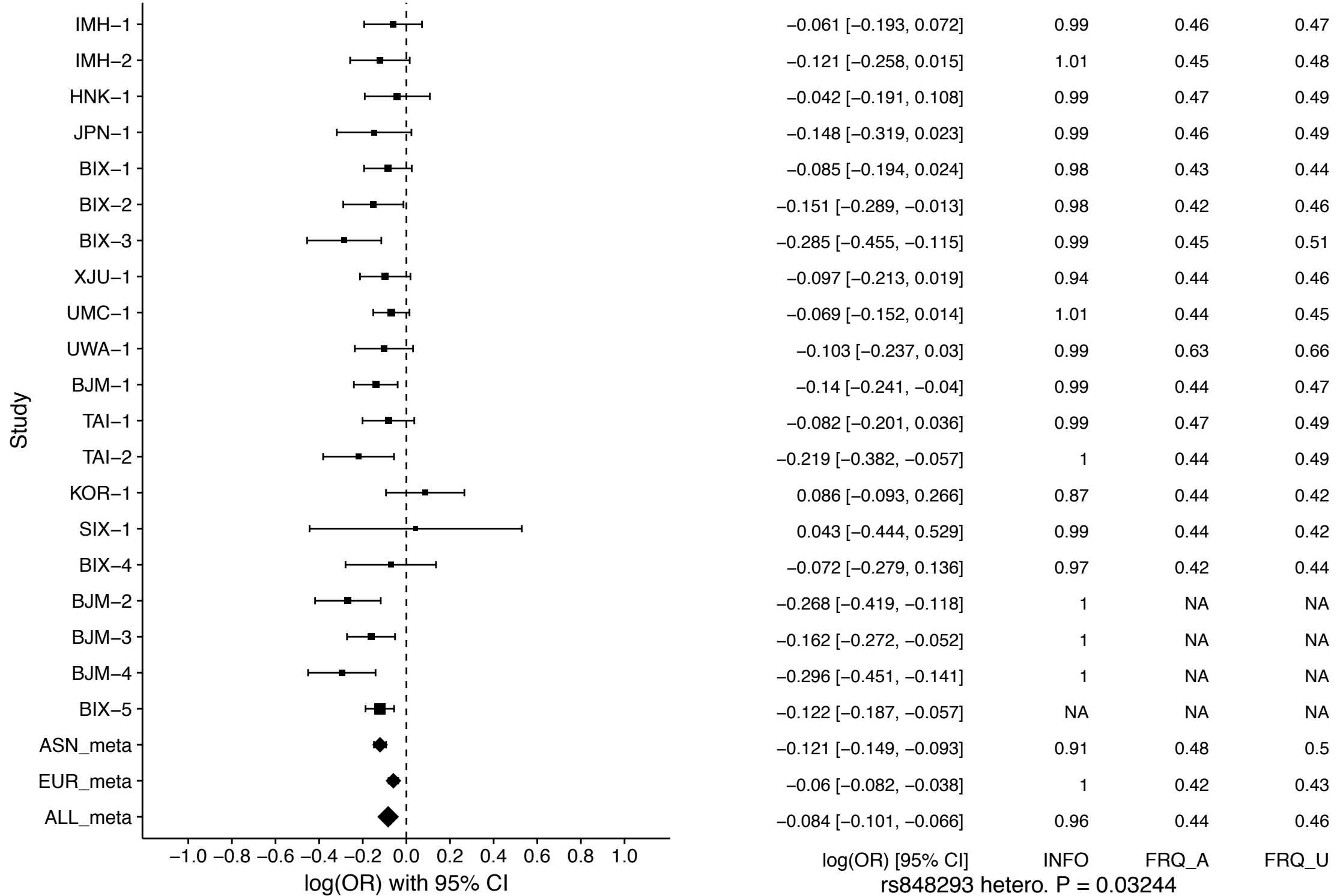
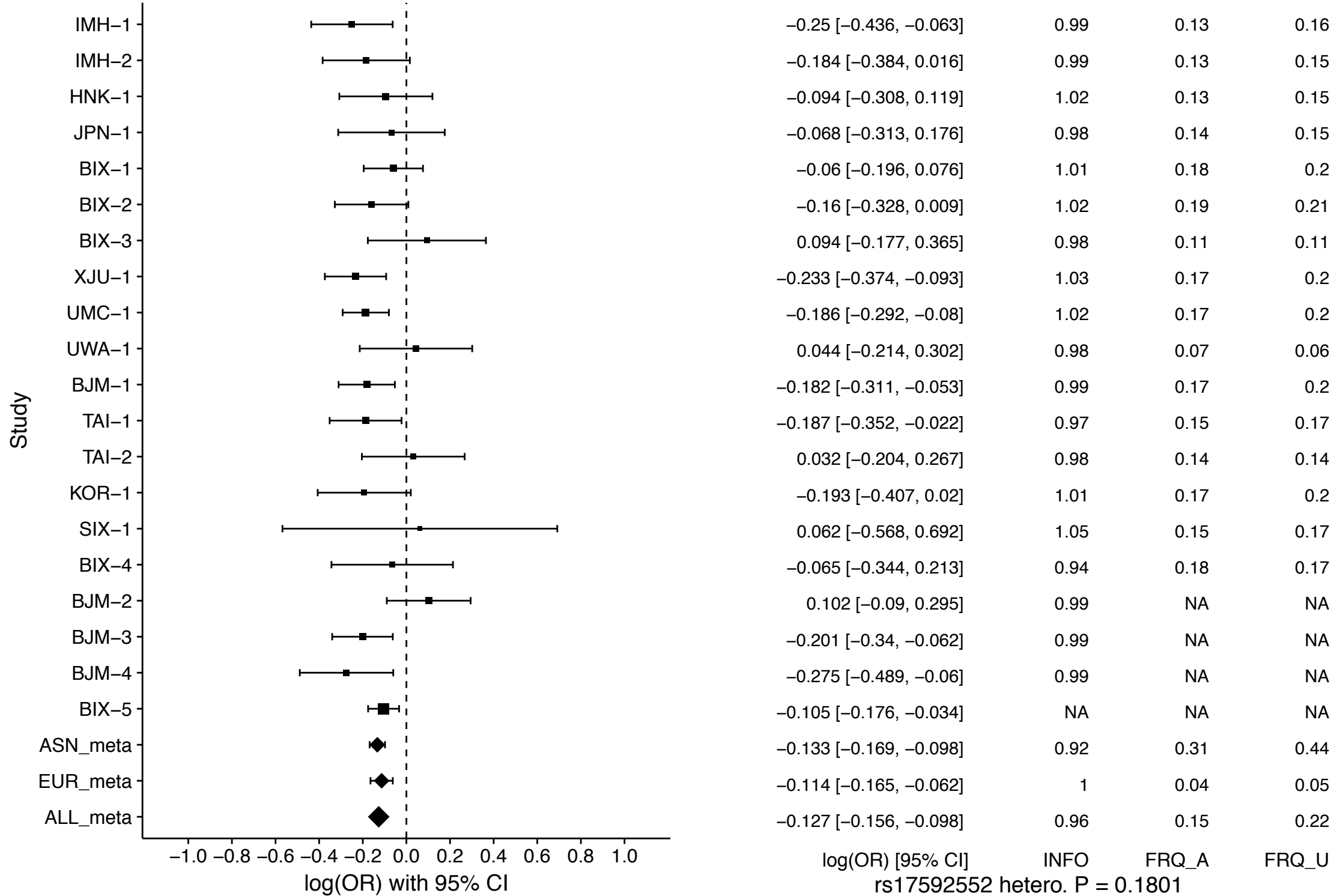
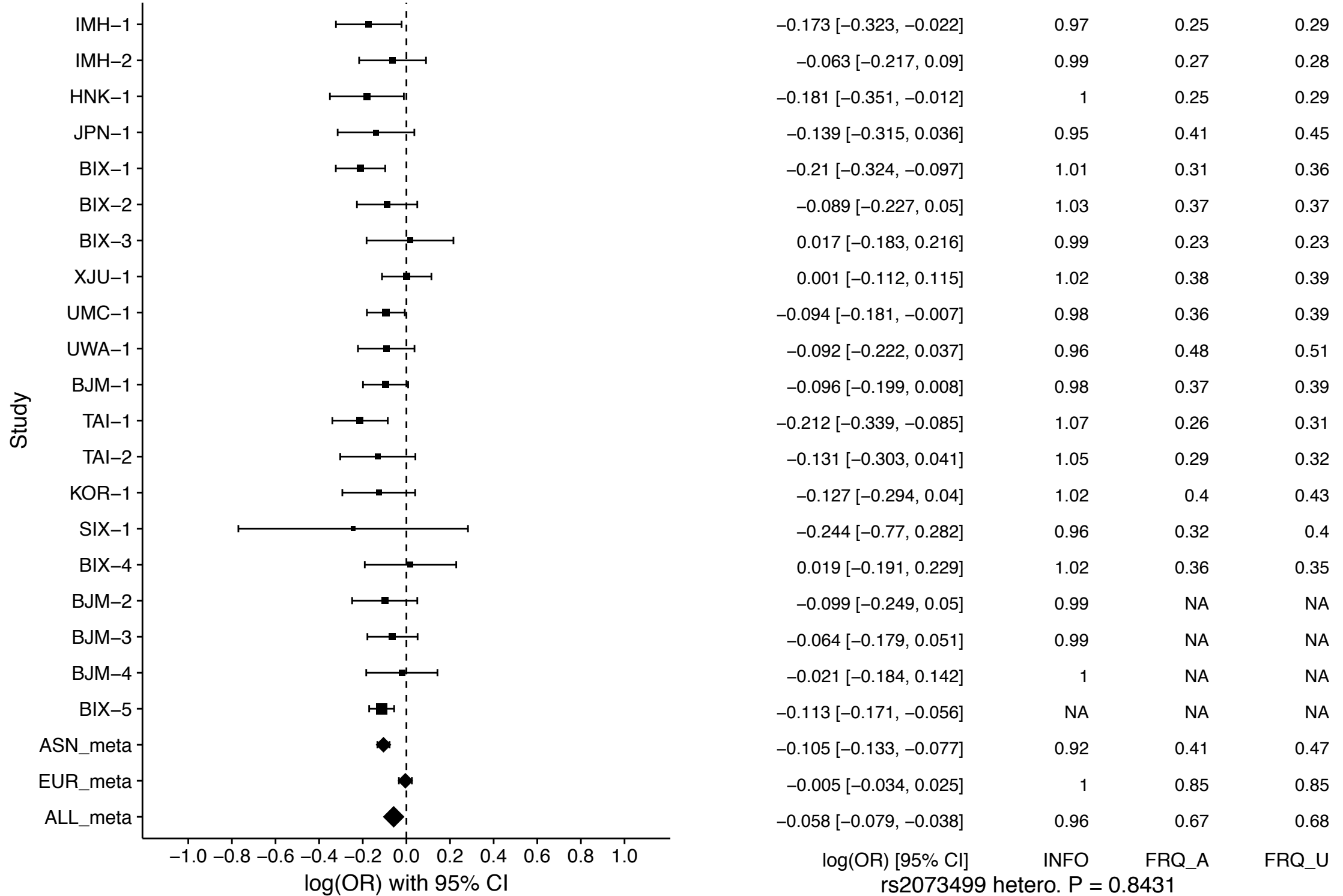


