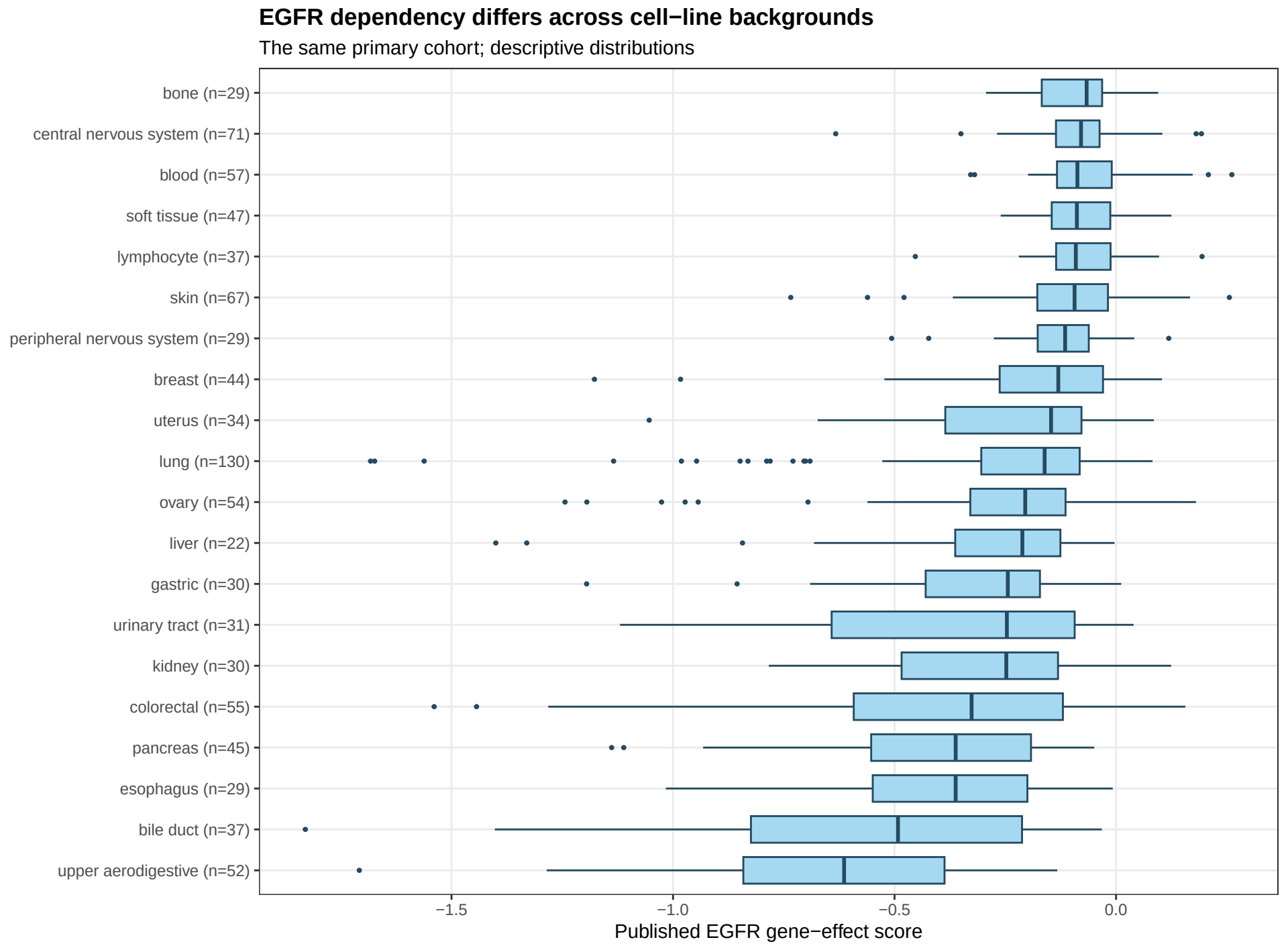
EGFR dependency differs across cell−line backgrounds
The same primary cohort; descriptive distributions
bone (n=29)
central nervous system (n=71)
blood (n=57)
soft tissue (n=47)
lymphocyte (n=37)
skin (n=67)
peripheral nervous system (n=29)
breast (n=44)
uterus (n=34)
lung (n=130)
ovary (n=54)
liver (n=22)
gastric (n=30)
urinary tract (n=31)
kidney (n=30)
colorectal (n=55)
pancreas (n=45)
esophagus (n=29)
bile duct (n=37)
upper aerodigestive (n=52)
−1.5
−1.0
−0.5
0.0
Published EGFR gene−effect score