

GET-Evidence - Feature #496

Import and process raw reads for Jay Flatley genome

05/13/2010 01:11 PM - Madeleine Ball

Status:	New	Start date:	
Priority:	Low	Due date:	
Assigned To:	Tom Clegg	% Done:	0%
Category:		Estimated time:	0.00 hour
Target version:		Totalhours:	
Billable:		Resolution:	
Estimatedhours:			
Hours:			

Description

<http://developer.amazonwebservices.com/connect/entry.jspa?externalID=3357&categoryID=279>

We should import and process reads for this genome.

"This data set contains the raw export files of the first genome sequenced by Illumina Individual Genome Service using Illumina's Genome Analyzer technology of paired 75-base reads. 92,254,659,274 bases were used to generate a consensus sequence with coverage of 32x average depth. The genome was obtained via peripheral blood of Jay Flatley, CEO of Illumina."

History

#1 - 02/15/2011 03:17 PM - Ward Vandewege

- Project changed from External to GET-Evidence
- Category deleted (GET-Evidence)

#2 - 02/16/2011 02:28 PM - Madeleine Ball

- Priority changed from Normal to Low