log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs4660761 hetero. P = 0.4028

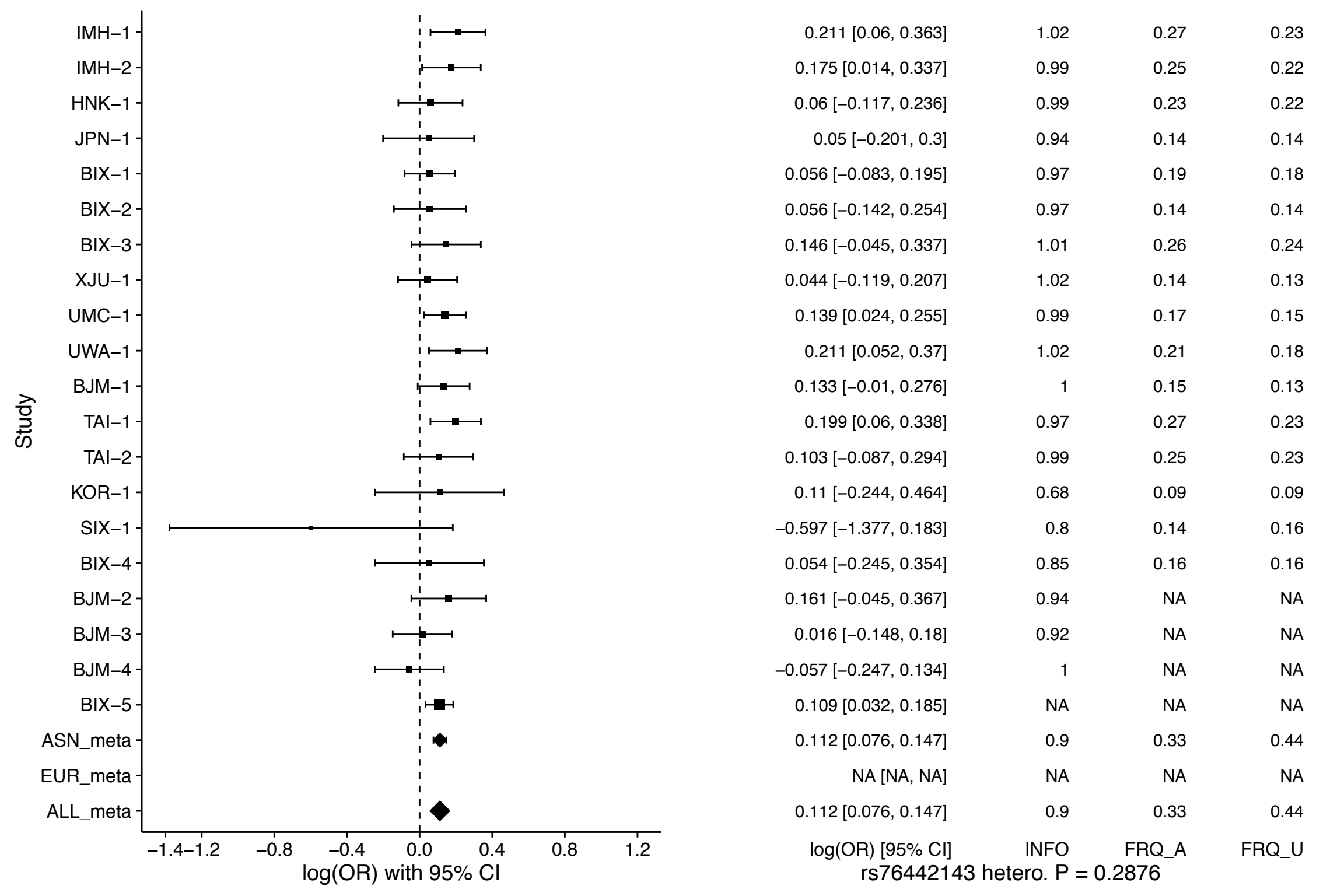


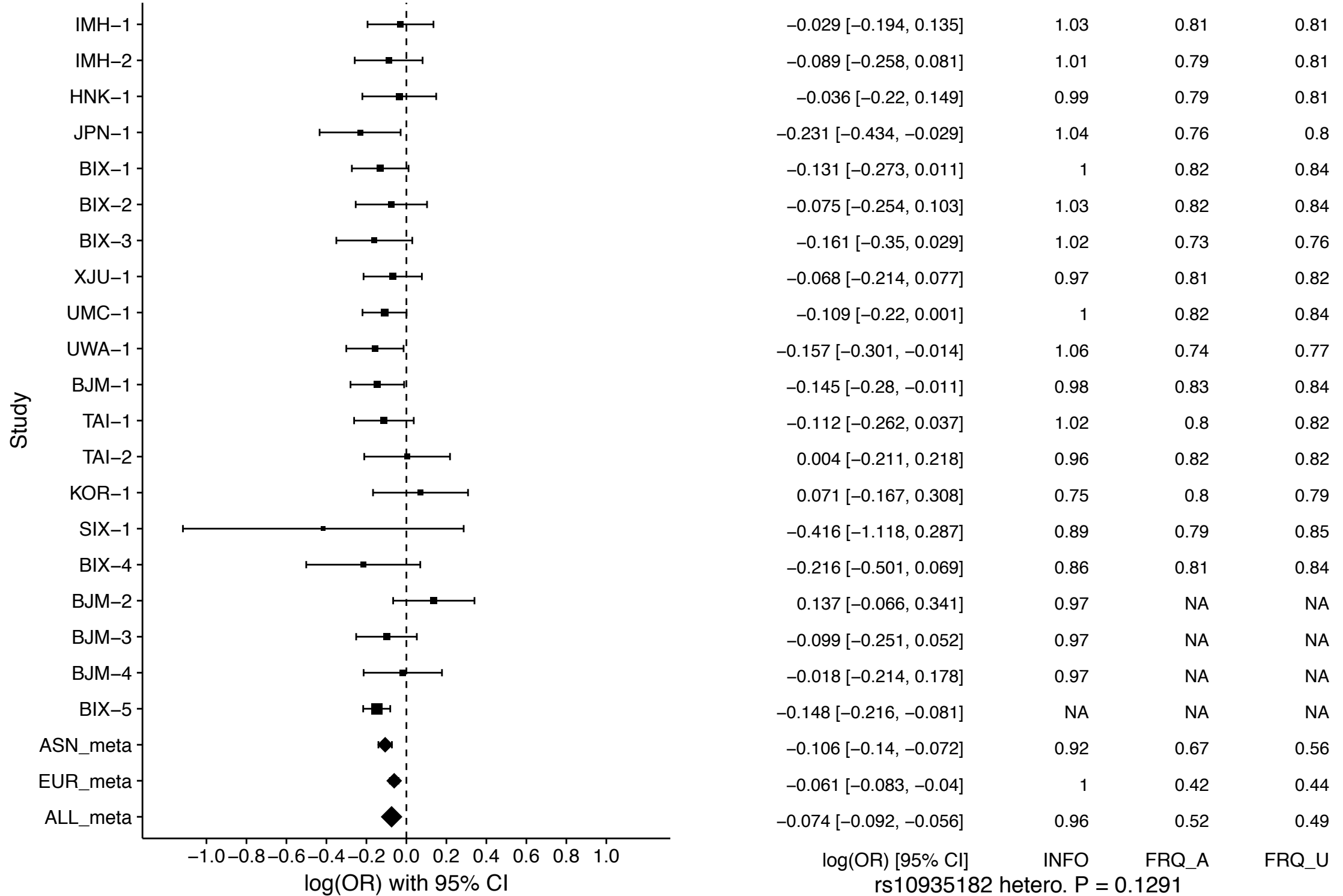
log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs848293 hetero. P = 0.03244

