

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 10, 2026

## FusionSeq

RRID:SCR\_013329

Type: Tool

### Proper Citation

FusionSeq (RRID:SCR\_013329)

### Resource Information

**URL:** <http://archive.gersteinlab.org/proj/rnaseq/fusionseq/>

**Proper Citation:** FusionSeq (RRID:SCR\_013329)

**Description:** A modular framework for finding gene fusions by analyzing Paired-End RNA-Sequencing data.

**Abbreviations:** FusionSeq

**Resource Type:** software resource

**Resource Name:** FusionSeq

**Resource ID:** SCR\_013329

**Alternate IDs:** OMICS\_01351

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20260808T190021+0000

### Ratings and Alerts

No rating or validation information has been found for FusionSeq.

No alerts have been found for FusionSeq.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

### Usage and Citation Metrics

We found 7 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

Omar M, et al. (2024) Semi-Supervised, Attention-Based Deep Learning for Predicting TMPRSS2:ERG Fusion Status in Prostate Cancer Using Whole Slide Images. Molecular cancer research : MCR, 22(4), 347.

Wei T, et al. (2021) Re-Evaluate Fusion Genes in Prostate Cancer. Cancer informatics, 20, 11769351211027592.

Beg S, et al. (2021) Integration of whole-exome and anchored PCR-based next generation sequencing significantly increases detection of actionable alterations in precision oncology. Translational oncology, 14(1), 100944.

Pisapia DJ, et al. (2020) Fusions involving BCOR and CREBBP are rare events in infiltrating glioma. Acta neuropathologica communications, 8(1), 80.

Beltran H, et al. (2016) Divergent clonal evolution of castration-resistant neuroendocrine prostate cancer. Nature medicine, 22(3), 298.

Latysheva NS, et al. (2016) Discovering and understanding oncogenic gene fusions through data intensive computational approaches. Nucleic acids research, 44(10), 4487.

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.