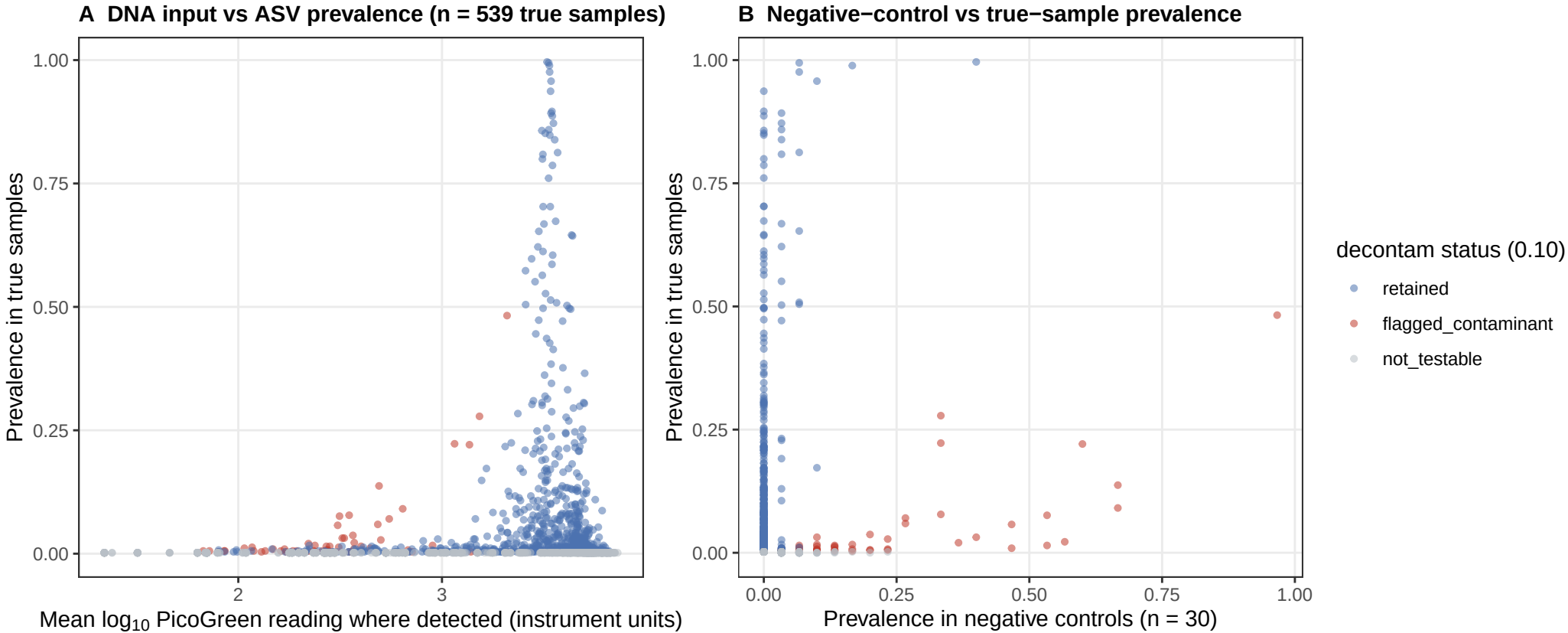
A DNA input vs ASV prevalence (n = 539 true samples)
B Negative−control vs true−sample prevalence
Prevalence in true samples
1.00
0.75
0.50
0.25
0.00
2
3
Mean log10 PicoGreen reading where detected (instrument units)
0.00
0.25
0.50
0.75
1.00
Prevalence in negative controls (n = 30)
decontam status (0.10)
retained
flagged_contaminant
not_testable