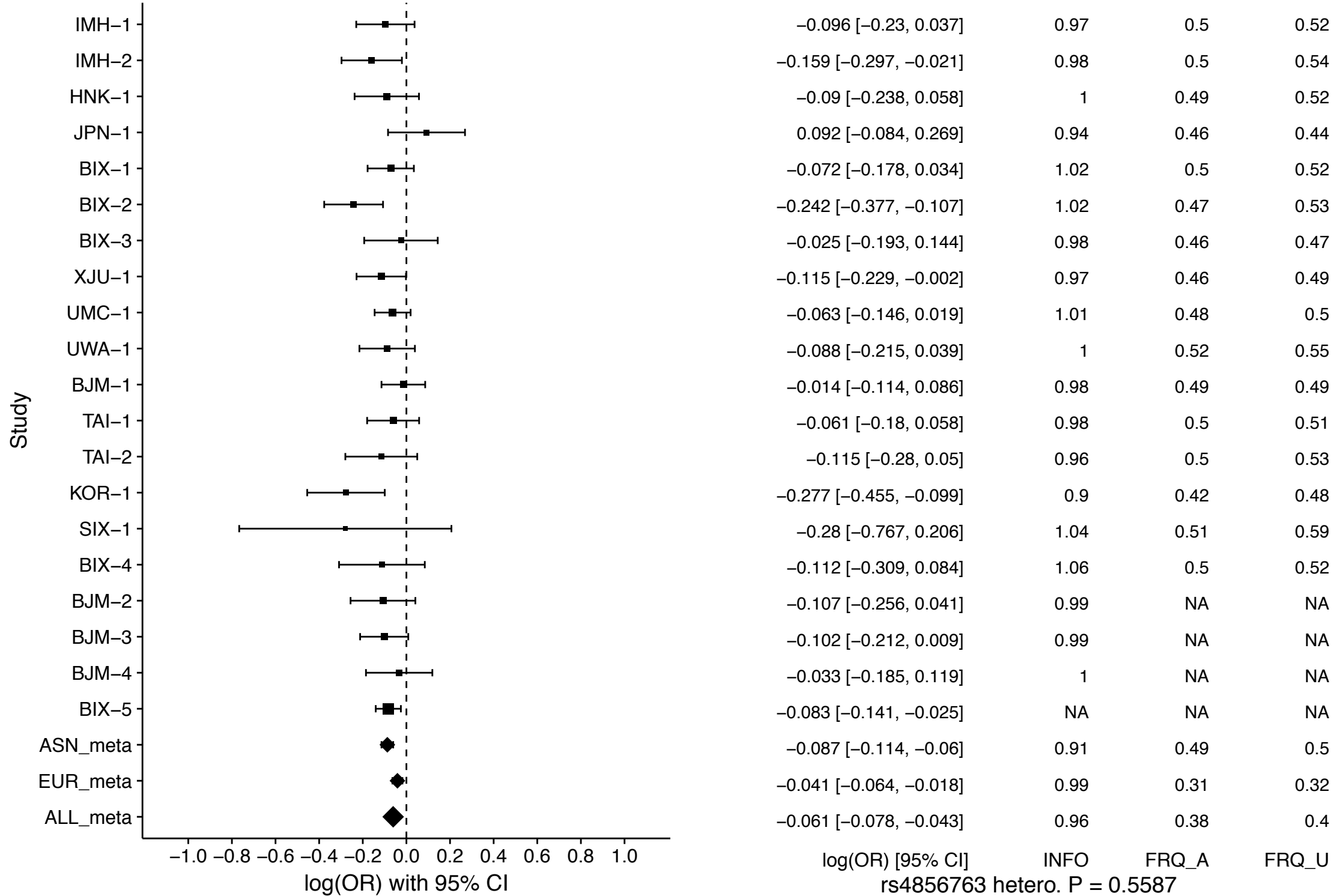




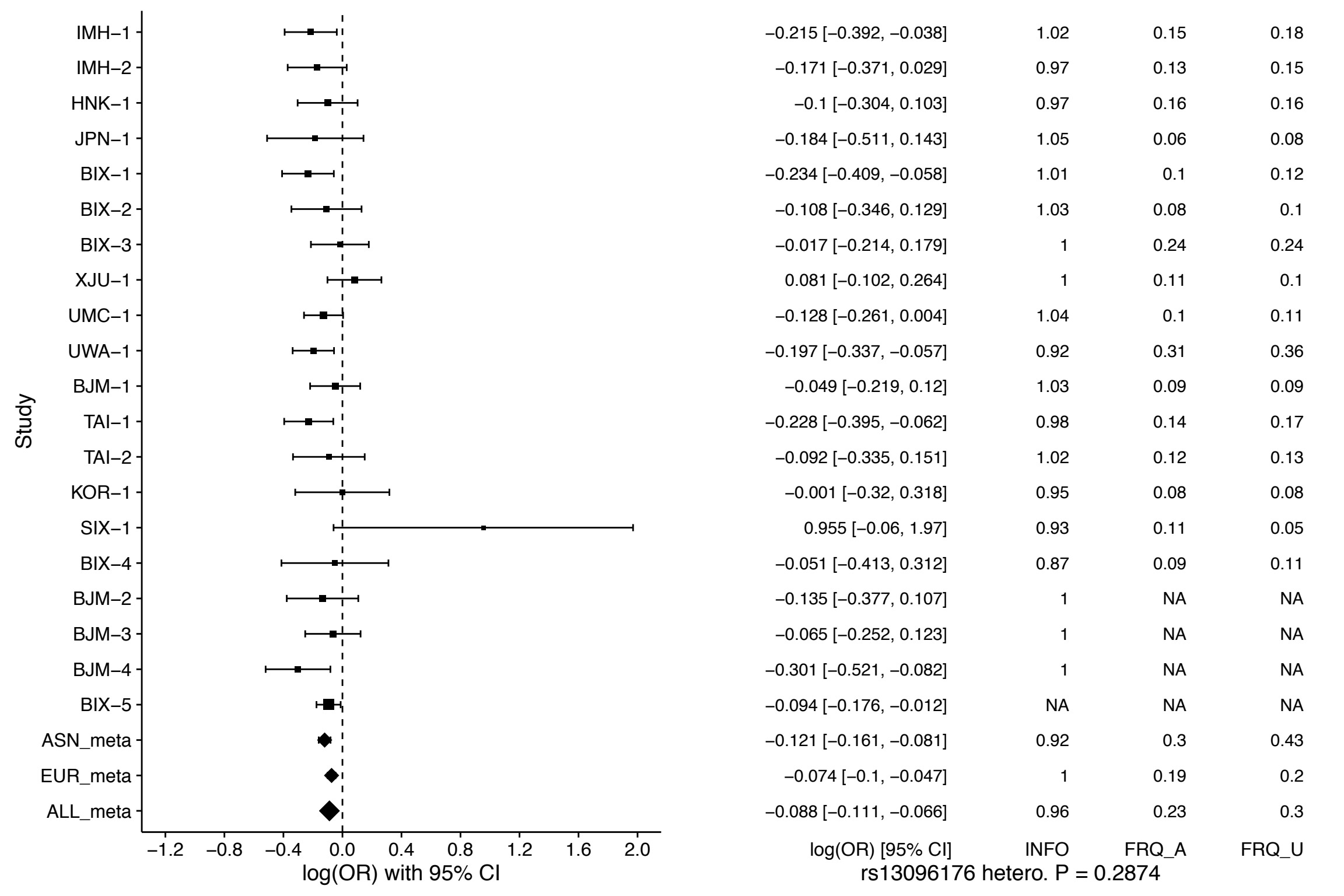
log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs2073499 hetero. P = 0.8431

