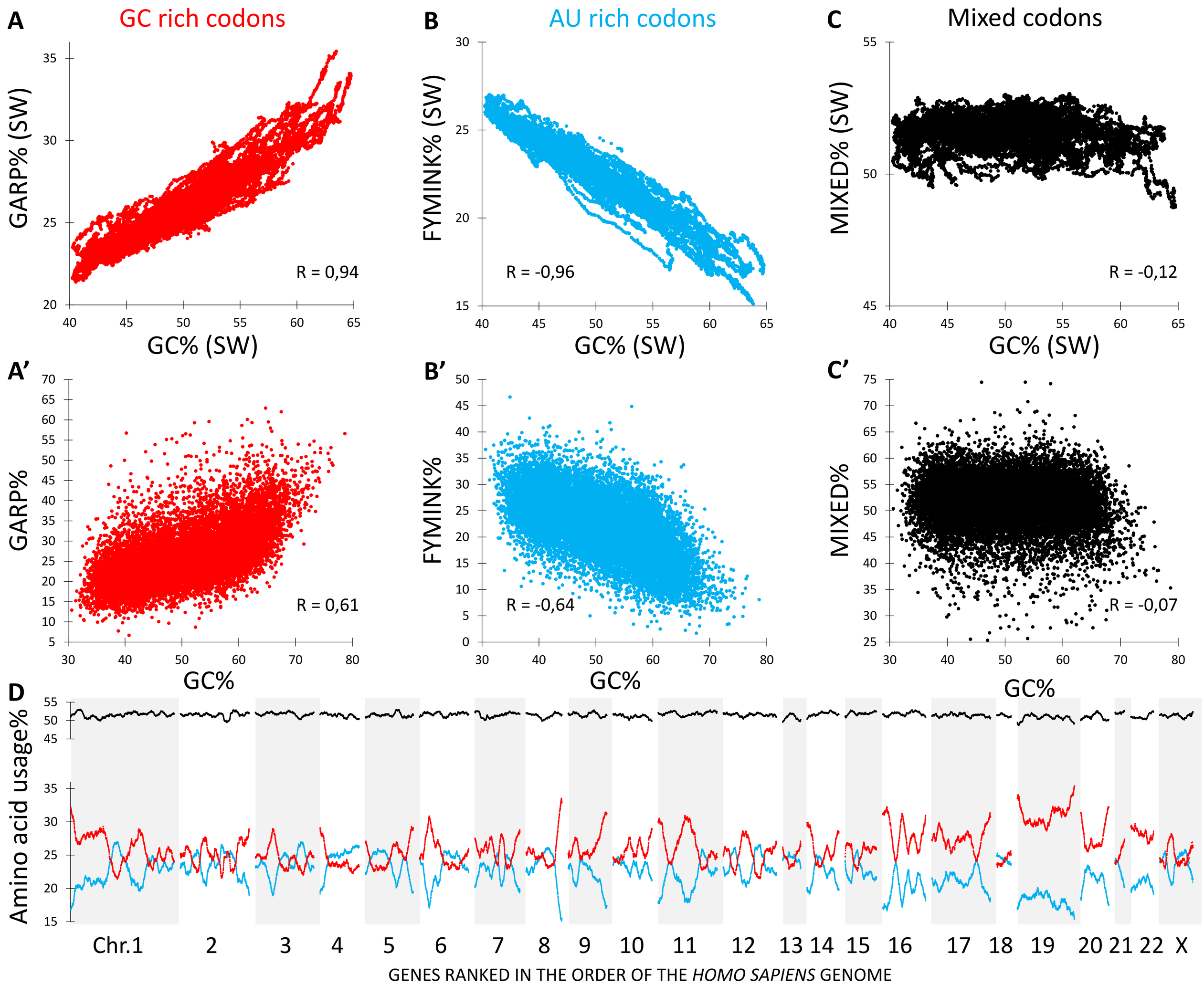
A
GC rich codons
GARP% (SW)
GC% (SW)
R = 0,94
B
AU rich codons
FYMINK% (SW)
GC% (SW)
R = -0,96
C
Mixed codons
MIXED% (SW)
GC% (SW)
R = -0,12
A'
GARP%
GC%
R = 0,61
B'
FYMINK%
GC%
R = -0,64
C'
MIXED%
GC%
R = -0,07
D
Amino acid usage%
Chr.1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 X
GENES RANKED IN THE ORDER OF THE *HOMO SAPIENS* GENOME