

where are we...

updates using pompjax, and the empirical prevalences from the literature.

ABM inferences

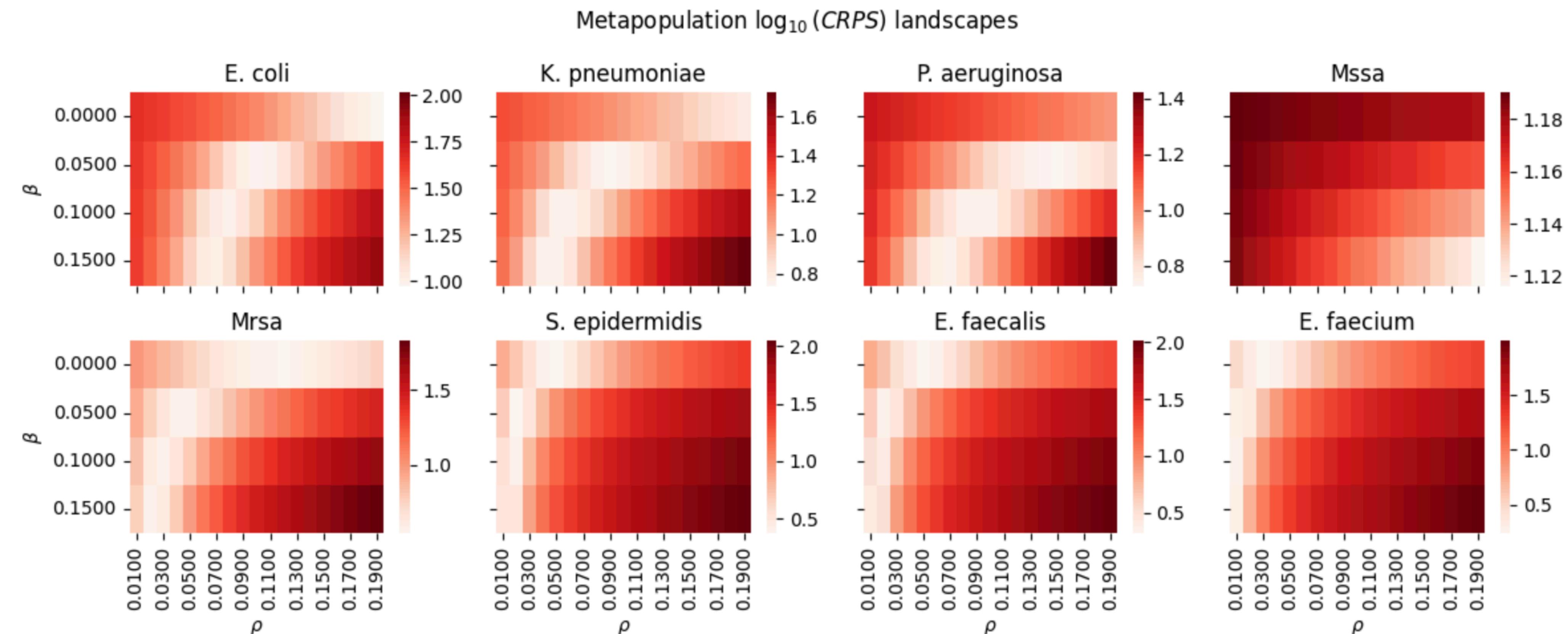
recap

- Process model at daily time scales.
- Same study period as before 01/Feb/2020 to 28/Feb/2021 (weird drop in culture after feb 2021)
- we decided to drop *C. albicans* and include *E. faecalis*, also in the top 15 of microorganism detected.

ABM inferences

Inference on synthetic data.

- as the definition of ρ is a little blurry and hard to find references I did some grid search and computed the fit to the hospital level data to ballpark the range for the different importation rates (AMRO prevalences).



ABM inferences

Inference on synthetic data.