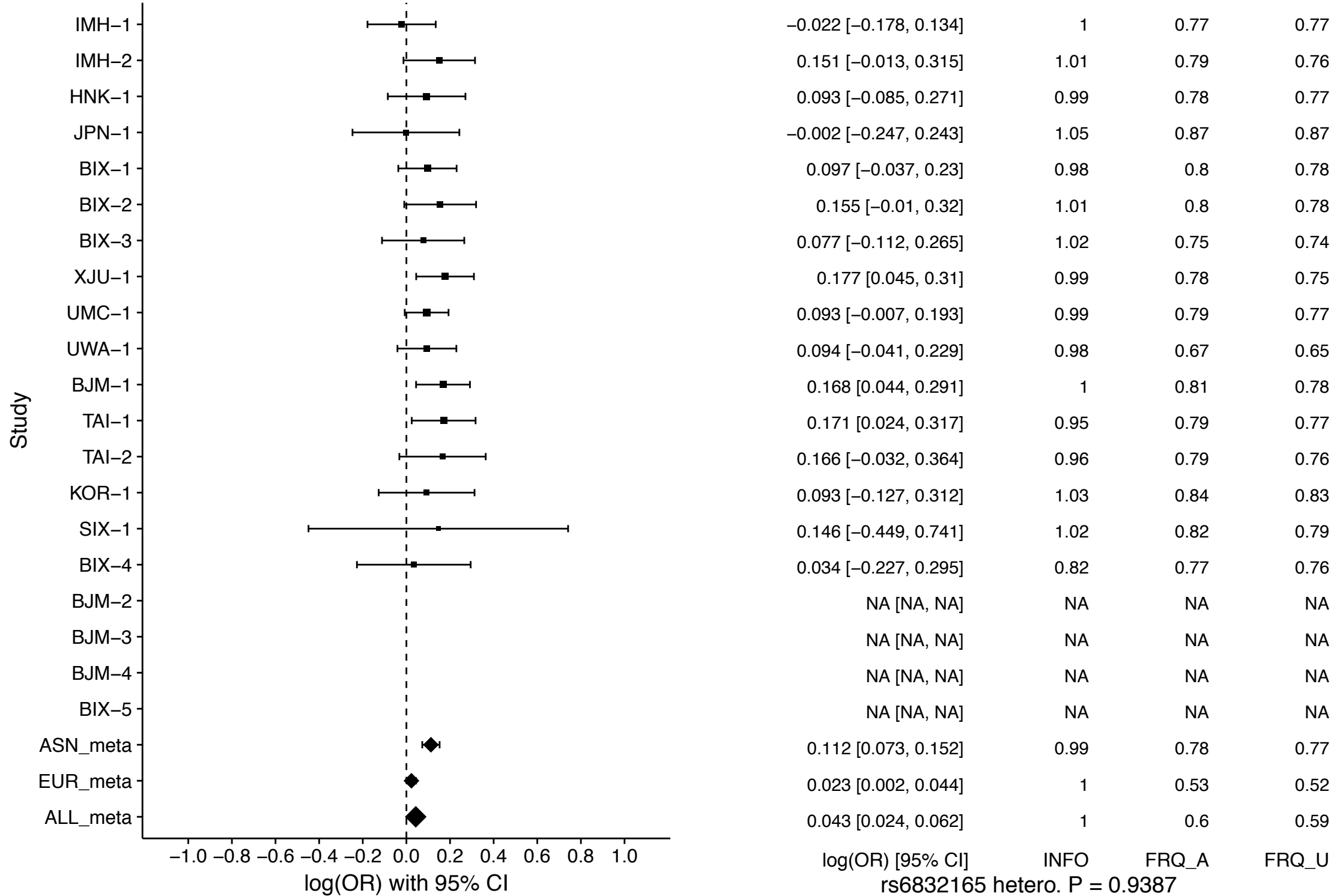




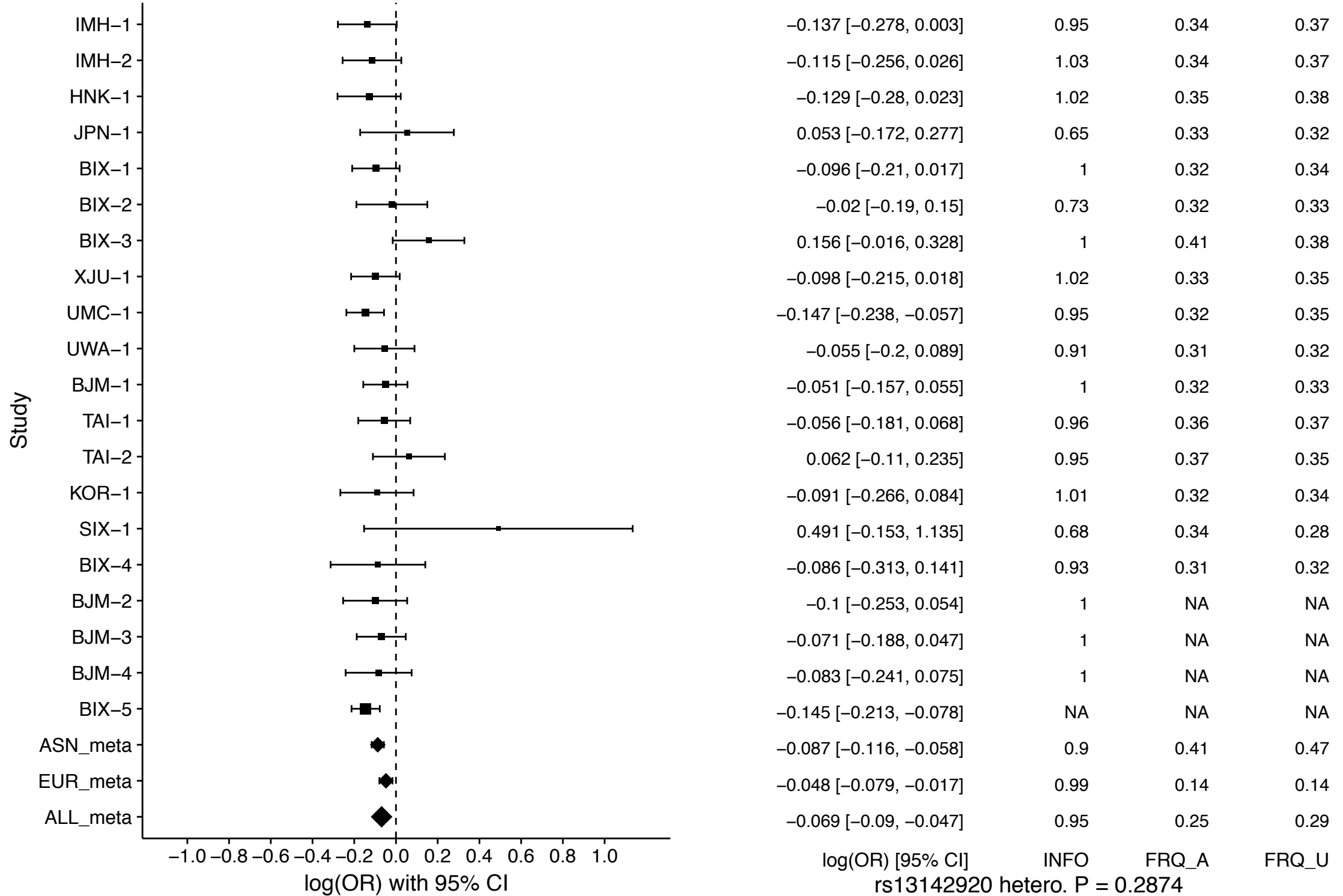


log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs4856763 hetero. P = 0.5587





log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs6832165 hetero. P = 0.9387



-1.0 -0.8 -0.6 -0.4 -0.2 0 0.2 0.4 0.6 0.8 1.0

log(OR) with 95% CI

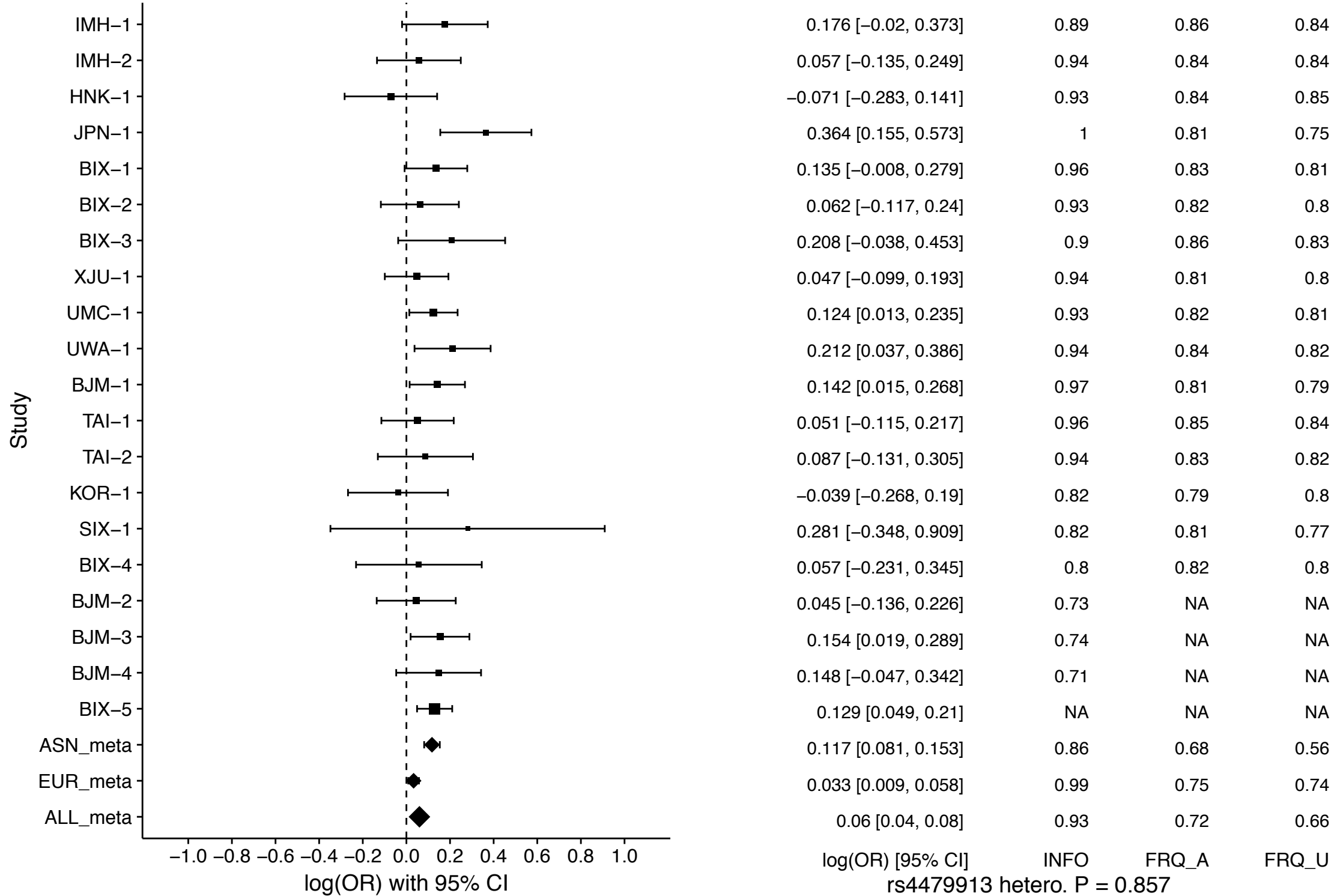
log(OR) [95% CI]

