



Supplementary Figure 11: Comparison of replicate correlations and various sample quality metrics in direct library replicates. In A - E, for each replicate pair we show a given quality metric for both replicates, on the y-axis we show correlation between these pairs; each replicate pair is connected by a dashed line. A) Average template length. B) GC contents. C) Average exome coverage. D) rRNA depletion quality. E) Median per-sample TIN. F) Comparison of per-transcript median TIN between library replicate pairs, showing high consistency in transcript quality.