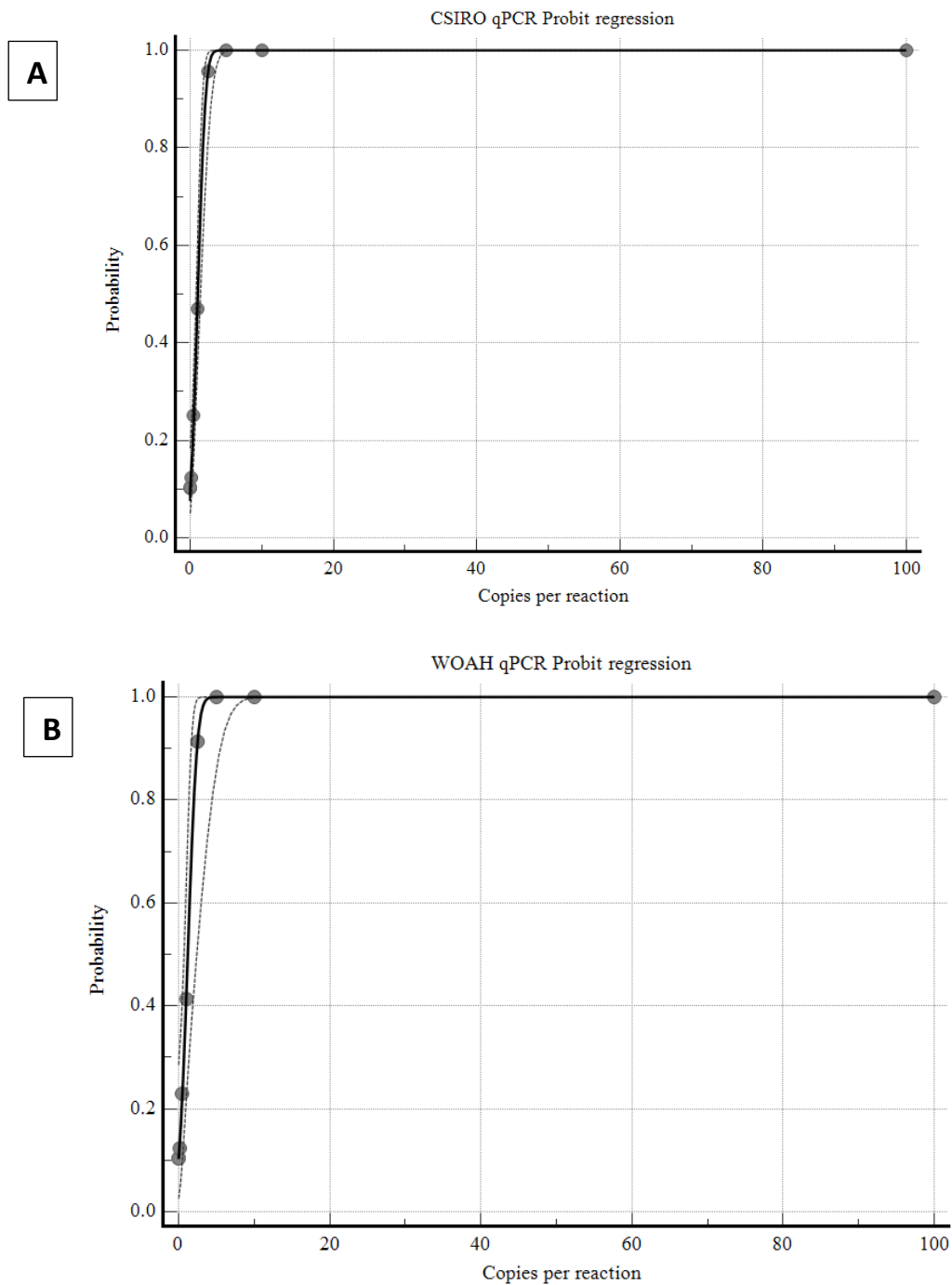


Table S1. Bayesian latent class model outputs for three conditionally dependent tests in three defined populations for WSSV ( $M_0$ , saturated model) including sensitivity analysis results.

