Wurmlab

Community: Journal table of contents highlights

published between 1st and last day of **JANUARY**** (in print *and* early access). **Finish it by February 12th**. Please add everything as nested bullet-points.

CTRL-K for links

CTRL-SHIFT-V paste w/o format

Look at the table of content duties here (github)

Am Nat

- Dridi & Erol Akçay. <u>Learning to Cooperate: The Evolution of Social Rewards in Repeated Interactions</u>
- Diane C. Wiernasz and Blaine J. Cole <u>Offspring Size and Reproductive Allocation in</u> <u>Harvester Ants</u>

Bioinformatics

- DART: a fast and accurate RNA-seq mapper with a partitioning strategy. Claims to be faster than other RNA-seq mappers at a higher or comparable sensitivity and accuracy. Speed is improved by using a "seed and cluster" approach to alignment instead of "seed and extend" (dynamic programming).
- <u>CCmiR: a computational approach for competitive and cooperative microRNA binding</u>
 <u>prediction</u>. Claims to be the only software that takes into account relative abundance of
 miRNAs and mRNAs for predicting binding sites.
- <u>SNPDelScore</u>: combining multiple methods to score deleterious effects of noncoding mutations in the human genome. Database of deleterious mutations and their deleteriousness. Provided by NCBI. Currently only for humans.

Cell

Current Biology

 Krupenye, C., Hare, B., n.d. Bonobos Prefer Individuals that Hinder Others over Those that Help. Current Biology 28, 280–286.e5. https://doi.org/10.1016/j.cub.2017.11.061

• <u>eLife</u>

 Ruby, J.G., Smith, M., Buffenstein, R., 2018. Naked mole-rat mortality rates defy Gompertzian laws by not increasing with age. eLife 7, e31157.
 https://doi.org/10.7554/eLife.31157

Evolution

Petri Rautiala, Heikki Helanterä, Mikael Puurtinen. <u>The evolutionary dynamics of adaptive virginity, sex-allocation, and altruistic helping in haplodiploid animals.</u> Doi: 10.1111/evo.13399.

• GBE

Genome Biol

- Michał Szpak et al <u>FineMAV</u>: prioritizing candidate genetic variants driving local adaptations in human populations https://doi.org/10.1186/s13059-017-1380-2
- Gerben Menschaert et al <u>The proBAM and proBed standard formats: enabling a seamless</u> integration of genomics and proteomics data https://doi.org/10.1186/s13059-017-1377-x

Genome Res

 Han et al. <u>SIDR: simultaneous isolation and parallel sequencing of genomic DNA and total</u> <u>RNA from single cells</u> doi: 10.1101/gr.223263.117

Insectes Sociaux

Journal of Evolutionary Biology

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 Voillemot et al (2018), <u>The divergence history of the perennial plant Linaria cavanillesii</u> confirms a recent loss of self-incompatibility. doi:10.1111/jeb.13209

MBE

- Mol Ecol
- Myrmecological News
- Nature
 - Moreno-Mayar et al. <u>Terminal Pleistocene Alaskan genome reveals first founding population</u> of Native Americans. doi:10.1038/nature25173.
 - Bakhoum et al. <u>Chromosomal instability drives metastasis through a cytosolic DNA</u> <u>response.</u> doi:10.1038/nature25432.
- Nature Comms
- Nature Genetics
 - Smith, J.J., Timoshevskaya, N., Ye, C., Holt, C., Keinath, M.C., Parker, H.J., Cook, M.E., Hess, J.E., Narum, S.R., Lamanna, F., Kaessmann, H., Timoshevskiy, V.A., Waterbury, C.K.M., Saraceno, C., Wiedemann, L.M., Robb, S.M.C., Baker, C., Eichler, E.E., Hockman, D., Sauka-Spengler, T., Yandell, M., Krumlauf, R., Elgar, G., Amemiya, C.T., 2018. The sea lamprey germline genome provides insights into programmed genome rearrangement and vertebrate evolution. Nature Genetics. https://doi.org/10.1038/s41588-017-0036-1
 - Hidden genetic variation shapes the structure of functional elements in Drosophila doi:10.1038/s41588-017-0010-v
 - Shared genetic effects on chromatin and gene expression indicate a role for enhancer priming in immune response doi:10.1038/s41588-018-0046-7
- Nature Methods
- Plos Biol
 - o Günther *et al.* Population genomics of Mesolithic Scandinavia: Investigating early postglacial migration routes and high-latitude adaptation. https://doi.org/10.1371/journal.pbio.2003703.
- Plos Comp Biol
- Plos Genet
 - Rohan Maddamsetti & Richard E. Lenski <u>Analysis of bacterial genomes from an evolution</u> <u>experiment with horizontal gene transfer shows that recombination can sometimes</u> <u>overwhelm selection https://doi.org/10.1371/journal.pgen.1007199</u>
 - Matt Ravenhall et al <u>Novel genetic polymorphisms associated with severe malaria and under selective pressure in North-eastern Tanzania</u> https://doi.org/10.1371/journal.pgen.1007172

PNAS

 Evolutionary transition from blood feeding to obligate nonbiting in a mosquito https://doi.org/10.1073/pnas.1717502115

• Proc B

- Blumstein, D.T., Williams, D.M., Lim, A.N., Kroeger, S., Martin, J.G.A., 2018. Strong social relationships are associated with decreased longevity in a facultatively social mammal. Proc Biol Sci 285. https://doi.org/10.1098/rspb.2017.1934
- <u>Science</u>
- TREE
- Trends in Genetics
 - Sarah Elizabeth Albritton, Sevinç Ercan. <u>Caenorhabditis elegans Dosage Compensation:</u>
 Insights into Condensin-Mediated Gene Regulation. https://doi.org/10.1016/j.tig.2017.09.010.
- Other papers/tools that you feel someone should have seen:
 - o BioRxiv
 - o Genetics

- o Heredity
- o Aging
- o Mol Cell Biol
- o J Gerontol
- o Methods in Ecology and Evolution
- o Current Opinion in Insect Science