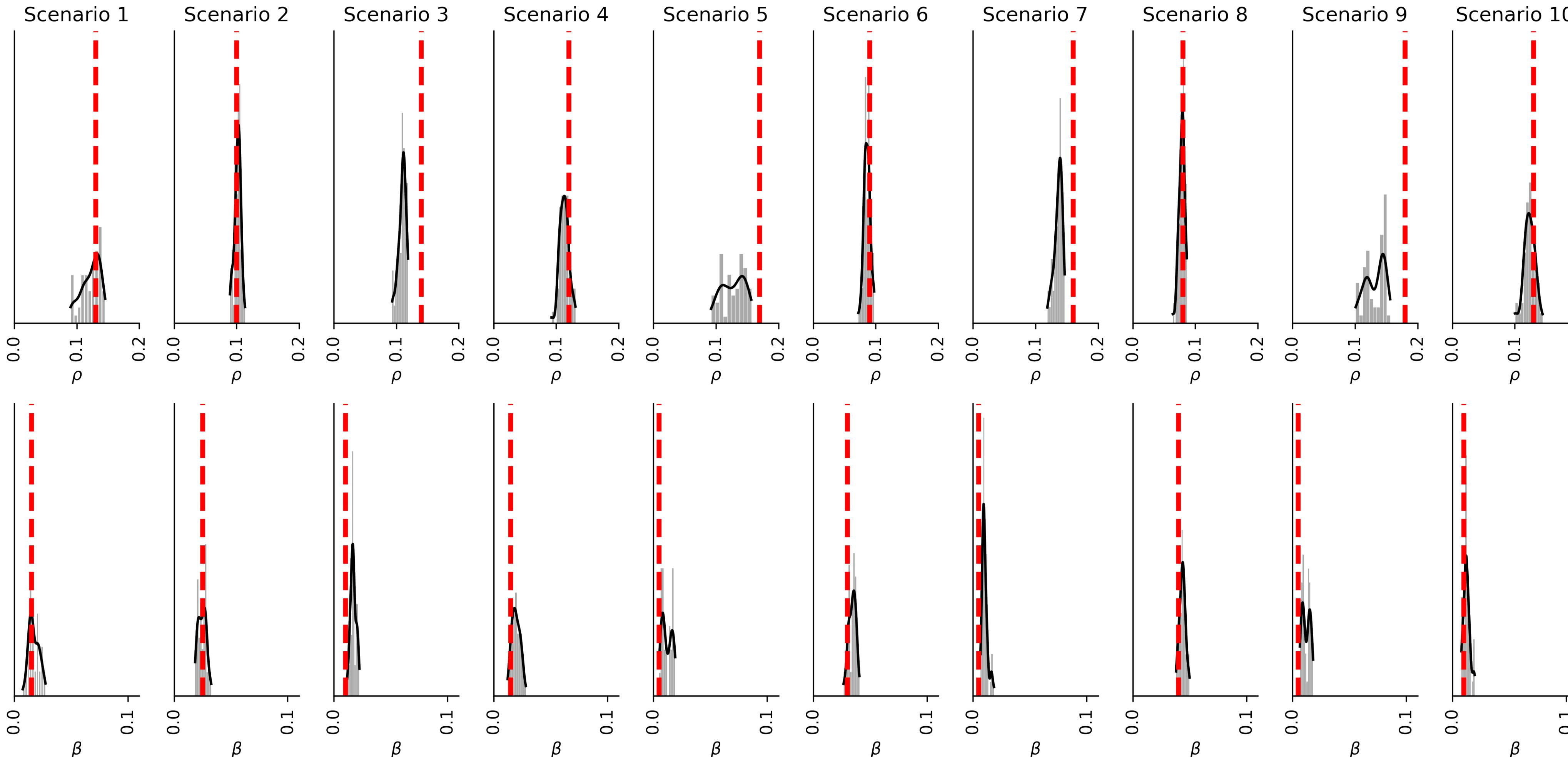
- Using the ranges in the grid search, I selected values that already minimize the CRPS to the hospital-level observations and ran inferences on that.
 - I checked the convergence plots and the inferences seem to be working/converging.
 - The empirical prevalences range from 0.4% (MRSA), to 60.3% (E. coli), so in general, we are covering a huge range of importations from all the AMRO to test in synthetic inferences.
 - Only MRSA and *P. aeruginosa* have importations lower than 15%
 - **Results:** in general it seems that the inference is working very well (truths are falling in the posterior - even shrinking the variance). However, for low importation rates (lower than 15%) the system struggles in both β and ρ .
 - Next slides shows the posterior for the 10 scenarios for all the AMROs. Prior ranges for the inference are shown as the limit of the x-axis.
 - **Next:** If we want to report CI maybe I could remove the shrinking between iterations and see how those posterior look like.

ABM inferences on synthetic data

Inferences on simulated observations

Two observational models;

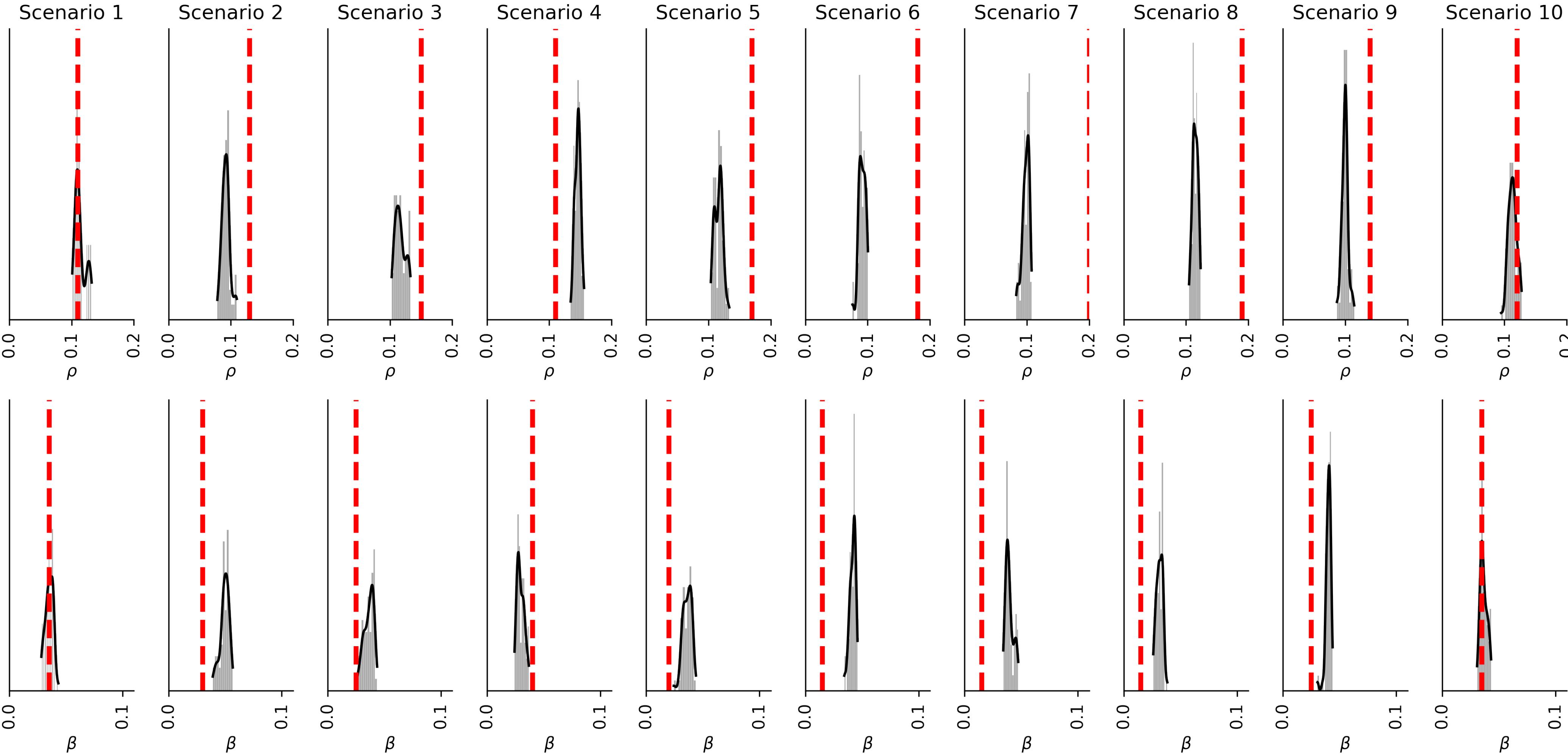
Posteriors for E. coli, $\gamma = 63.0\%$