| <b>Assay</b>               | <b>Parameter</b>  | <b>Posterior median (95% CrI)<br/>Informative priors (Table 1)</b> | <b>Sensitivity analysis 1<br/>Flat Se1, Sp1, Se2, Sp2</b> | <b>Sensitivity analysis 2<br/>Flat Se1, Sp1, Se2, Sp2<br/>Prev1, Prev2, Prev3</b> |
|----------------------------|-------------------|--|---|---|
| <b>Test A (WOAH qPCR)</b>  | Sensitivity (Se1) | 0.772 (0.718, 0.821)   | 0.807 (0.740, 0.863)                                      | 0.806 (0.739, 0.862)  |
|                            | Specificity (Sp1) | 0.996 (0.987, 0.999)   | 0.972 (0.919, 0.999)                                      | 0.974 (0.915, 0.999)  |
| <b>Test B (CSIRO qPCR)</b> | Sensitivity (Se2) | 0.916 (0.877, 0.947)   | 0.967 (0.926, 0.994)                                      | 0.967 (0.926, 0.992)  |
|                            | Specificity (Sp2) | 0.995 (0.984, 0.999)   | 0.974 (0.919, 0.998)                                      | 0.975 (0.921, 0.998)  |
| <b>Test C (SMP)</b>        | Sensitivity (Se3) | 0.949 (0.898, 0.981)   | 0.976 (0.933, 1.000)                                      | 0.974 (0.934, 0.999)  |
|                            | Specificity (Sp3) | 0.994 (0.965, 1.000)   | 0.992 (0.952, 1.000)                                      | 0.992 (0.956, 1.000)  |
| <b>Inferred prevalence</b> | POP1 (Prev1)      | 0.531 (0.433, 0.627)   | 0.526 (0.430, 0.622)                                      | 0.525 (0.428, 0.620)  |
|                            | POP2 (Prev2)      | 0.371 (0.280, 0.468)   | 0.360 (0.271, 0.459)                                      | 0.359 (0.270, 0.453)  |
|                            | POP3 (Prev3)      | 0.905 (0.827, 0.957)   | 0.885 (0.798, 0.947)                                      | 0.897 (0.808, 0.955)  |
| <b>Test correlation</b>    | $\rho_{D^+,12}$   | 0.370 (0.160, 0.538)   | 0.263 (0.062, 0.460)                                      | 0.258 (0.065, 0.450)  |
|                            | $\rho_{D^+,13}$   | 0.168 (0.018, 0.349)   | 0.200 (−0.062, 0.495)                                     | 0.200 (−0.026, 0.450)   |
|                            | $\rho_{D^+,23}$   | 0.215 (0.012, 0.550)   | 0.126 (−0.450, 0.635)                                     | 0.130 (−0.275, 0.577)   |
|                            | $\rho_{D^-,12}$   | 0.490 (−0.811, 0.978)  | 0.631 (−0.259, 0.975)                                     | 0.641 (−0.324, 0.975)   |
|                            | $\rho_{D^-,13}$   | 0.235 (−0.883, 0.964)  | 0.286 (−0.763, 0.948)                                     | 0.297 (−0.717, 0.953)   |
|                            | $\rho_{D^-,23}$   | 0.056 (−0.944, 0.948)  | −0.088 (−0.951, 0.912)                                    | −0.102 (−0.951, 0.915)  |



(continued on next page)

Figure S1. Limit of detection (LOD) analysis results shown for CSIRO and WOH qPCR assays (A and B), and SMP WSSV (C). LOD results for (A) CSIRO qPCR is 2.4 copies per reaction, (B) WOH qPCR is 2.8 copies per reaction, and SMP WSSV is 2.9 copies per reaction. Twenty replicates were tested at each concentration (A + B) and four replicates tested at each concentration (C + D). Titration points assessed: (A + B) 100 / 10 / 5 / 2.5 / 1 / 0.5 / 0.1 / 0.01, and (C + D) 500 / 250 / 125 / 64 / 62.5 / 16 / 4 / 1 / 0.25.