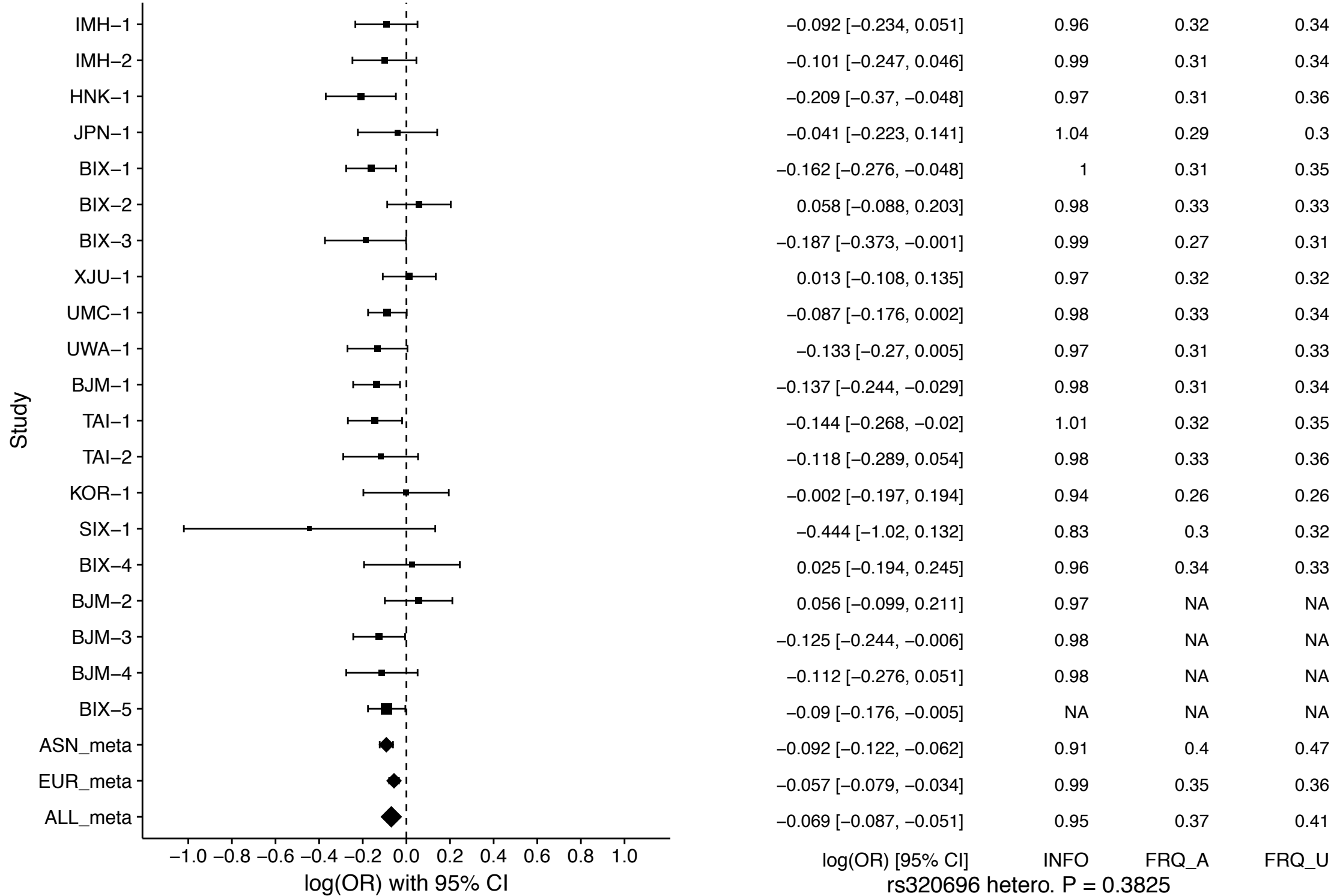
INFO

FRQ_A

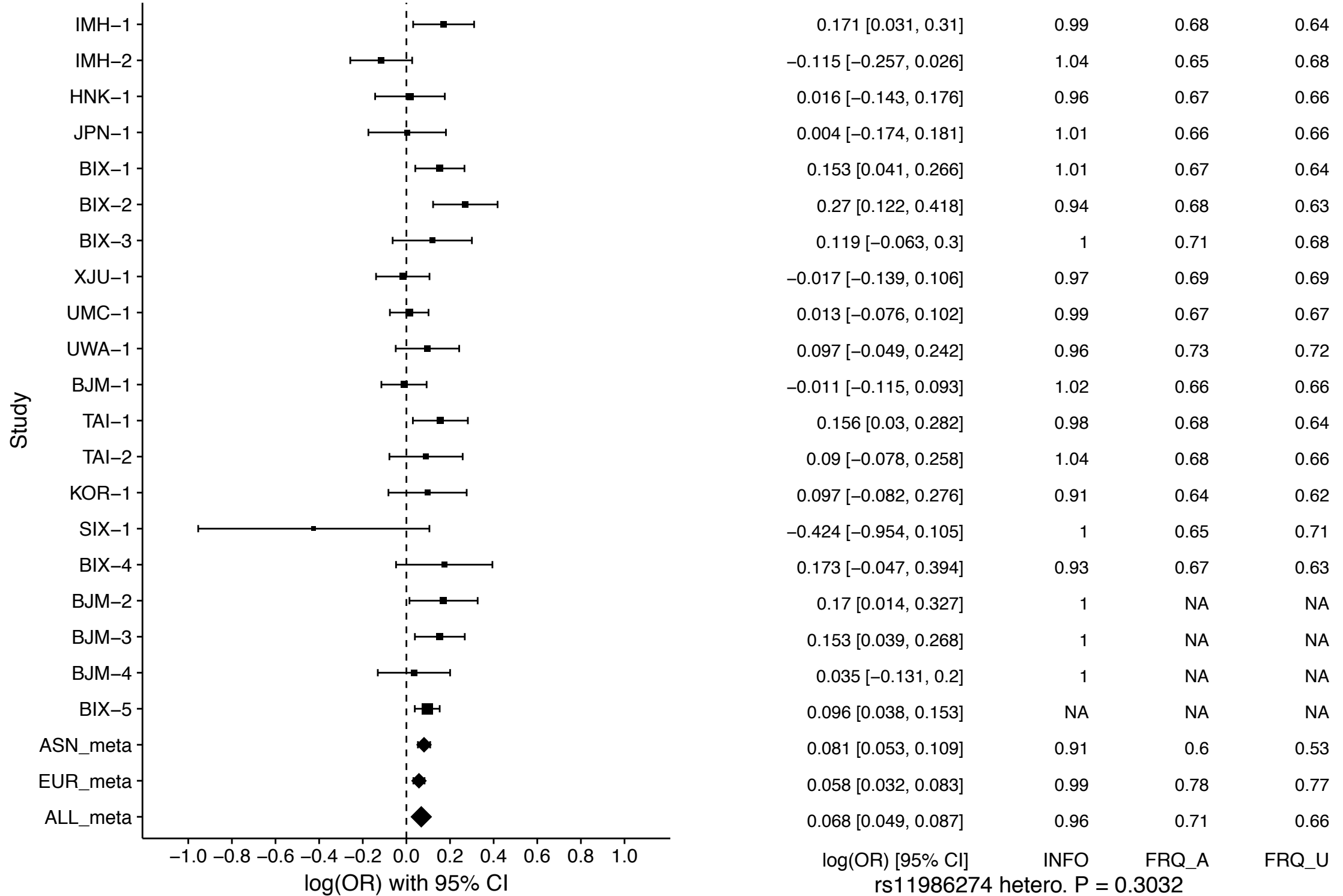
FRQ_U

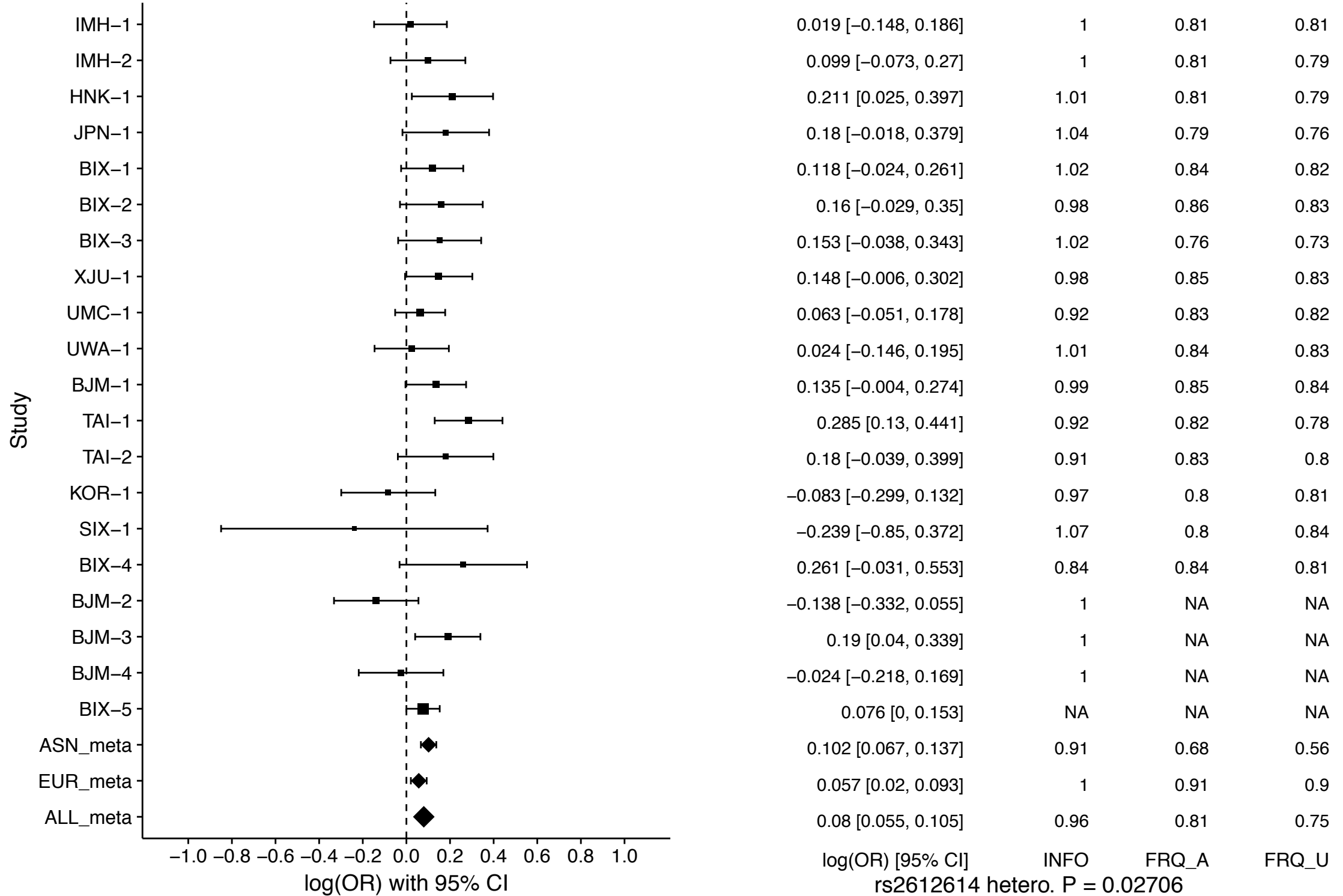
rs13142920 hetero. P = 0.2874



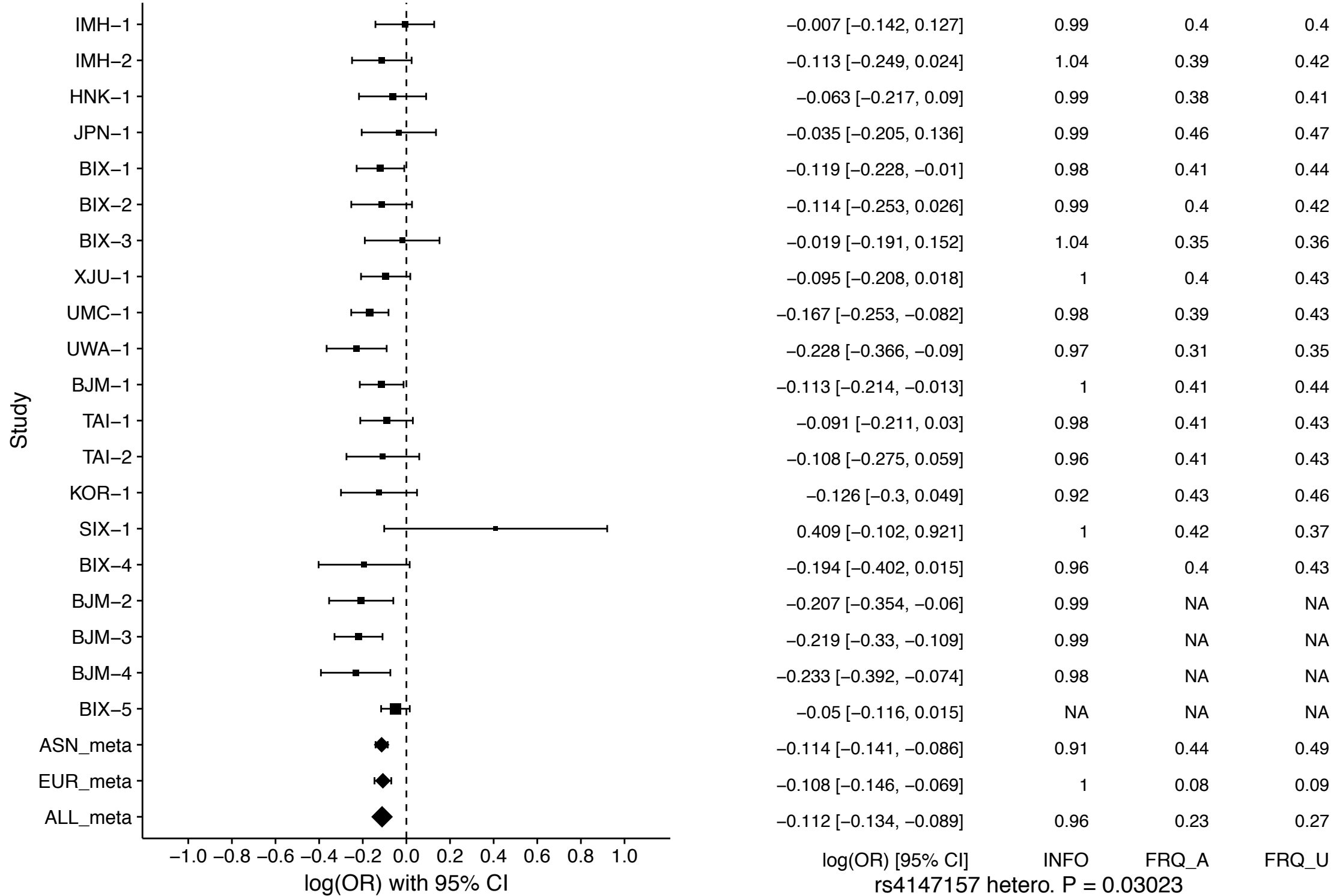


