

06.02.2017	time	student	paper
	11:00-11:20	María del Rocío Chamorro González	Dopico XC, Evangelou M, Ferreira RC, Guo H, Pekalski ML, et al. (2015) Widespread seasonal gene expression reveals annual differences in human immunity and physiology. <i>Nat Commun</i> 6: 7000.
	11:20-11:40	Nazli Güllü	Sancar A, Lindsey-Boltz LA, Kang TH, Reardon JT, Lee JH, et al. (2010) Circadian clock control of the cellular response to DNA damage. <i>FEBS Lett</i> 584: 2618-2625.
	11:40-12:00	Yi Bei	Wu S, Powers S, Zhu W, Hannun YA (2016) Substantial contribution of extrinsic risk factors to cancer development. <i>Nature</i> 529: 43-47.
	12:00-12:20	David Pride	Yu X, Rollins D, Ruhn KA, Stubblefield JJ, Green CB, et al. (2013) TH17 cell differentiation is regulated by the circadian clock. <i>Science</i> 342: 727-730.
BREAK			
	13:40-14:00	Eleonora Usatikova	Gordan R, Shen N, Dror I, Zhou T, Horton J, et al. (2013) Genomic regions flanking E-box binding sites influence DNA binding specificity of bHLH transcription factors through DNA shape. <i>Cell Rep</i> 3: 1093-1104.
	14:00-14:20	Sakurako Wong	Hu Y, Shmygelska A, Tran D, Eriksson N, Tung JY, et al. (2016) GWAS of 89,283 individuals identifies genetic variants associated with self-reporting of being a morning person. <i>Nat Commun</i> 7: 10448.
	14:20-14:40	Vahid Asimi	Koike N, Yoo SH, Huang HC, Kumar V, Lee C, et al. (2012) Transcriptional architecture and chromatin landscape of the core circadian clock in mammals. <i>Science</i> 338: 349-354.
	14:40-15:00	Alžbita Ressnerová	Archer SN, Laing EE, Moller-Levet CS, van der Veen DR, Bucca G, et al. (2014) Mistimed sleep disrupts circadian regulation of the human transcriptome. <i>Proc Natl Acad Sci U S A</i> 111: E682-691.
BREAK			
	15:20-15:40	Marta Hernández Justicia	Wallach T, Schellenberg K, Maier B, Kalathur RK, Porras P, et al. (2013) Dynamic circadian protein-protein interaction networks predict temporal organization of cellular functions. <i>PLoS Genet</i> 9: e1003398.
	15:40-16:00	Pooja Joshi	Plocik AM, Graveley BR (2013) New insights from existing sequence data: generating breakthroughs without a pipette. <i>Mol Cell</i> 49: 605-617.
	16:00-16:20	Shailey Twamley	He E, Kapuy O, Oliveira RA, Uhlmann F, Tyson JJ, et al. (2011) System-level feedbacks make the anaphase switch irreversible. <i>Proc Natl Acad Sci U S A</i> 108: 10016-10021.
07.02.2017			
	9:00-9:20	Christoforos Dimitropoulos	Hawkins RD, Larjo A, Tripathi SK, Wagner U, Luu Y, et al. (2013) Global chromatin state analysis reveals lineage-specific enhancers during the initiation of human T helper 1 and T helper 2 cell polarization. <i>Immunity</i> 38: 1271-1284.
	9:20-9:40	Gerard Arrey Tané	Hendriks GJ, Gaidatzis D, Aeschmann F, Grosshans H (2014) Extensive oscillatory gene expression during <i>C. elegans</i> larval development. <i>Mol Cell</i> 53: 380-392
	9:40-10:00	Trendelina Rrustemi	Xu Y, Ma P, Shah P, Rokas A, Liu Y, et al. (2013) Non-optimal codon usage is a mechanism to achieve circadian clock conditionality. <i>Nature</i> 495: 116-120.
	10:00-10:20	Kyung Hwan Lee	Thaiss CA, Levy M, Korem T, Dohnalova L, Shapiro H, et al. (2016) Microbiota Diurnal Rhythmicity Programs Host Transcriptome Oscillations. <i>Cell</i> 167: 1495-1510 e1412.
BREAK			
	10:40-11:00	Minze Xu	Schwanhauser B, Busse D, Li N, Dittmar G, Schuchhardt J, et al. (2011) Global quantification of mammalian gene expression control. <i>Nature</i> 473: 337-342.
	11:00-11:20	Ugne Dubonyte	Eden E, Geva-Zatorsky N, Issaeva I, Cohen A, Dekel E, et al. (2011) Proteome half-life dynamics in living human cells. <i>Science</i> 331: 764-768.
	11:20-11:40	Georgia Lattanzi	Klinger B, Sieber A, Fritsche-Guenther R, Witzel F, Berry L, et al. (2013) Network quantification of EGFR signaling unveils potential for targeted combination therapy. <i>Mol Syst Biol</i> 9: 673.
	11:40-12:00	Ángel Gil Nolskog	Lauriola M, Enuka Y, Zeisel A, D'Uva G, Roth L, et al. (2014) Diurnal suppression of EGFR signalling by glucocorticoids and implications for tumour progression and treatment. <i>Nat Commun</i> 5: 5073.