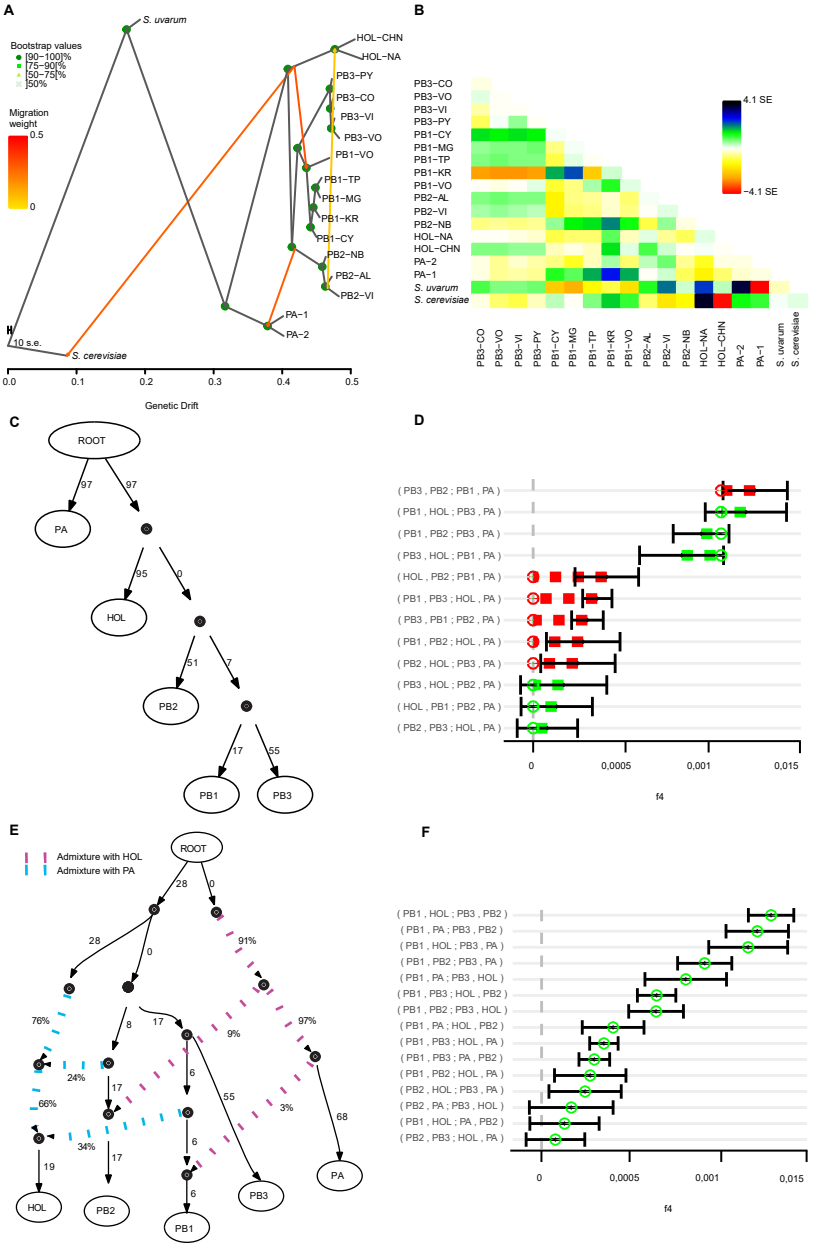
A
S. uvarum
Bootstrap values
[90−100]%
[75−90]%
[50−75]%
50%
Migration weight
0.5
0
HOL−CHN
HOL−NA
PB3−PY
PB3−CO
PB3−VI
PB3−VO
PB1−VO
PB1−TP
PB1−MG
PB1−KR
PB1−CY
PB2−NB
PB2−AL
PB2−VI
PA−1
PA−2
10 s.e.
S. cerevisiae
0.0
0.1
0.2
0.3
0.4
0.5
Genetic Drift
B
PB3−CO
PB3−VO
PB3−VI
PB3−PY
PB1−CY
PB1−MG
PB1−TP
PB1−KR
PB1−VO
PB2−AL
PB2−VI
PB2−NB
HOL−NA
HOL−CHN
PA−2
PA−1
S. uvarum
S. cerevisiae
4.1 SE
−4.1 SE
C
ROOT
97
97
PA
95
0
HOL
51
7
PB2
17
55
PB1
PB3
D
(PB3 , PB2 ; PB1 , PA)
(PB1 , HOL ; PB3 , PA)
(PB1 , PB2 ; PB3 , PA)
(PB3 , HOL ; PB1 , PA)
(HOL , PB2 ; PB1 , PA)
(PB1 , PB3 ; HOL , PA)
(PB3 , PB1 ; PB2 , PA)
(PB1 , PB2 ; HOL , PA)
(PB2 , HOL ; PB3 , PA)
(PB3 , HOL ; PB2 , PA)
(HOL , PB1 ; PB2 , PA)
(PB2 , PB3 ; HOL , PA)
0
0,0005
0,001
0,015
f4
E
Admixture with HOL
Admixture with PA
ROOT
28
0
28
0
91%
76%
8
17
9%
97%
24%
17
6
66%
55
3%
68
34%
17
6
19
6
HOL
PB2
PB1
PB3
PA
F
(PB1 , HOL ; PB3 , PB2)
(PB1 , PA ; PB3 , PB2)
(PB1 , HOL ; PB3 , PA)
(PB1 , PB2 ; PB3 , PA)
(PB1 , PA ; PB3 , HOL)
(PB1 , PB3 ; HOL , PB2)
(PB1 , PB2 ; PB3 , HOL)
(PB1 , PA ; HOL , PB2)
(PB1 , PB3 ; HOL , PA)
(PB1 , PB3 ; PA , PB2)
(PB1 , PB2 ; HOL , PA)
(PB2 , HOL ; PB3 , PA)
(PB2 , PA ; PB3 , HOL)
(PB1 , HOL ; PA , PB2)
(PB2 , PB3 ; HOL , PA)
0
0,0005
0,001
0,015
f4

Figure 5