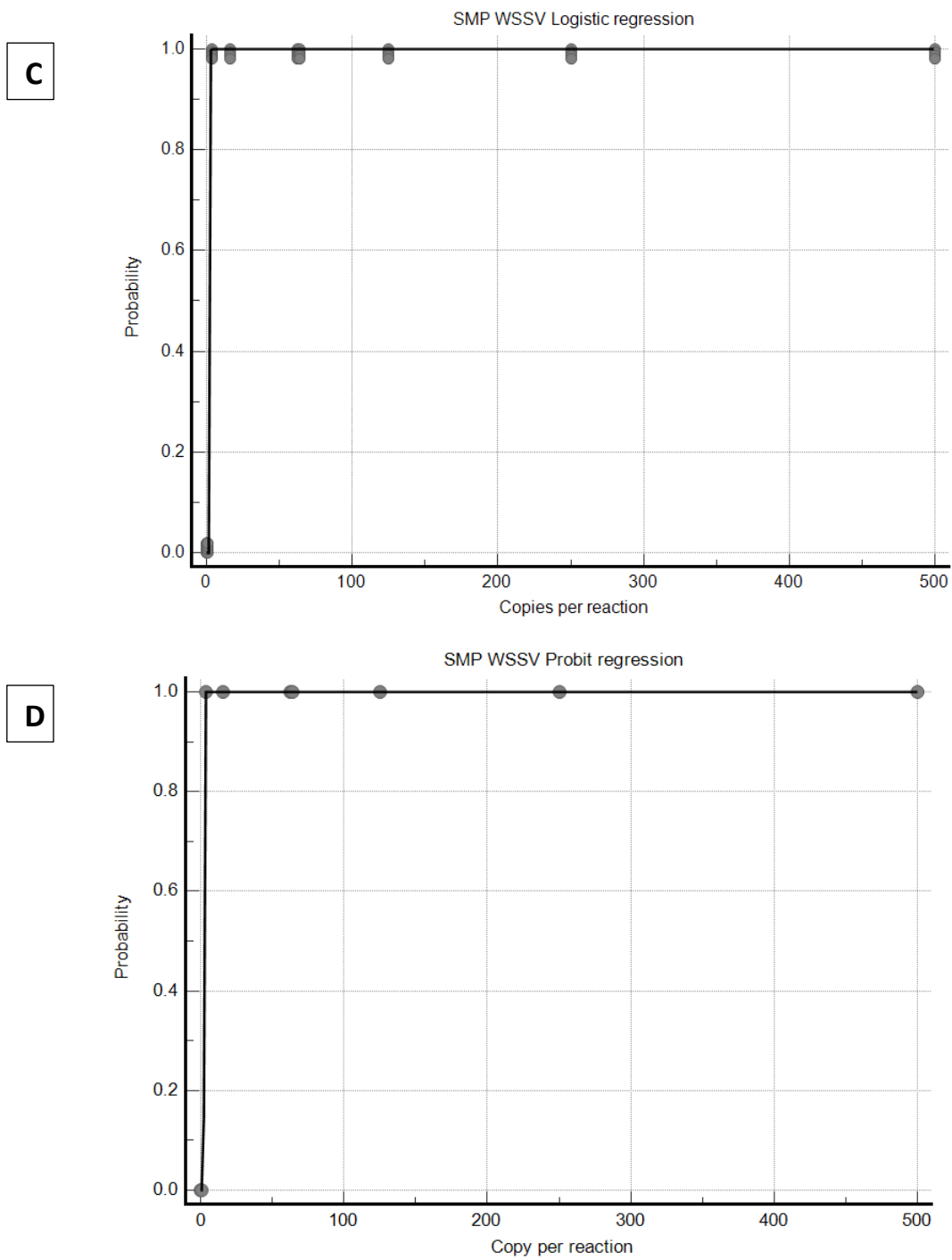


Figure S1. (continued)