Inferences on simulated observations

Two observational models;

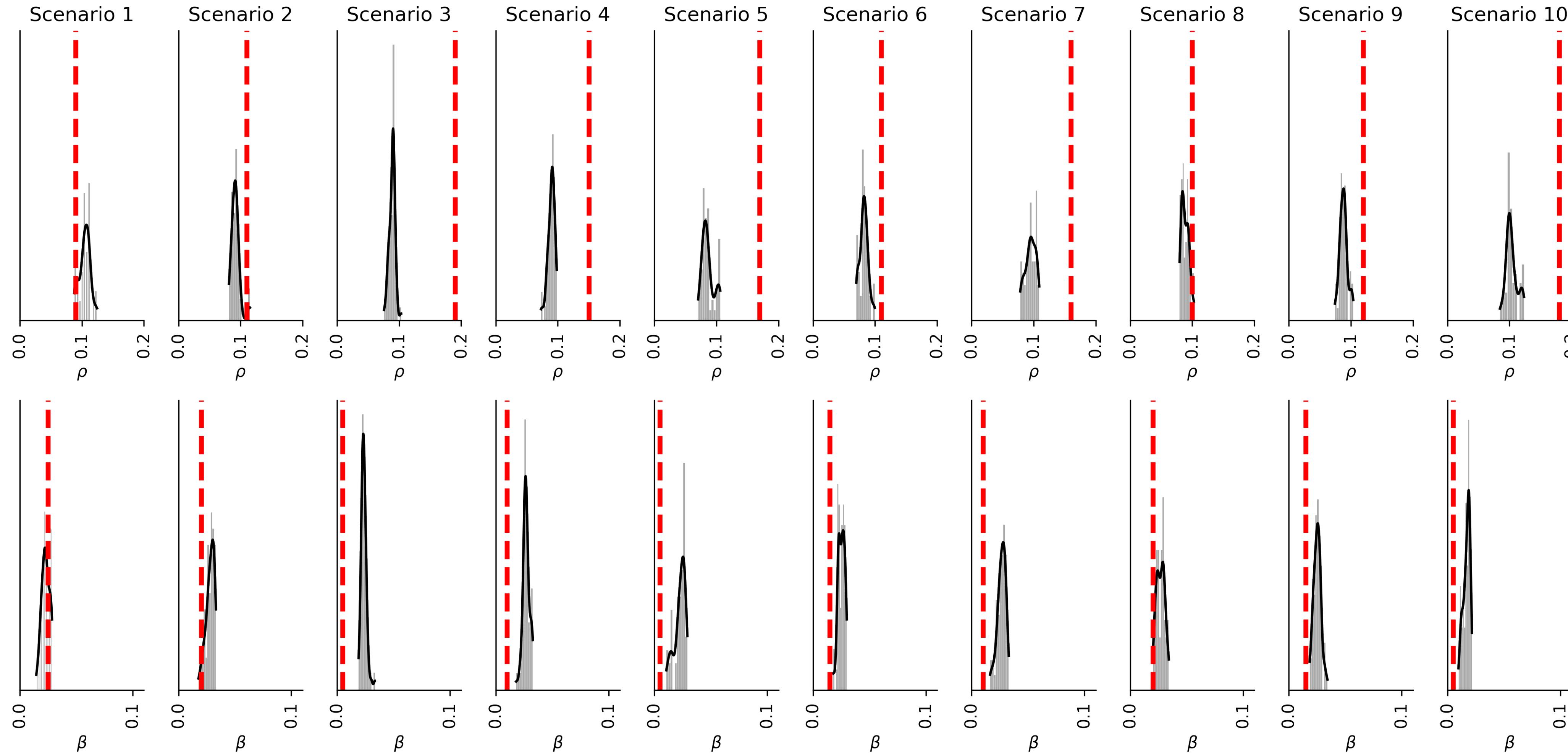
Posteriors for *P. aeruginosa*, $\gamma = 11.6\%$



Inferences on simulated observations

Two observational models;

