YJL144W
TIM17
YAK1
swap region A limits
known RM/BY variants
new variant site
Allele frequency (reference / total)
1.0
0.8
0.6
0.4
0.2
0.0
Sequenced populations based on strains built from the GFP collection
low protein (n=15)
high protein (n=15)
other populations (n=45)
Sequenced populations based on reconstructed strains (NOT from the GFP collection)
low protein (n=10)
high protein (n=10)
other populations (n=30)
147
148
149
150
Position on chromosome X (kb)