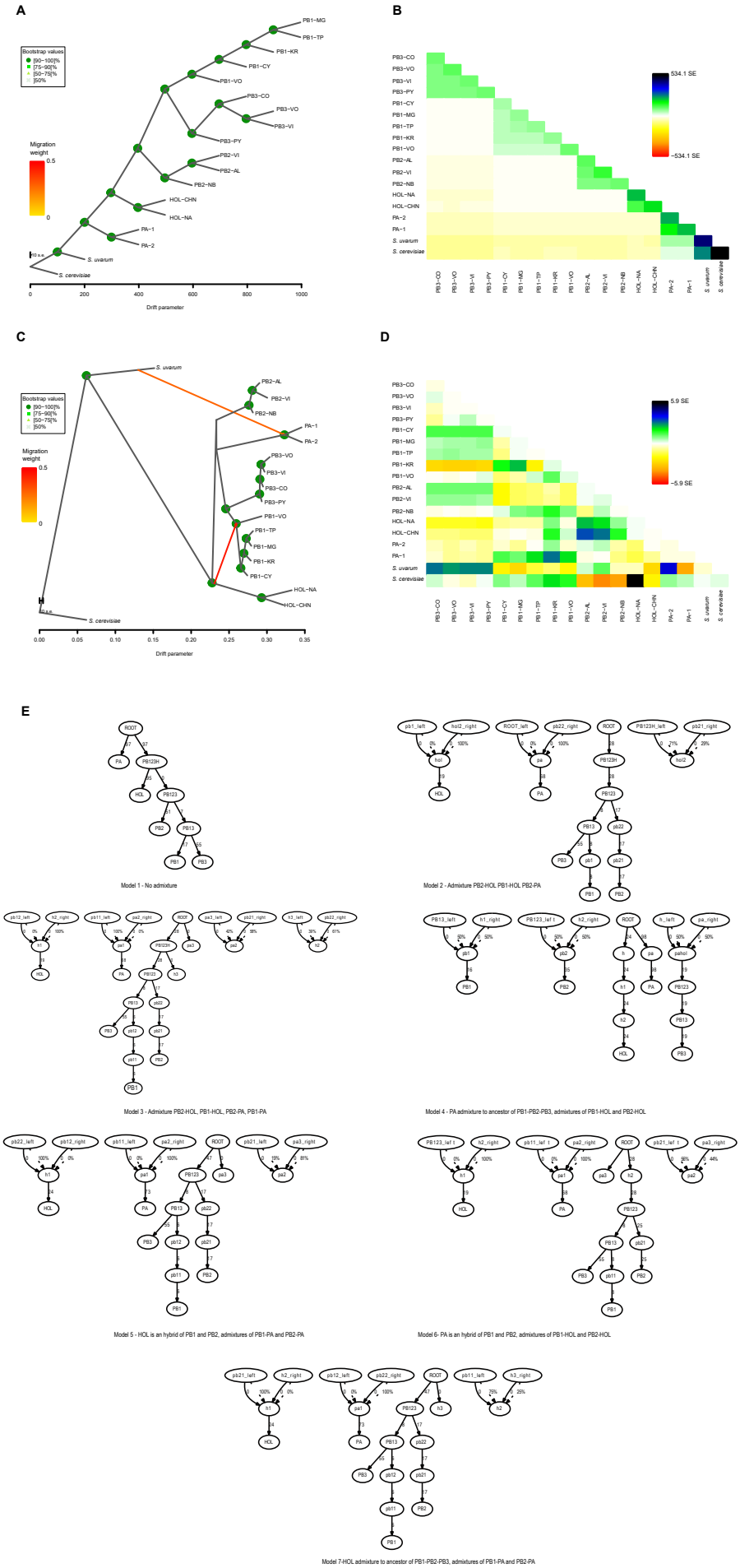
A
Bootstrap values
[90−100]%
[75−90]%
[50−75]%
]50%
Migration weight
0.5
0
PB1−MG
PB1−TP
PB1−KR
PB1−CY
PB1−VO
PB3−CO
PB3−VO
PB3−VI
PB3−PY
PB2−VI
PB2−AL
PB2−NB
HOL−CHN
HOL−NA
PA−1
PA−2
S. uvarum
S. cerevisiae
0
200
400
600
800
1000
Drift parameter
B
534.1 SE
−534.1 SE
PB3−CO
PB3−VO
PB3−VI
PB3−PY
PB1−CY
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
C
Bootstrap values
[90−100]%
[75−90]%
[50−75]%
]50%
Migration weight
0.5
0
S. uvarum
PB2−AL
PB2−VI
PB2−NB
PA−1
PA−2
PB3−VO
PB3−VI
PB3−CO
PB3−PY
PB1−VO
PB1−TP
PB1−MG
PB1−KR
PB1−CY
HOL−NA
HOL−CHN
S. cerevisiae
0.00
0.05
0.10
0.15
0.20
0.25
0.30
0.35
Drift parameter
D
5.9 SE
−5.9 SE
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
E
Model 1 - No admixture
Model 2 - Admixture PB2-HOL PB1-HOL PB2-PA
Model 3 - Admixture PB2-HOL, PB1-HOL, PB2-PA, PB1-PA
Model 4 - PA admixture to ancestor of PB1-PB2-PB3, admixtures of PB1-HOL and PB2-HOL
Model 5 - HOL is an hybrid of PB1 and PB2, admixtures of PB1-PA and PB2-PA
Model 6- PA is an hybrid of PB1 and PB2, admixtures of PB1-HOL and PB2-HOL
Model 7-HOL admixture to ancestor of PB1-PB2-PB3, admixtures of PB1-PA and PB2-PA

Figure 4.