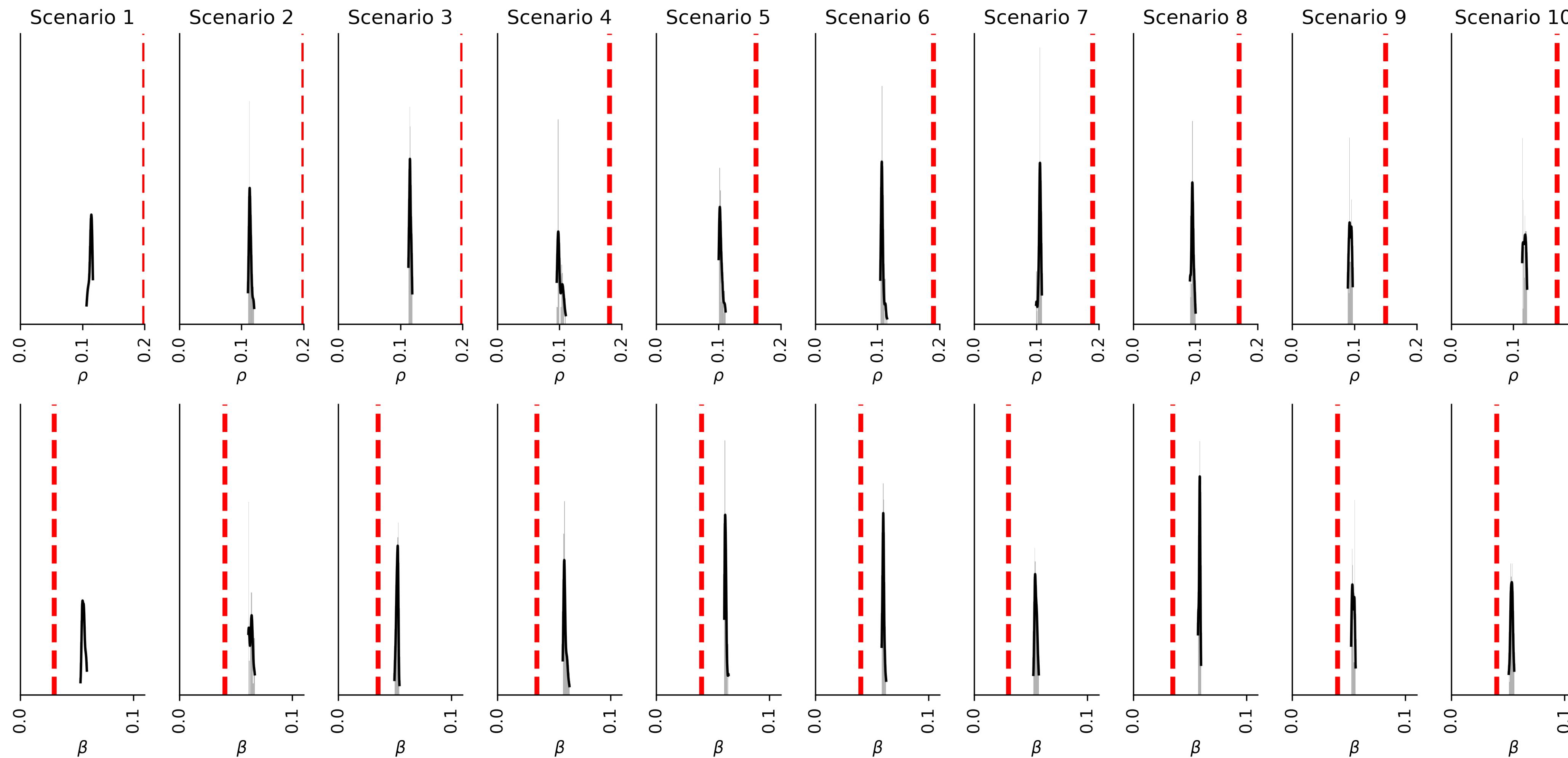
Posteriors for $K. pneumoniae$, $\gamma = 23.0\%$



Inferences on simulated observations

Two observational models;

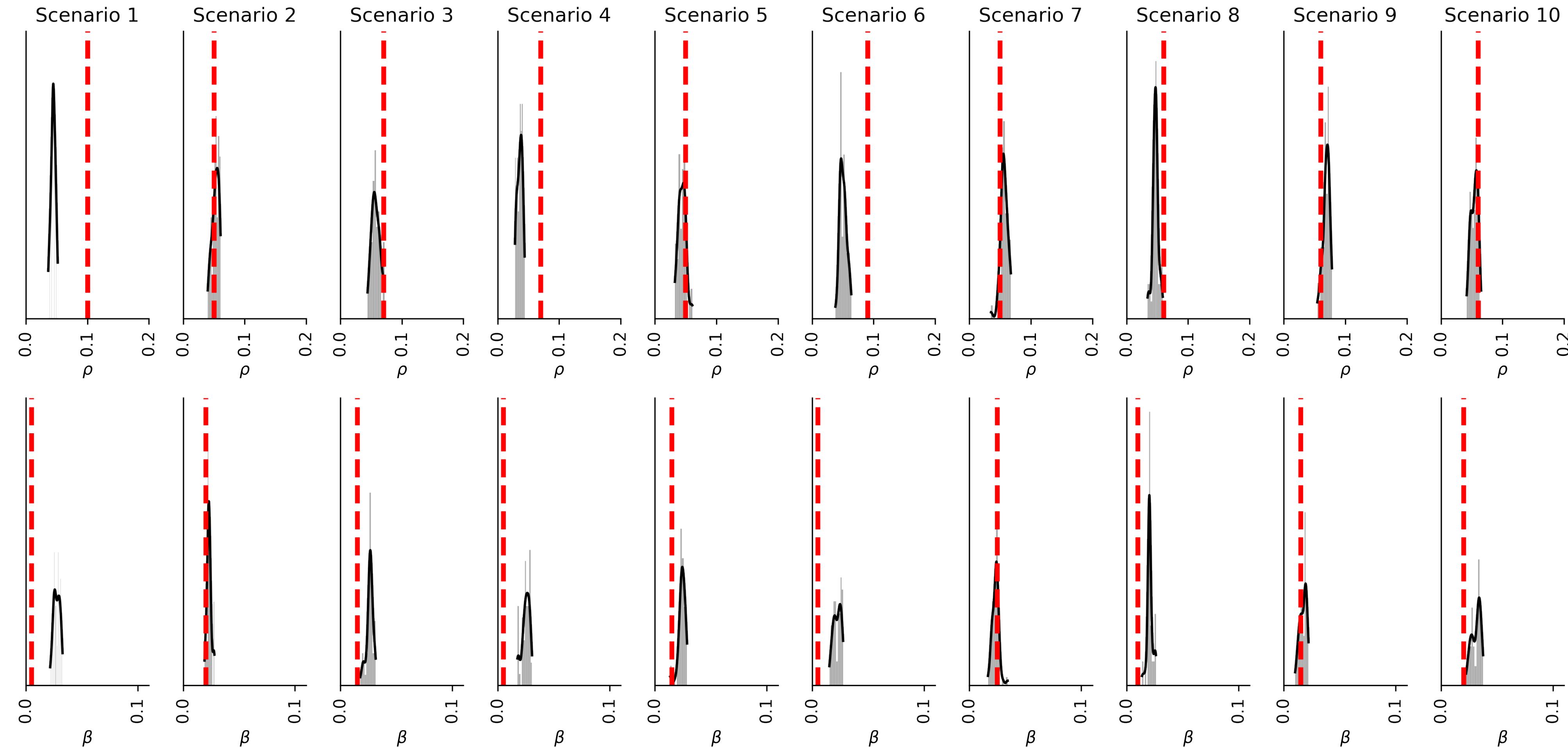
Posteriors for MSSA, $\gamma = 0.4\%$



Inferences on simulated observations

Two observational models;

