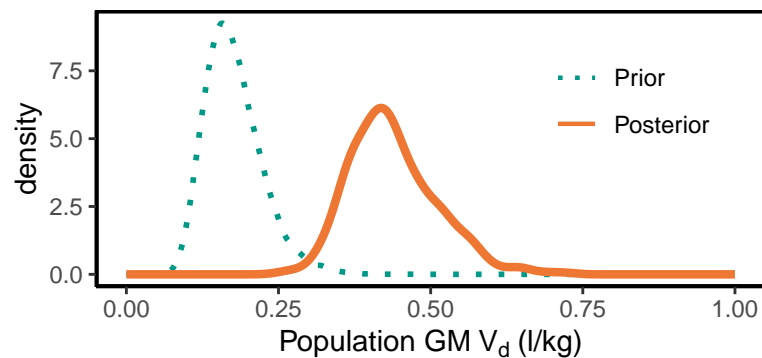


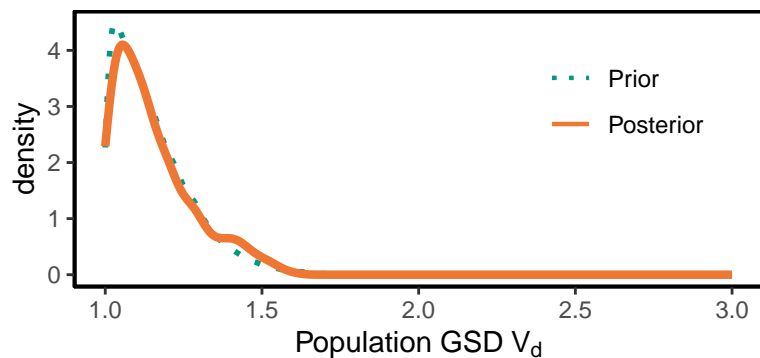
A: PFOA V_d Population GM

Posterior Median (95% CI): 0.428 (0.322–0.593)



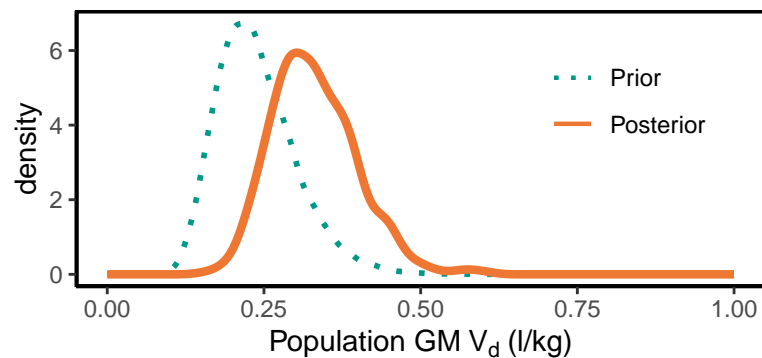
B: PFOA V_d Population GSD

Posterior Median (95% CI): 1.12 (1.01–1.47)



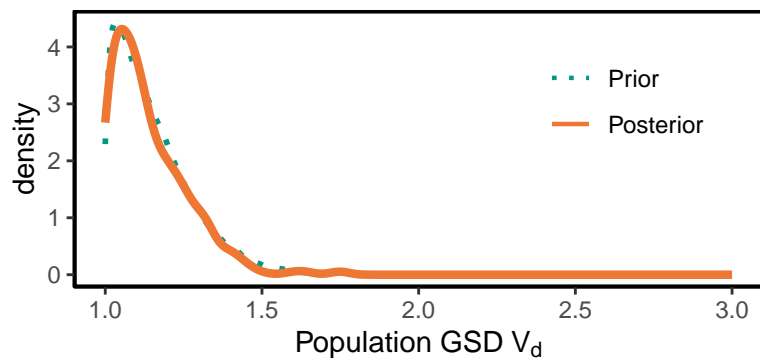
C: PFOS V_d Population GM

Posterior Median (95% CI): 0.322 (0.221–0.47)



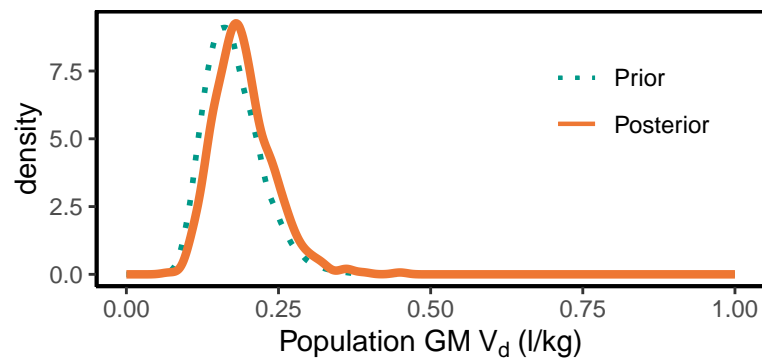
D: PFOS V_d Population GSD

Posterior Median (95% CI): 1.11 (1–1.41)



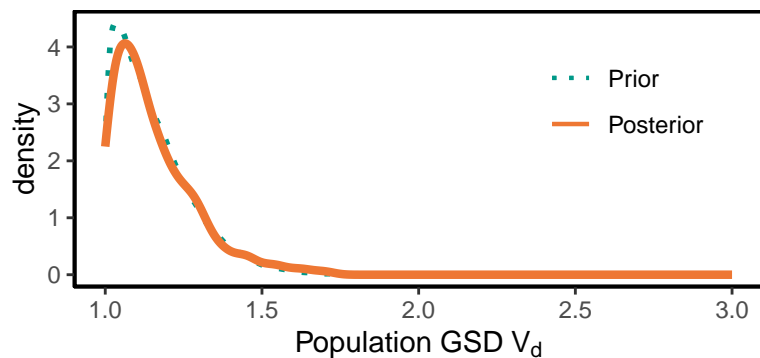
E: PFNA V_d Population GM

Posterior Median (95% CI): 0.186 (0.113–0.302)



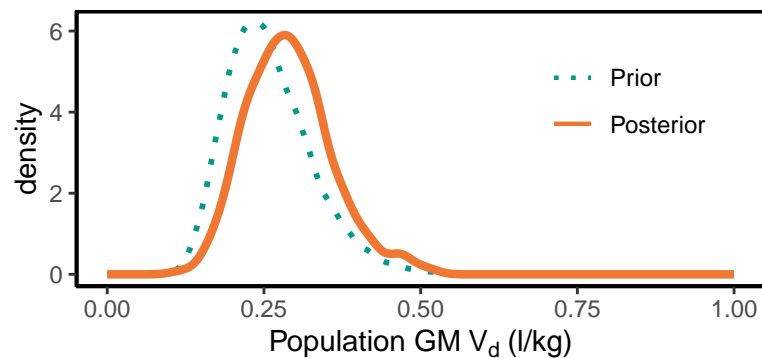
F: PFNA V_d Population GSD

Posterior Median (95% CI): 1.12 (1.01–1.51)



G: PFHxS V_d Population GM

Posterior Median (95% CI): 0.286 (0.173–0.456)



H: PFHxS V_d Population GSD

Posterior Median (95% CI): 1.11 (1–1.45)