log(OR) [95% CI] INFO FRQ_A FRQ_U
rs320696 hetero. P = 0.3825

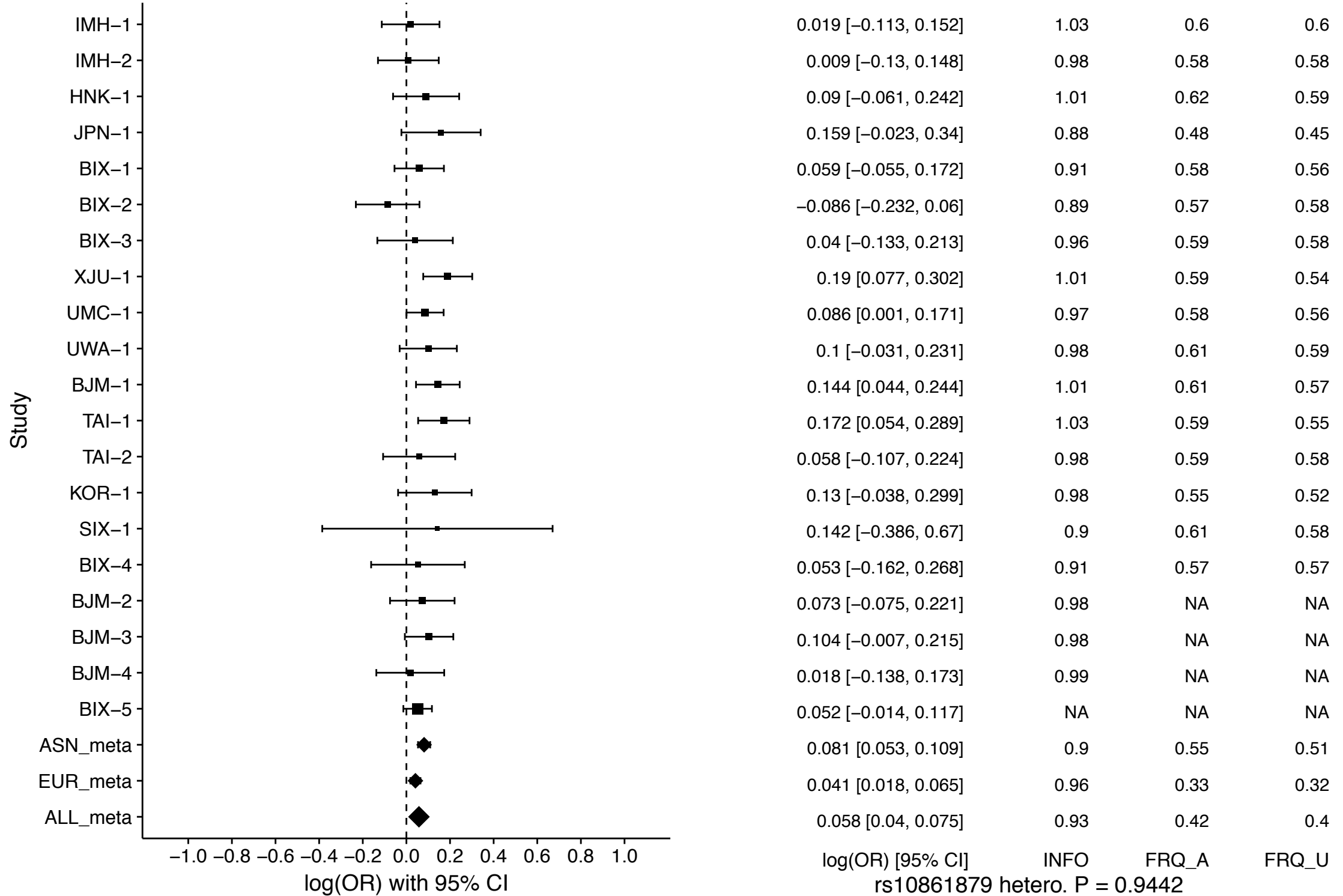


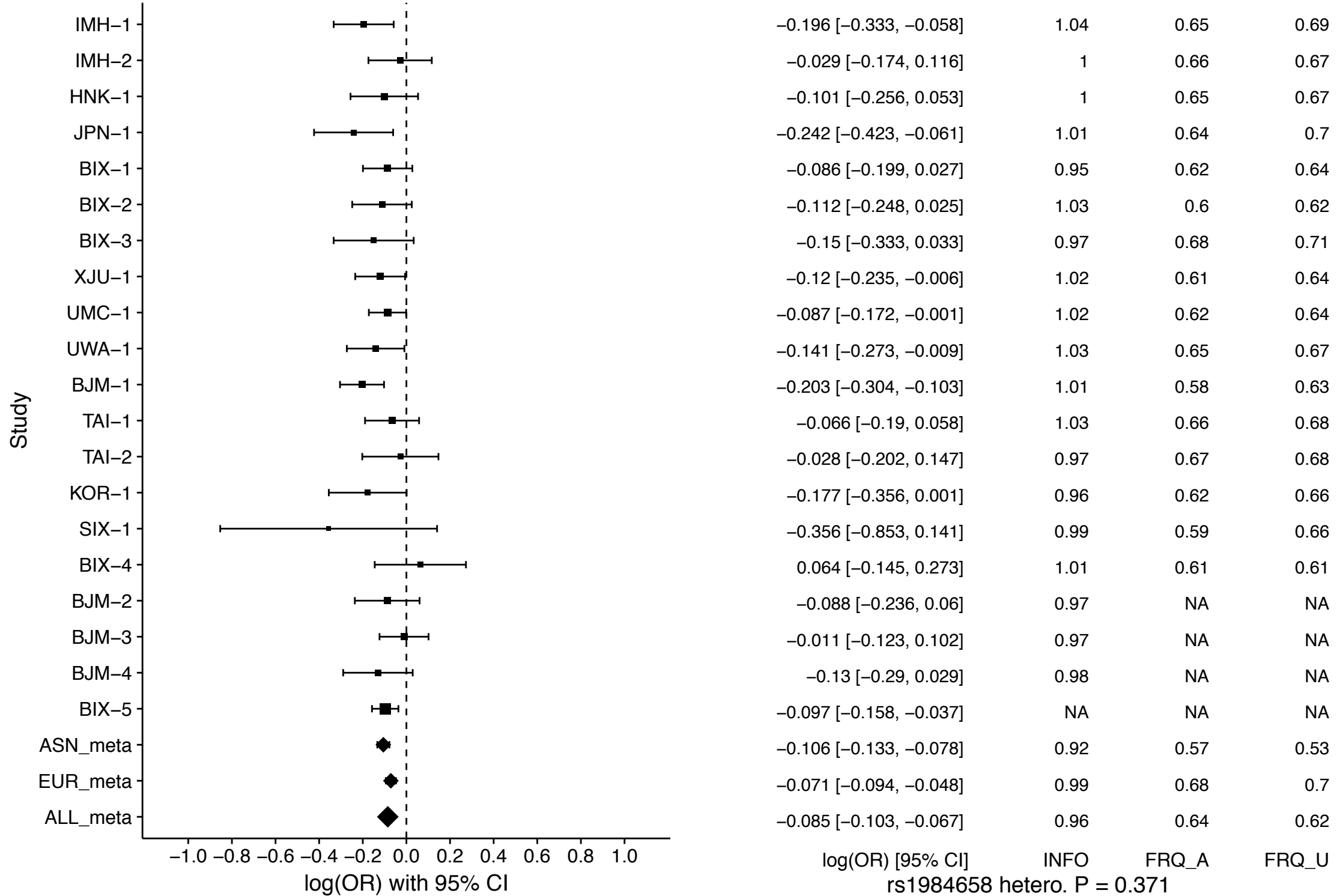


log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs2612614 hetero. P = 0.02706

