Anfo  
RRID:SCR_015973  
Type: Tool  

Proper Citation  
Anfo (RRID:SCR_015973)  

Resource Information  

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

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 Name: Anfo  
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Resource Type: Resource  
Resource ID: SCR_015973  
Website Status: Last checked up  

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.