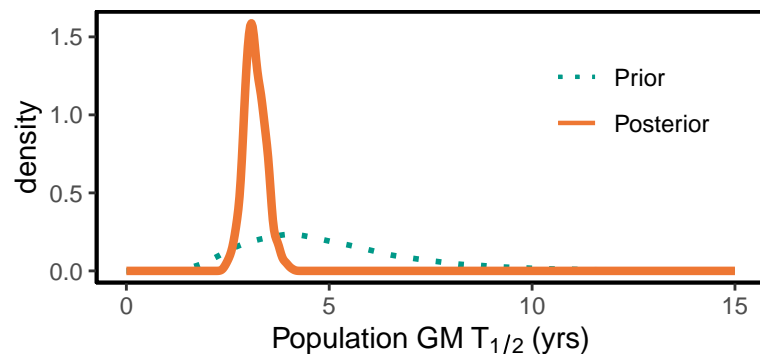


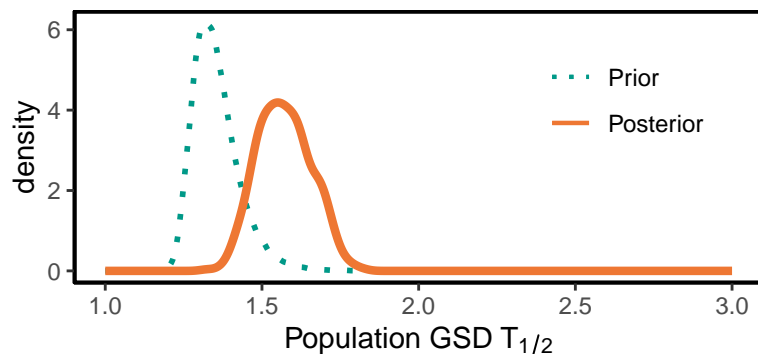
A: PFOA  $T_{1/2}$  Population GM

Posterior Median (95% CI): 3.14 (2.69–3.73)



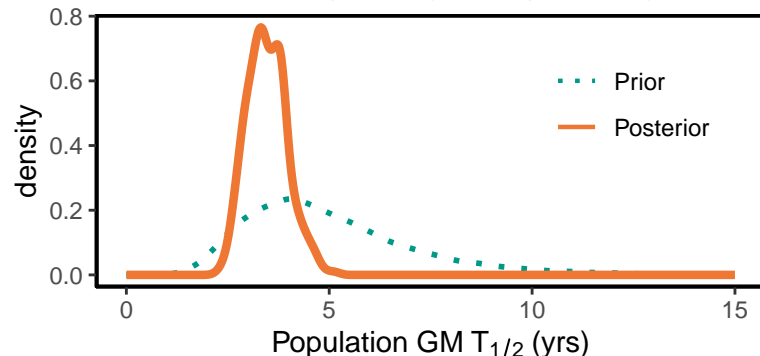
B: PFOA  $T_{1/2}$  Population GSD

Posterior Median (95% CI): 1.57 (1.42–1.73)



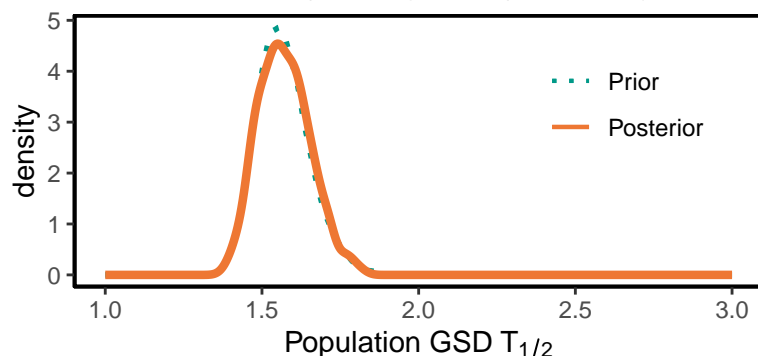
C: PFOS  $T_{1/2}$  Population GM

Posterior Median (95% CI): 3.42 (2.62–4.5)



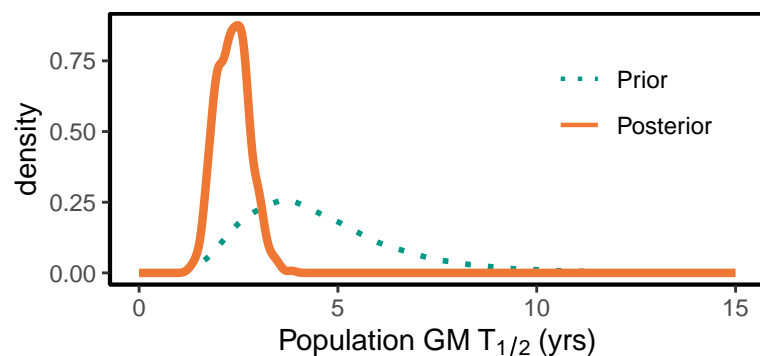
D: PFOS  $T_{1/2}$  Population GSD

Posterior Median (95% CI): 1.57 (1.43–1.75)



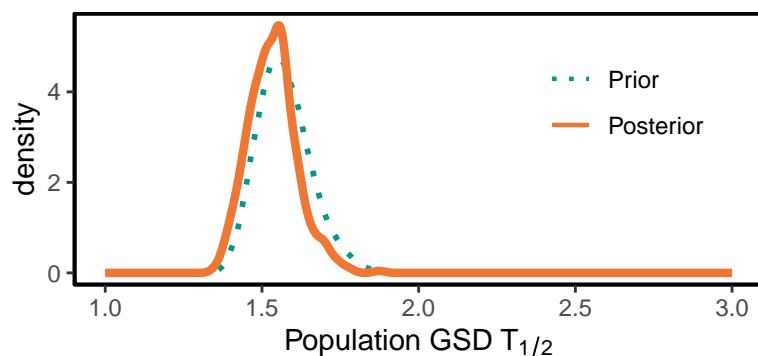
E: PFNA  $T_{1/2}$  Population GM

Posterior Median (95% CI): 2.35 (1.65–3.16)



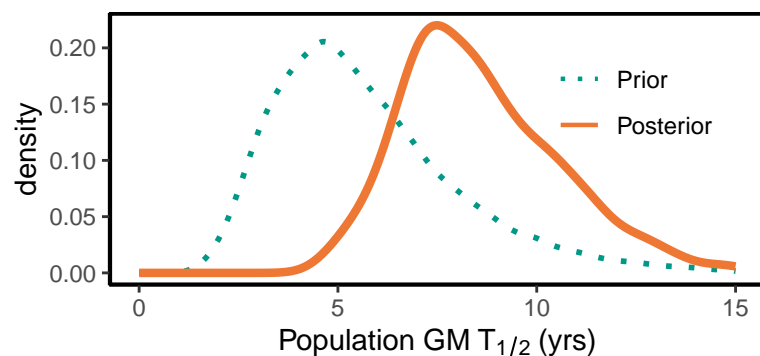
F: PFNA  $T_{1/2}$  Population GSD

Posterior Median (95% CI): 1.53 (1.4–1.7)



G: PFHxS  $T_{1/2}$  Population GM

Posterior Median (95% CI): 8.3 (5.38–13.5)



H: PFHxS  $T_{1/2}$  Population GSD

Posterior Median (95% CI): 1.57 (1.42–1.77)