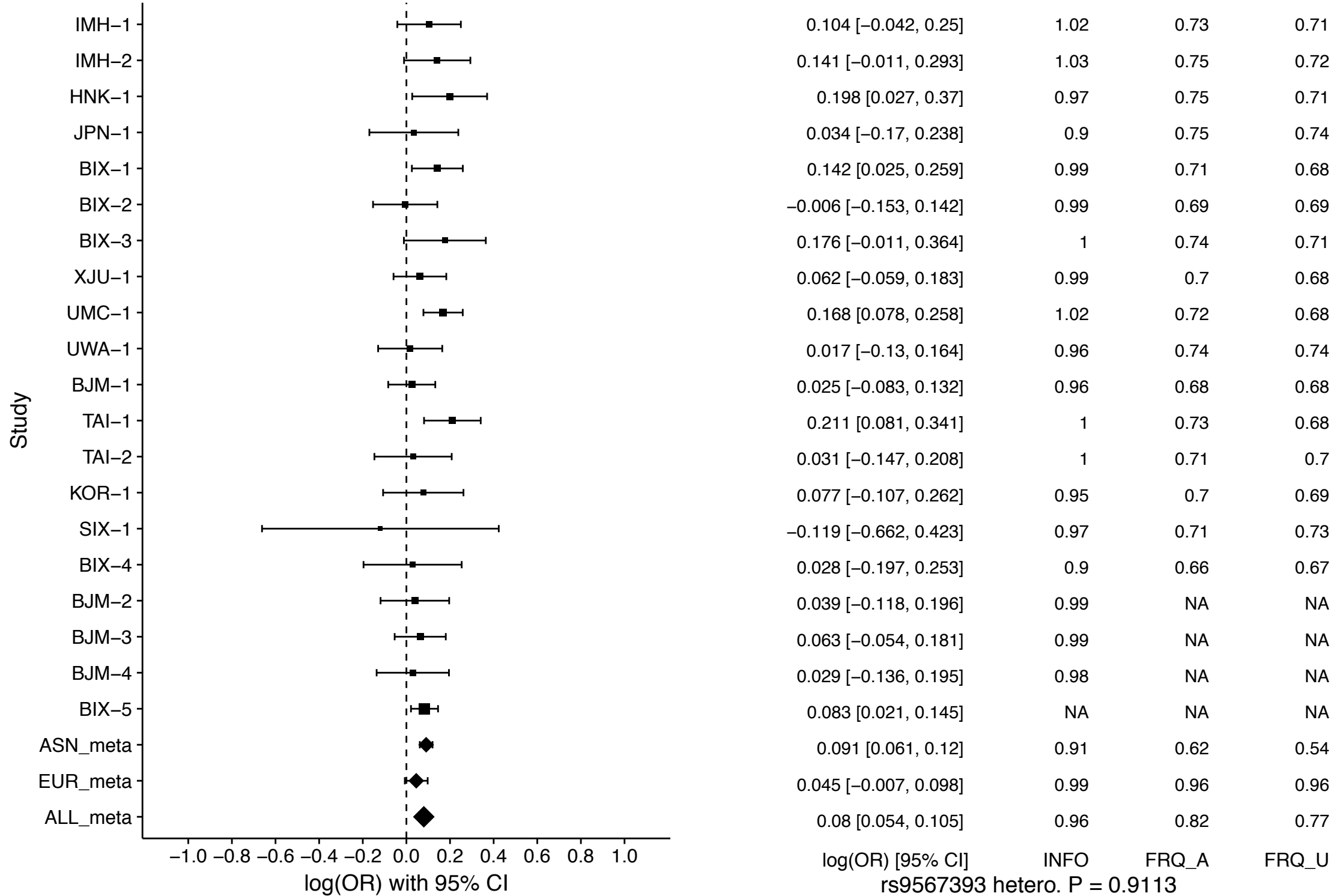


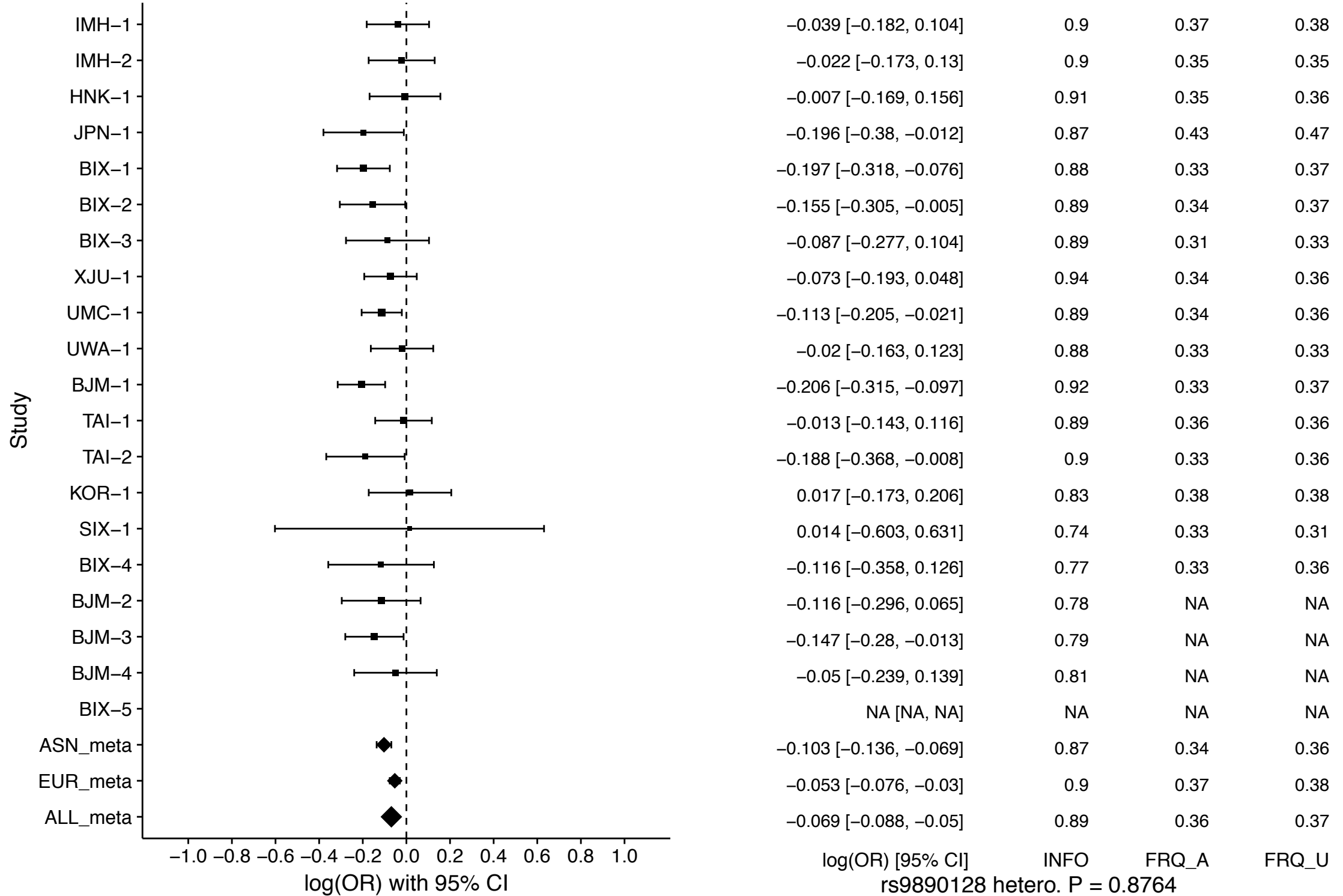
rs4147157 hetero. P = 0.03023





log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs1984658 hetero. P = 0.371



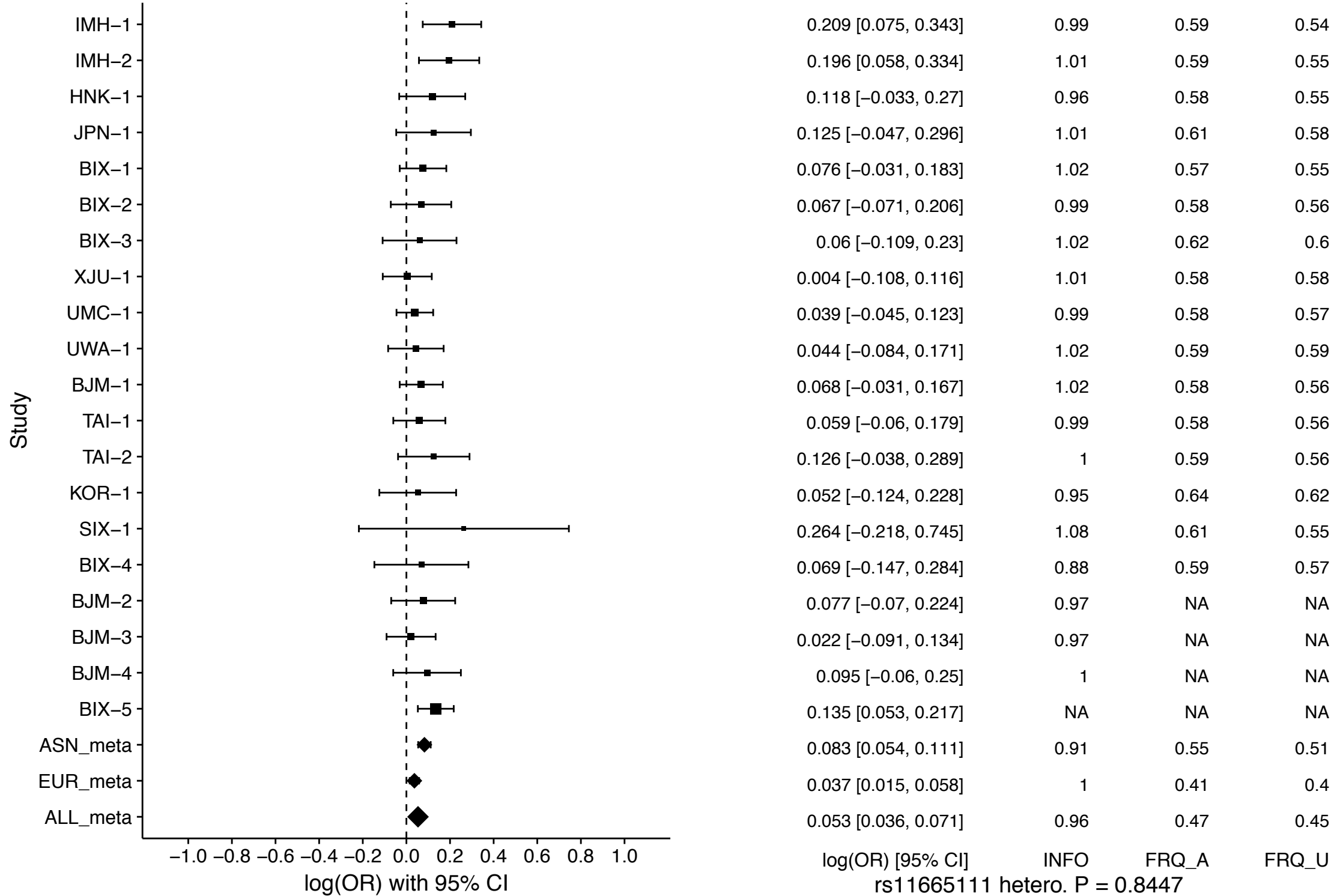


