Anfo
RRID:SCR_015973
Type: Tool

Proper Citation

Anfo (RRID:SCR_015973)

Resource Information

URL: https://bioinf.eva.mpg.de/anfo/

Proper Citation: Anfo (RRID:SCR_015973)

Description: Anfo is a mapper in the spirit of Soap/Maq/Bowtie, but its implementation takes more after BLAST/BLAT. It’s most useful for the alignment of sequencing reads where the DNA sequence is somehow modified (think ancient DNA or bisulphite treatment) and/or there is more divergence between sample and reference than what fast mappers will handle gracefully (say the reference genome is missing and a related species is used instead).

Resource Type: Resource

Website Status: Last checked up

Resource Name: Anfo

Resource ID: SCR_015973

Ratings and Alerts

No rating or validation information has been found for Anfo.

No alerts have been found for Anfo.

Data and Source Information

Source: SciCrunch Registry
Usage and Citation Metrics

We have not found any literature mentions for this resource.