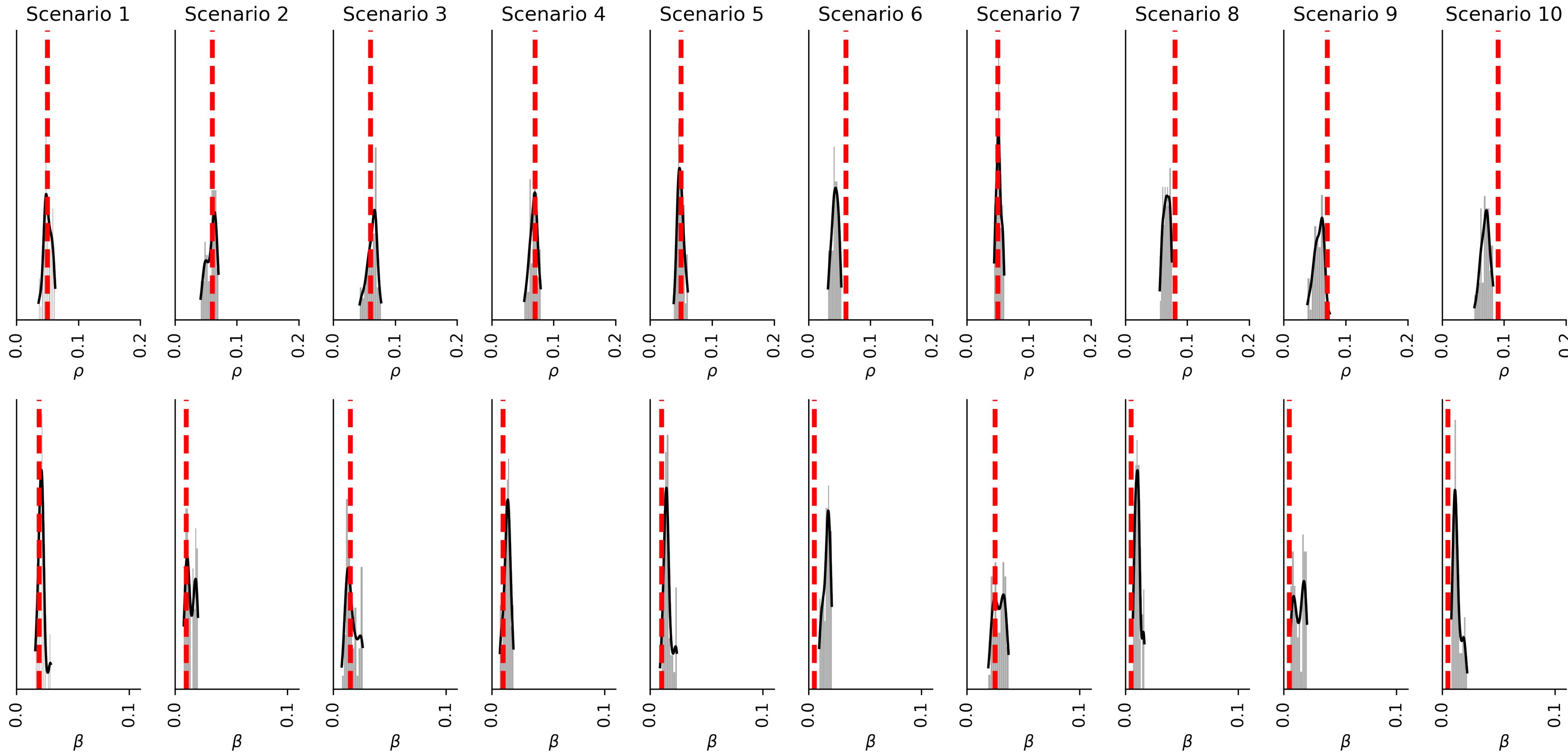
Posteriors for MRSA, $\gamma = 25.0\%$



Inferences on simulated observations

Two observational models;

