



	NO WAIT TIMES	NO STORAGE QUOTAS	NO JOB SUBMISSION LIMITS	NO DATA TRANSFER BOTTLENECKS	NO IT EXPERIENCE REQUIRED	NO REQUIRED INFRASTRUCTURE
GALAXY MAIN	X	X	X	X		
LOCAL GALAXY	?	?	?		X	X
CLOUD GALAXY		Ø		X	X	
SLIPSTREAM GALAXY						

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TOOLS	TASK	DATA	RUN-TIME
Bowtie 2	Mapping whole human genome	204 million paired-end 100bp Illumina reads	2 Hours 44 Minutes
SAMTools	SAM-BAM conversion	127GB SAM (41GB resulting BAM)	2 Hours 7 Minutes
TopHat 2	RNA-Seq mapping	24 million 100bp Illumina reads	1 Hours 24 Minutes
Cufflinks 2	Differential Expression Analysis	4.3 GB SAM File	0 Hours 11 Minutes



INTEL® XEON® E5 PROCESSORS (16 CORES)

384 GB RAM

100GB SOLID STATE DRIVE

16TB INTEGRATED HIGH-SPEED STORAGE

FLEXIBLE LINUX-BASED SERVER ARCHITECTURE



GET RESULTS FASTER WITH A DEDICATED GALAXY SERVER

SAVETIME AND MONEY	AVERAGE LAB CAN SAVE > 1 MONTH IN DEPLOYMENT TIME AND >\$20,000 IN START-UP COSTS		
HARDWARE SPECIFICATIONS Designed for data intensive analysis	CPUs, RAM, network capability, storage, and more are optimally selected and configured		
EFFICIENT RESOURCE MANAGEMENT Decreases time to results	SGE and Galaxy installed and configured with appropriate resource management		
PLUG-AND-PLAY APPLIANCE Helps to get up and running quickly	Galaxy and underlying computational tools are pre-installed on the appliance		
OPTIMIZED DATA TRANSFER Improves performance	Galaxy software and SlipStream hardware have been tuned for faster uploads and downloads		
AUTOMATED UPDATES Minimizes IT management overhead	Galaxy and underlying computational tools can be automatically updated		
PRE-LOADED DATASETS Makes data analysis more convenient	Commonly used reference datasets for use with Galaxy are pre-loaded on the appliance		

SEEKING THOUGHT LEADERS FOR EARLY ACCESS PROGRAM

Limited Availability - CONTACT NOW TO JOIN

slipstream-galaxy@bioteam.net www.bioteam.net/slipstream/galaxy-edition