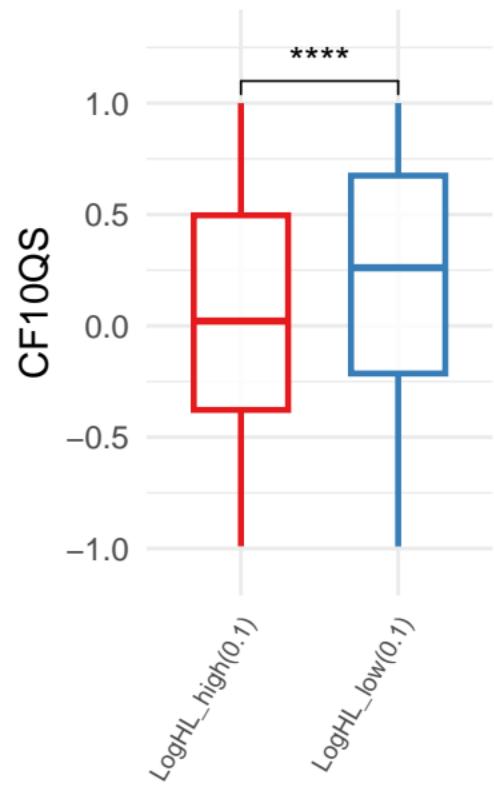
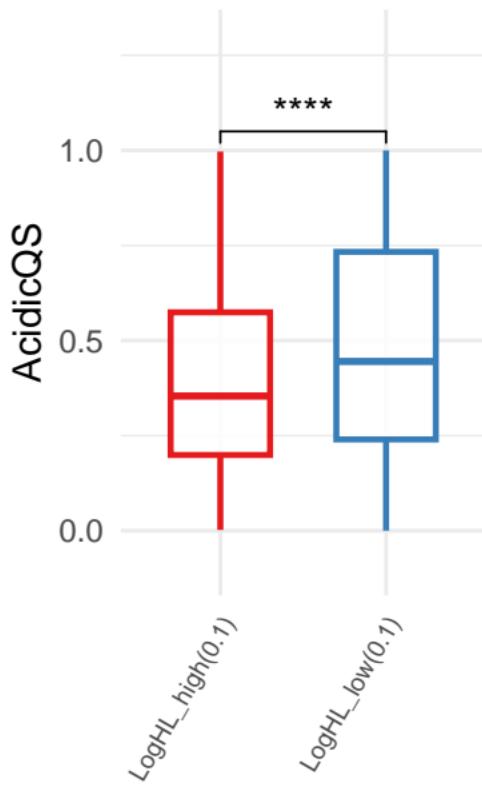


Group  LogHL_high(0.1)  LogHL_low(0.1)

CF10QS 11750 sites



AcidicQS 11750 sites



ub_acc 11750 sites