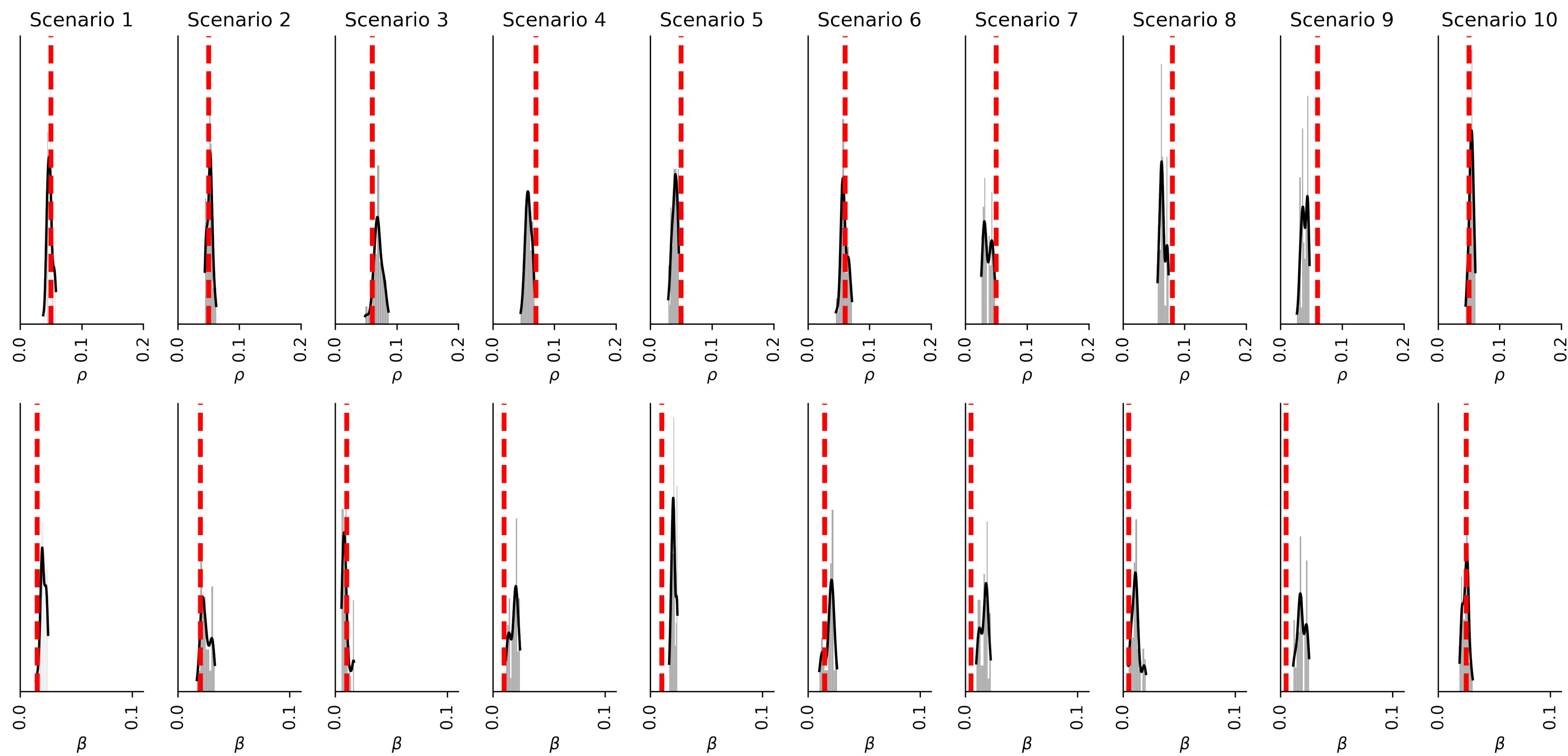
Posteriors for *S. epidermidis*, $\gamma = 58.0\%$



Inferences on simulated observations

Two observational models;

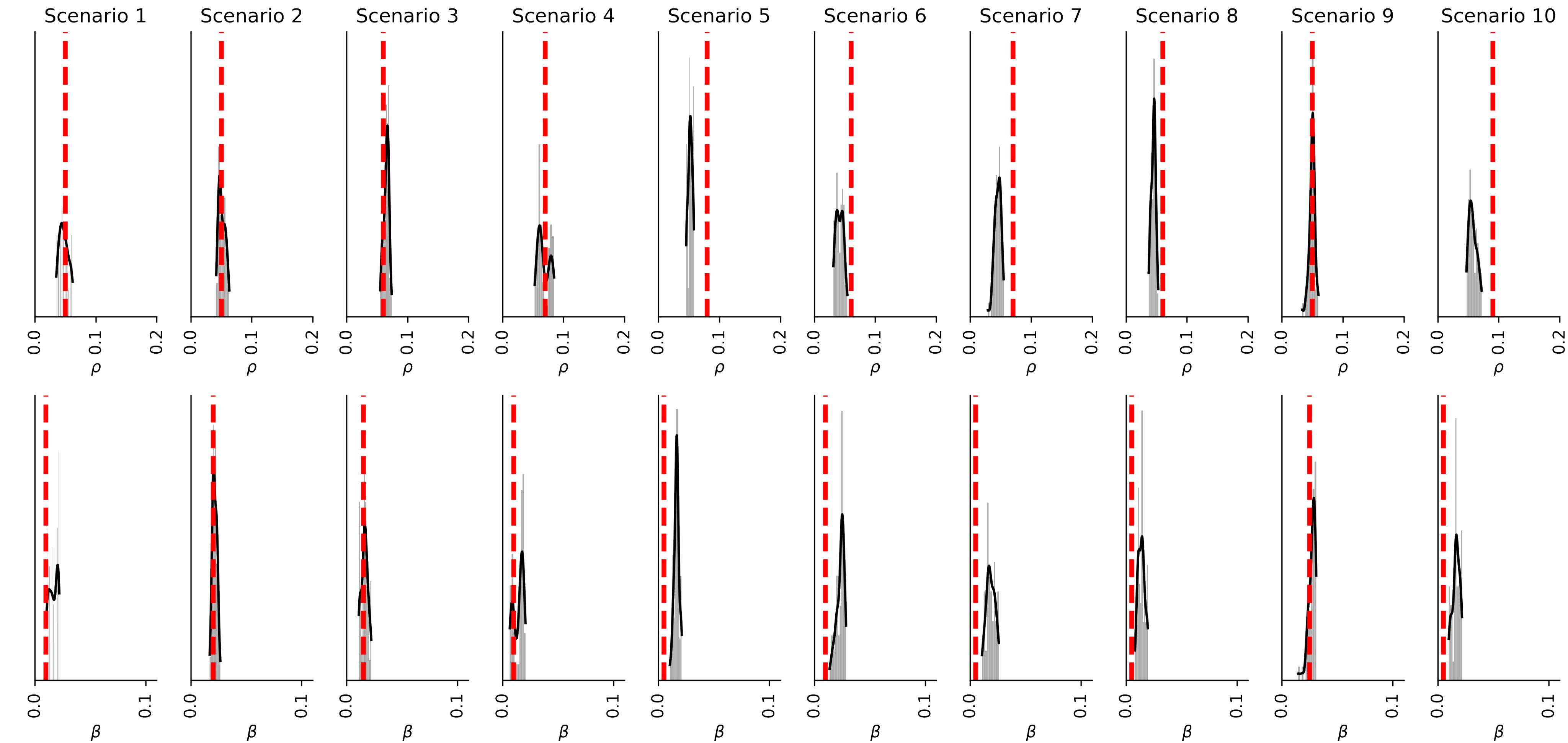
Posteriors for $E. faecalis$, $\gamma = 47.6\%$



Inferences on simulated observations

Two observational models;