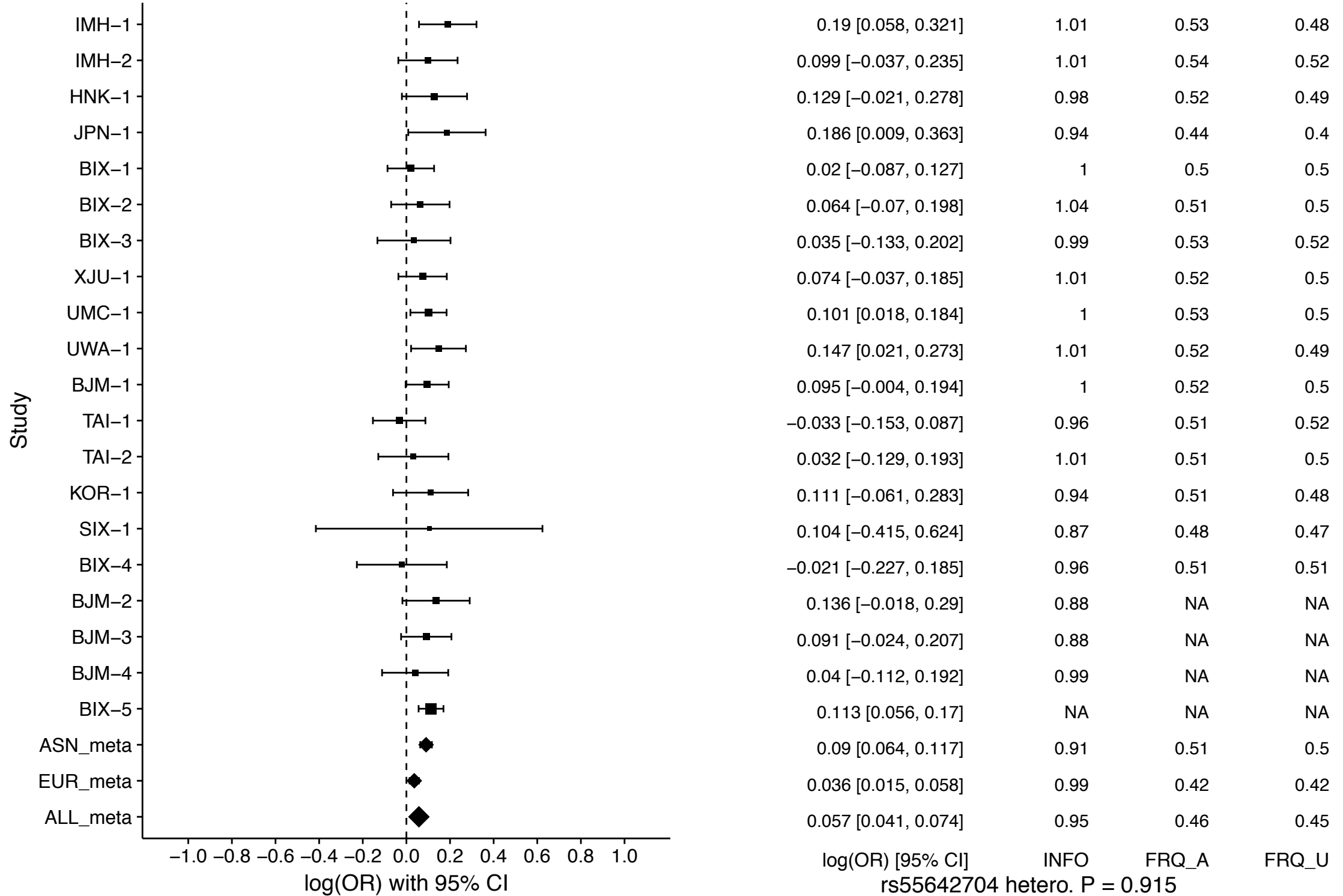
-1.0 -0.8 -0.6 -0.4 -0.2 0.0 0.2 0.4 0.6 0.8 1.0

log(OR) with 95% CI

log(OR) [95% CI] INFO FRQ_A FRQ_U

rs9890128 hetero. P = 0.8764





log(OR) [95% CI] INFO FRQ_A FRQ_U
 rs55642704 hetero. P = 0.915