Plate 2: MiSeq and qPCR Bd reads by sample (n = 93)

65% total matches
95% total matches if MiSeq reads < 100 are assumed 0
60% total qPCR−/MiSeq− matches (not displayed)
34% (32 samples) qPCR−/MiSeq+ for all reads
3% (3 samples) qPCR−/MiSeq+ if MiSeq reads < 100 are assumed 0

MiSeq count

qPCR
Bd−
Bd+

Sample ID