Posteriors for E. faecium, $\gamma = 40.6\%$

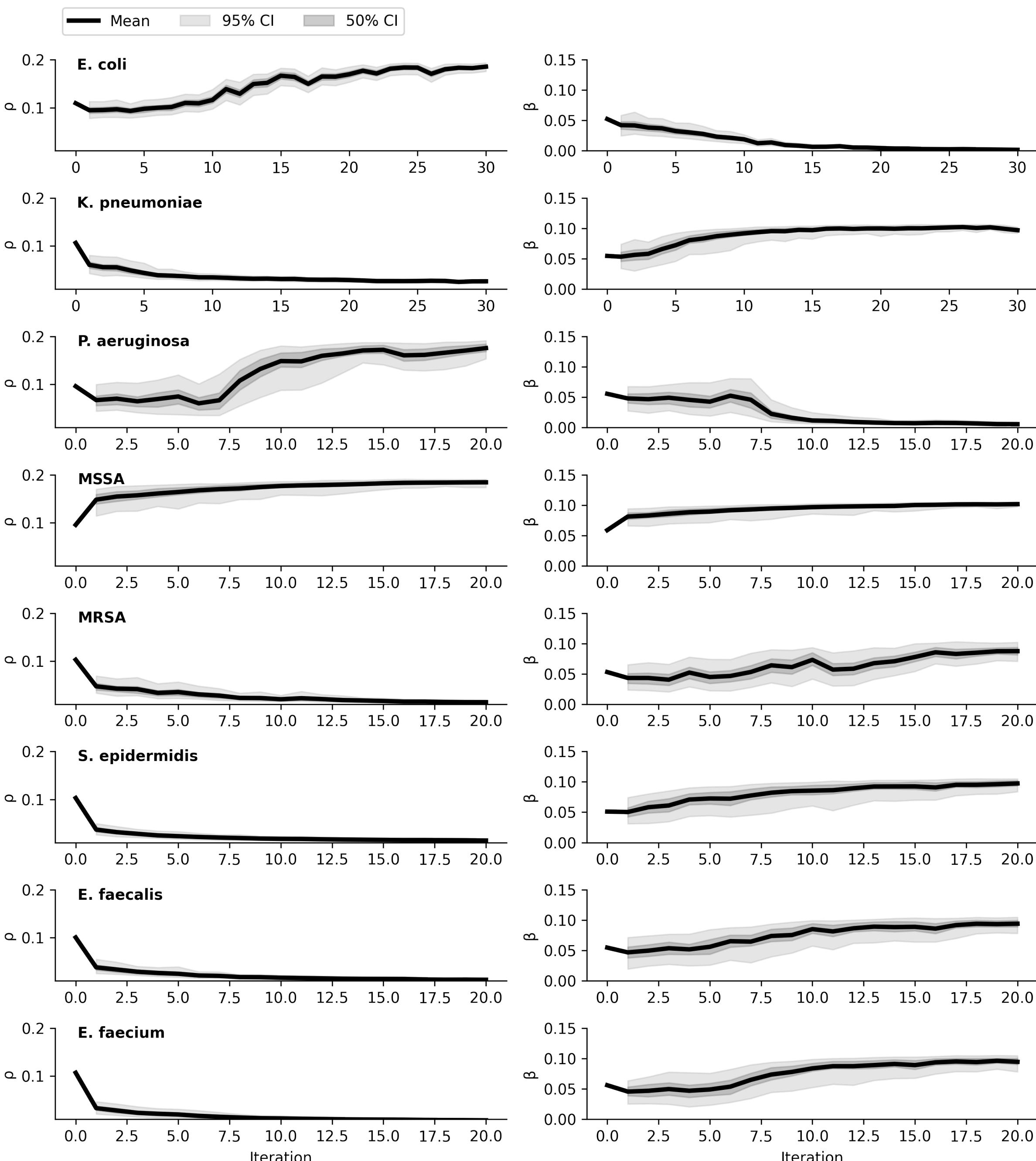


ABM inferences on real data

ABM inferences

Convergence plots - Individual level observational model

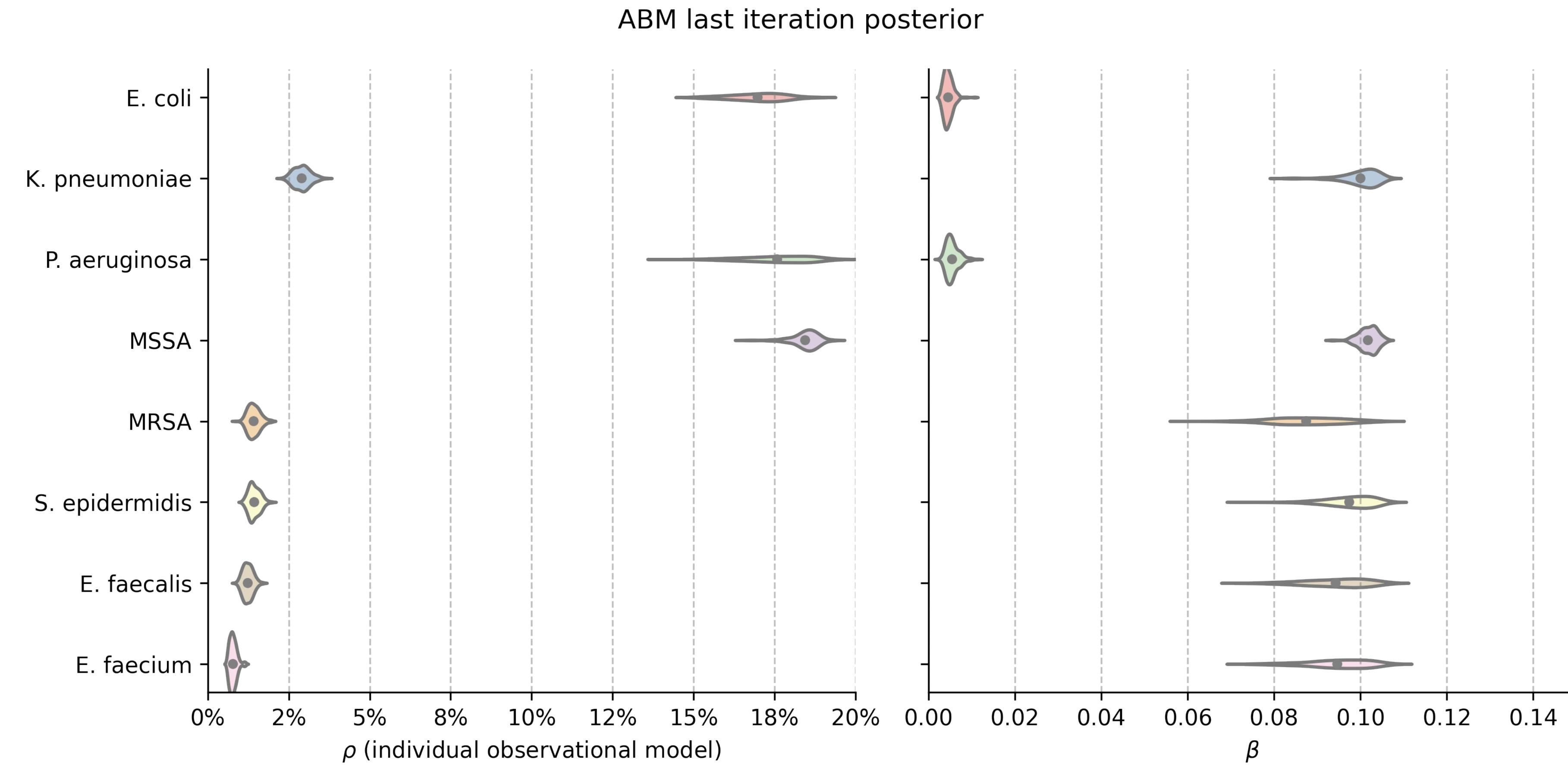
- Visually inspect convergence plots to see... convergence.
 - Convergences plots consistently look to asymptotically go to a specific value.



