PacBio IsoSeq (HQ Clustered Reads) Library Size per Sample 500000 450000 400000 type\_step FL\_RAW 350000 FL\_ALIGNED **S** 300000 FL\_ALIGNED.HQ FLNC\_RAW 250000 Pag 200000 FLNC\_ALIGNED FLNC\_ALIGNED.HQ CLUSTERED.HQ\_RAW 150000 CLUSTERED.HQ\_ALIGNED CLUSTERED.HQ\_ALIGNED.HQ 100000 50000 L.v1.r1 L.v1.r2 L.v2.r1 L.v2.r2 M.v1.r1 M.v1.r2 M.v2.r1 M.v2.r2 N.v1.r1 N.v1.r2 Sample