Approved	? Group #	Members	Date	Article title
Yes	1	Ben Hong, Olivia Dai, Max Jentzsch	Wed, Mar 1, 2017	No evidence for extensive horizontal gene transfer in the genome of the tardigrade Hypsibius dujardini
Yes	2	Molly McNamara, Michael Macarthur, Ilana Kelsey, Jason Lin	Wed, Mar 1, 2017	Bronchoalveolar Lavage Proteomics in Patients with Suspected Lung Cancer
Yes	3	Hector Perez, Taylor Sehein, Katia Chadaideh	Wed, Mar 1, 2017	Diet Dominates Host Genotype in Shaping the Murine Gut Microbiota
Yes	4	Ibrahim Diakite, Jonathan Batty	Mon, Mar 6, 2017	Meta-analysis of host response networks identifies a common core in tuberculosis
Yes	5	Jiwon Lee, Min-Jung Wang, Mutong Zhao, Qianyu Yuan, You Wu	Mon, Mar 6, 2017	Analysis of protein-coding genetic variation in 60,706 humans
Yes	6	Samantha Giffen, Eddie Irvine, Joanna Olivas, Greg Babunovic	Mon, Mar 6, 2017	Highly Parallel Genome-wide Expression Profiling of Individual Cells Using Nanoliter Droplets
Yes	7	Conor Foley, Isabelle Mieling, Alex Wu, Yongling Ye, Nancy Zhang	Mon, Mar 6, 2017	A network of conserved synthetic lethal interactions for exploration of precision cancer therapy
Yes	8	Alberta Wang, Kathy Lee, Amit Joshi	Wed, Mar 8, 2017	Association of host genome with intestinal microbial composition in a large healthy cohort
Yes	9	Qiuhao Zhang, Xinan Wang, Fangyuan Hong, Yuxi Liu	Wed, Mar 8, 2017	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer
Yes	10	Kaylyn Williamson, Andrew Goldfarb, Soun Lee, Marsha Wibowo	Wed, Mar 8, 2017	Cell-free DNA comprises an in vivo nucleosome footprint that informs its tissues-of-origin
Yes	11	Ibrahim Kurt, Hazreen Abdul Majid	Wed, Mar 8, 2017	HiChIP: efficient and sensitive analysis of protein-directed genome architecture