ABM inferences

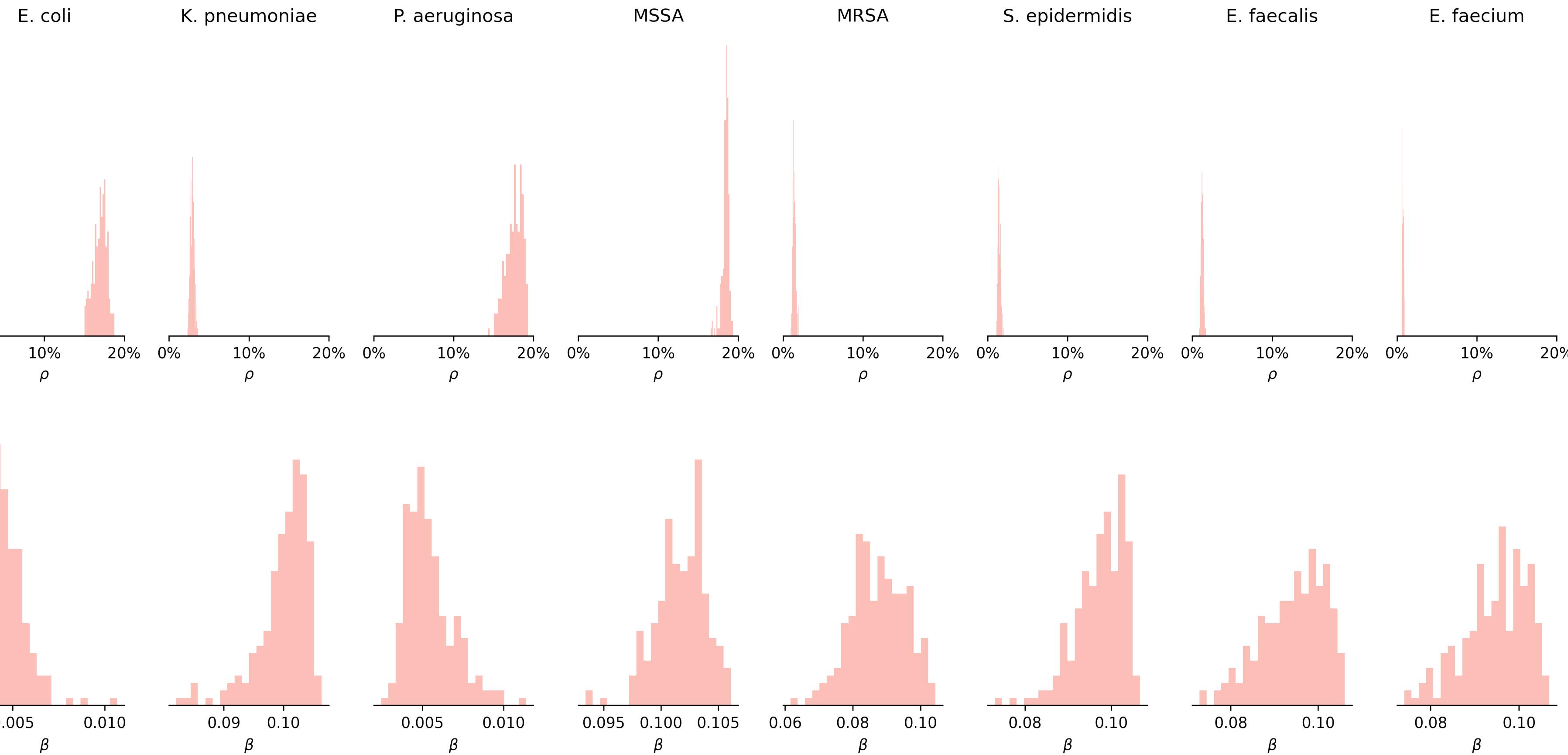
Posterior inferences

- Effective sensitivity around 1% for MRSA, *S. epidermidis*, *E. faecalis*, *E. Faecium*.
- 2.1% for *K. Pneumonia*
- Around 18% for *E. coli*, *P. aeruginosa* and MSSA



ABM inferences

Posterior joint distributions and marginals



- Just another way to viz the posterior inferences.