

Arvados - Support #14906

[SDK] Maintain Arvados SDKs in Bioconda

03/01/2019 02:58 PM - Peter Amstutz

Status:	New	Start date:	
Priority:	Normal	Due date:	
Assigned To:		% Done:	0%
Category:		Estimated time:	0.00 hour
Target version:	To Be Groomed		
Description			
<p>Bioconda is a channel for the conda package manager specializing in bioinformatics software. Conda is a rootless package manager that is popular in scientific computing environments where neither root nor containers are available. Bioconda packages are also automatically added to Biocontainers (http://biocontainers.pro) via Muled (http://biocontainers.pro/docs/developer-manual/developer-intro/)</p> <p>https://bioconda.github.io/</p> <p>Some Arvados components are already packaged for it, originally contributed by Brad Chapman:</p> <p>https://github.com/bioconda/bioconda-recipes/tree/master/recipes/arvados-cli</p> <p>https://github.com/bioconda/bioconda-recipes/tree/master/recipes/arvados-cwl-runner</p> <p>https://github.com/bioconda/bioconda-recipes/tree/master/recipes/arvados-python-client</p> <p>We should support these directly so that they don't get out of date.</p>			

History

#1 - 03/01/2019 02:58 PM - Peter Amstutz

- Status changed from New to In Progress

#2 - 03/01/2019 03:11 PM - Peter Amstutz

- Description updated

#3 - 03/01/2019 03:11 PM - Peter Amstutz

- Status changed from In Progress to New

#4 - 03/01/2019 03:11 PM - Peter Amstutz

- Subject changed from [SDK] Arvados SDKs in Bioconda to [SDK] Maintain Arvados SDKs in Bioconda

#5 - 03/01/2019 03:18 PM - Brad Chapman

This would be great and happy to help. You've already got the relevant recipes listed above. Bioconda is an open project so they'd be open to anyone contributing and docs are here:

<https://bioconda.github.io/contributing.html>

Let me know how I can help with this.

#6 - 03/01/2019 03:26 PM - Tom Morris

- Target version set to To Be Groomed