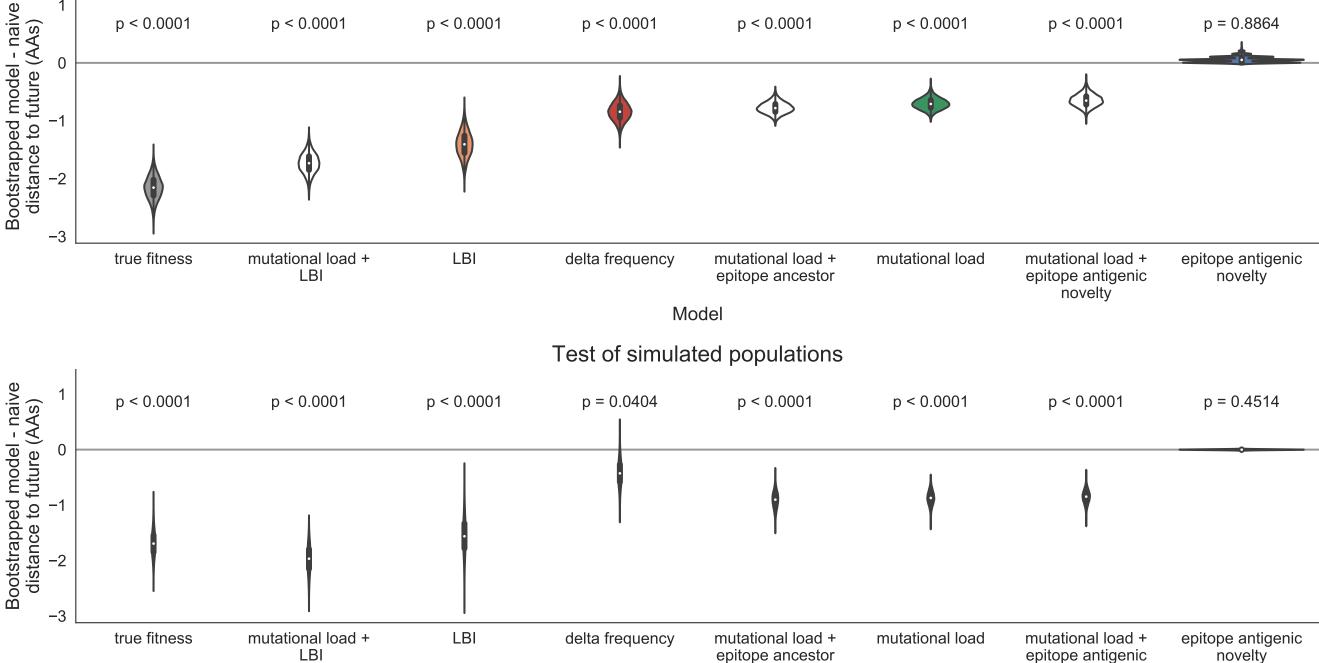
Validation of simulated populations p < 0.0001p < 0.0001p < 0.0001p < 0.0001p < 0.0001p = 0.8864LBI mutational load + mutational load delta frequency mutational load + epitope ancestor

novelty



p < 0.0001

p < 0.0001

Model