Figure S3: Cold-fixed samples generate reproducible expression profiles, highly correlated with those generated by frozen samples.

(a) Pearson correlation between frozen (F) and Cold-Fixed (CF) or Standard-Fixed (SF) samples, plus correlation between replicate samples from the same cold-fixed tissue. (b-e) Dot plots comparing, expression profiles of RNA from, respectively: frozen vs standard-fixed tissue (b), Frozen vs Cold-Fixed tissue of a representative CRC sample (c), frozen vs Cold-Fixed tissue of a representative breast cancer sample (d), Cold-fixed vs replicate sample extracted from the same cold-fixed tissue (e).

