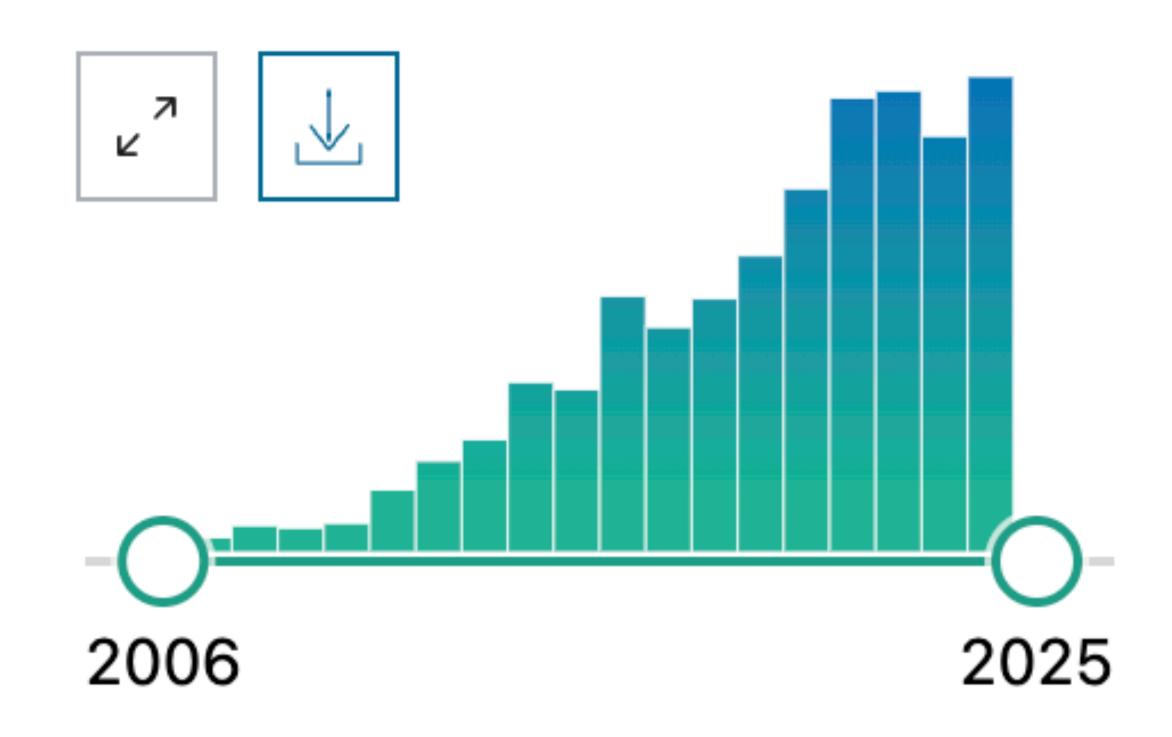
Bioinformatics tools and pipelines

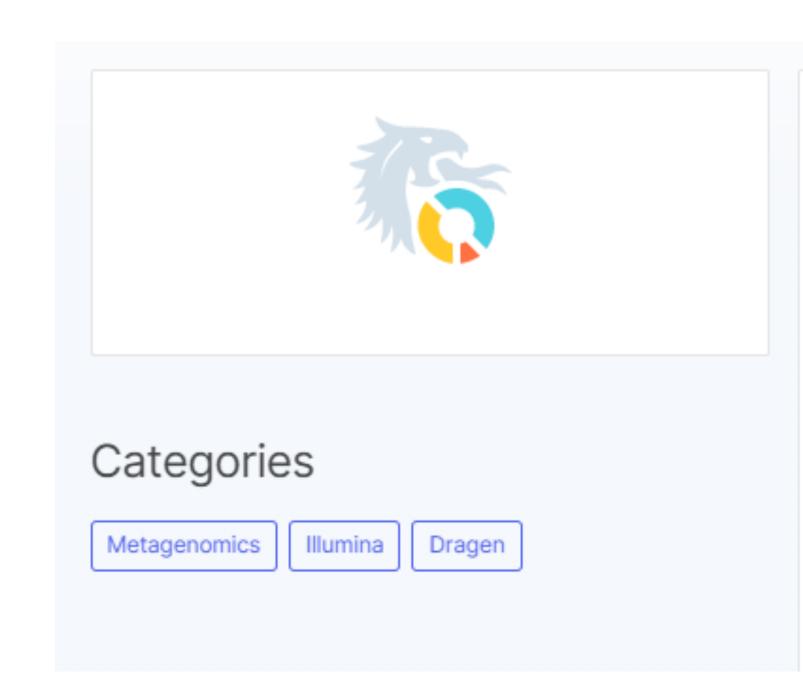
In Metagenomics

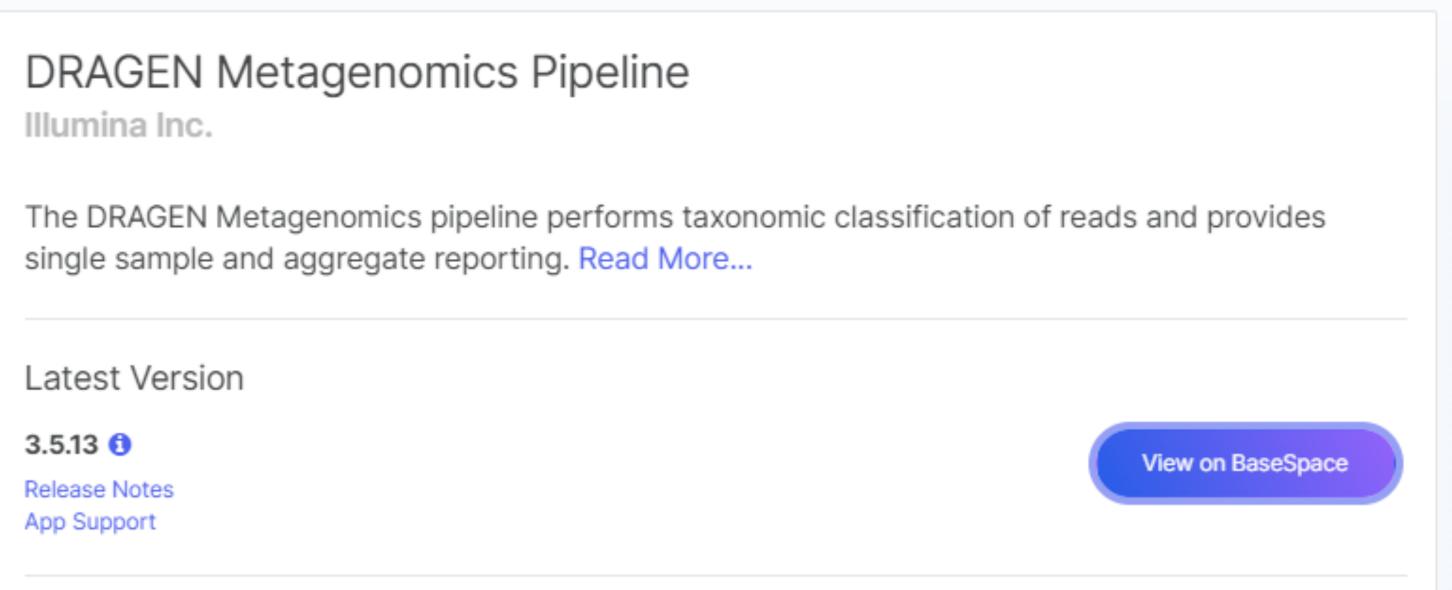
• Past, present, and future

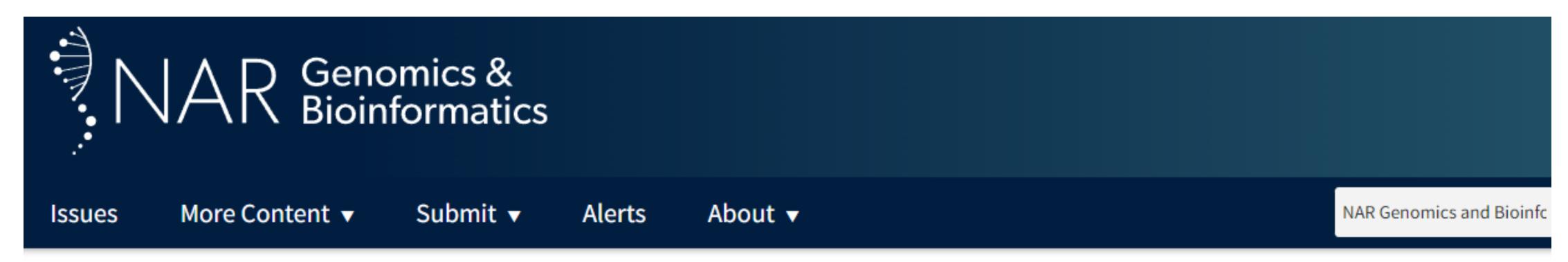
 Pubmed search for "metagenomics pipeline"

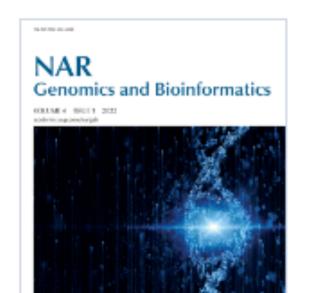
RESULTS BY YEAR











JOURNAL ARTICLE

nf-core/mag: a best-practice pipeline for metagenome hybrid assembly and binning 3

Sabrina Krakau ™, Daniel Straub, Hadrien Gourlé, Gisela Gabernet, Sven Nahnsen