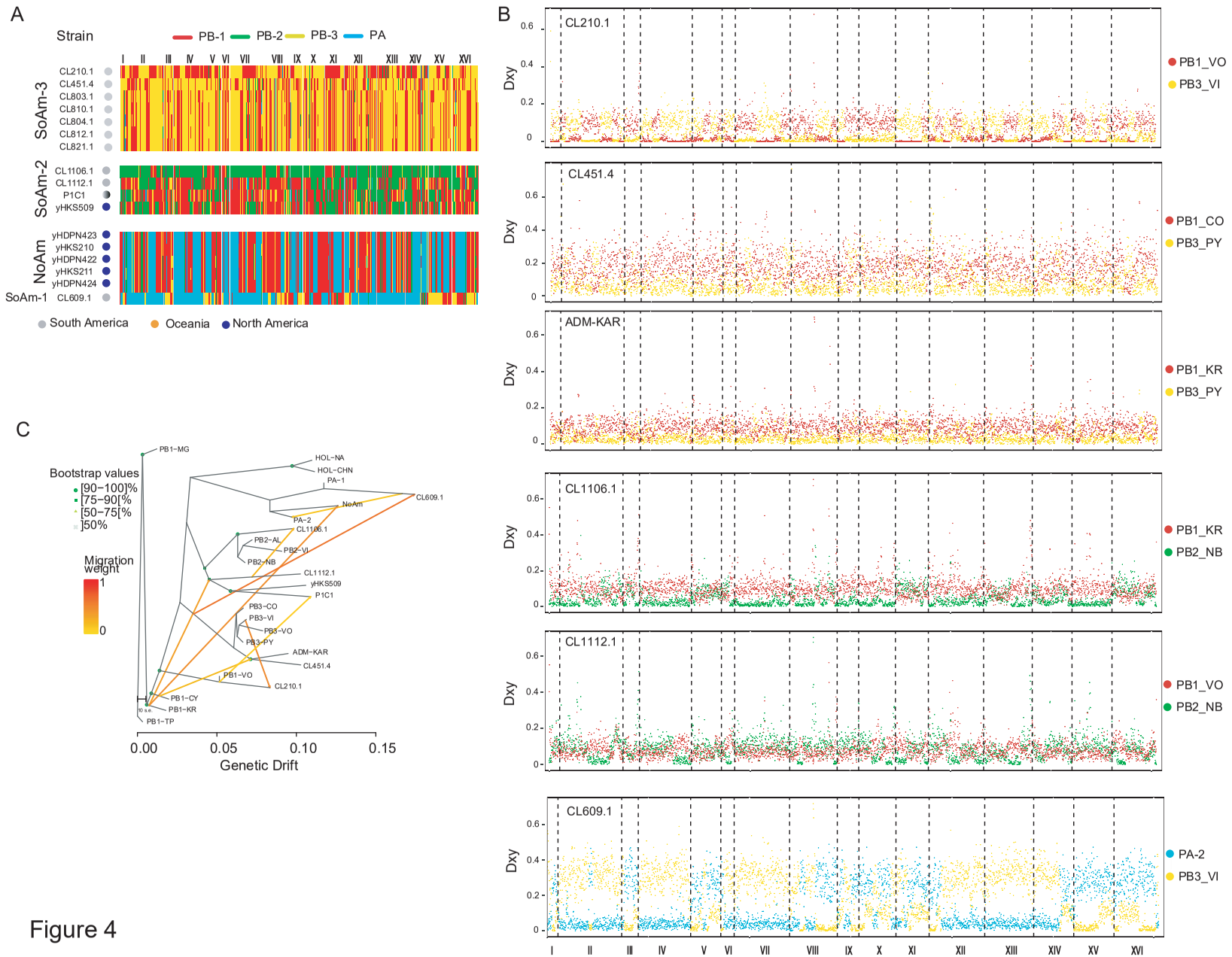
A
Strain
PB-1
PB-2
PB-3
PA
I II III IV V VI VII VIII IX X XI XII XIII XIV XV XVI
SoAm-3
CL210.1
CL451.4
CL803.1
CL810.1
CL804.1
CL812.1
CL821.1
SoAm-2
CL1106.1
CL1112.1
P1C1
yHKS509
NoAm
yHDPN423
yHKS210
yHDPN422
yHKS211
yHDPN424
SoAm-1
CL609.1
South America
Oceania
North America
B
CL210.1
PB1_VO
PB3_VI
CL451.4
PB1_CO
PB3_PY
ADM-KAR
PB1_KR
PB3_PY
CL1106.1
PB1_KR
PB2_NB
CL1112.1
PB1_VO
PB2_NB
CL609.1
PA-2
PB3_VI
Dxy
0
0.2
0.4
0.6
C
Bootstrap values
[90−100]%
[75−90[%
[50−75[%
]50%
Migration weight
1
0
PB1−MG
HOL−NA
HOL−CHN
PA−1
CL609.1
NoAm
PA−2
CL1106.1
PB2−AL
PB2−VI
PB2−NB
CL1112.1
yHKS509
P1C1
PB3−CO
PB3−VI
PB3−VO
PB3−PY
ADM−KAR
CL451.4
PB1−VO
CL210.1
PB1−CY
PB1−KR
PB1−TP
0.00
0.05
0.10
0.15
Genetic Drift

Figure 4