Figure S1. Validation of the correlation between CNV load and SNV in TCGA-breast cancer samples (n=962). (A) The correlation between CNV load and SNV count in TCGA. (B) The correlation between CNV load and the number of mutated genes in TCGA.

Figure S2. The IGV screenshot of *C12orf76::ZFAT* fusions in RNA-Seq.

Figure S3. The IGV screenshot of *CLTC::DHX40* fusions in RNA-Seq.

Figure S4. The IGV screenshot of *CKMT2-AS1::PDE4D* fusions in RNA-Seq.

Figure S5. The IGV screenshot of *NAV1::PKP1* fusions in RNA-Seq.

Figure S6. The IGV screenshot of *EXOC6::NSMCE4A* fusions in RNA-Seq.

Figure S7. The IGV screenshot of *RBM38::RAE1* fusions in RNA-Seq.

Figure S8. The IGV screenshot of *TOM1L2::TRPC4AP* fusions in RNA-Seq.

Figure S9. The IGV screenshot of *EIF3L::TRIOBP* fusions in RNA-Seq.