

| Journal | Title | PMID | Citation |
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| <i>Cell</i> | Genome-wide CRISPR Screens in Primary Human T Cells Reveal Key Regulators of Immune Function. | 30449619 | Shifrut E, Carnevale J, Tobin V, Roth TL, Woo JM, Bui CT, Li PJ, Diolaiti ME, Ashworth A, Marson A. Genome-wide CRISPR Screens in Primary Human T Cells Reveal Key Regulators of Immune Function. <i>Cell</i> . 2018 Nov 13. |
| <i>Nat Med</i> | A patient-level pooled analysis of treatment-shortening regimens for drug-susceptible pulmonary tuberculosis. | 30397355 | Imperial MZ, Nahid P, Phillips PPJ, Davies GR, Fielding K, Hanna D, Hermann D, Wallis RS, Johnson JL, Lienhardt C, Savic RM. A patient-level pooled analysis of treatment-shortening regimens for drug-susceptible pulmonary tuberculosis. <i>Nat Med</i> . 2018 Nov; 24(11):1708-1715. |
| <i>Nat Med</i> | Aurora kinase A drives the evolution of resistance to third-generation EGFR inhibitors in lung cancer. | 30478424 | Shah KN, Bhatt R, Rotow J, Rohrberg J, Olivas V, Wang VE, Hemmati G, Martins MM, Maynard A, Kuhn J, Galeas J, Donnella HJ, Kaushik S, Ku A, Dumont S, Krings G, Haringsma HJ, Robillard L, Simmons AD, Harding TC, McCormick F, Goga A, Blakely CM, Bivona TG, Bandyopadhyay S. Aurora kinase A drives the evolution of resistance to third-generation EGFR inhibitors in lung cancer. <i>Nat Med</i> . 2018 Nov 26. |
| <i>Nat Med</i> | Publisher Correction: A patient-level pooled analysis of treatment-shortening regimens for drug-susceptible pulmonary tuberculosis. | 30429542 | Imperial MZ, Nahid P, Phillips PPJ, Davies GR, Fielding K, Hanna D, Hermann D, Wallis RS, Johnson JL, Lienhardt C, Savic RM. Publisher Correction: A patient-level pooled analysis of treatment-shortening regimens for drug-susceptible pulmonary tuberculosis. <i>Nat Med</i> . 2018 Nov 14. |
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| <i>N Engl J Med</i> | AR101 Oral Immunotherapy for Peanut Allergy. | 30449234 | Vickery BP, Vereda A, Casale TB, Beyer K, du Toit G, Hourihane JO, Jones SM, Shreffler WG, Marcantonio A, Zawadzki R, Sher L, Carr WW, Fineman S, Greos L, Rachid R, Ibáñez MD, Tilles S, Assa'ad AH, Nilsson C, Rupp N, Welch MJ, Sussman G, Chinthrajah S, Blumchen K, Sher E, Spergel JM, Leickly FE, Zielen S, Wang J, Sanders GM, Wood RA, Cheema A, Bindslev-Jensen C, Leonard S, Kachru R, Johnston DT, Hampel FC, Kim EH, Anagnostou A, Pongracic JA, Ben-Shoshan M, Sharma HP, Stillerman A, Windom HH, Yang WH, Muraro A, Zubeldia JM, Sharma V, Dorsey MJ, Chong HJ, Ohayon J, Bird JA, Carr TF, Siri D, Fernández-Rivas M, Jeong DK, Fleischer DM, Lieberman JA, Dubois AEJ, Tsoumani M, Ciaccio CE, Portnoy JM, Mansfield LE, Fritz SB, Lanser BJ, Matz J, Oude Elberink HNG, Varshney P, Dilly SG, Adelman DC, Burks AW. AR101 Oral Immunotherapy for Peanut Allergy. <i>N Engl J Med</i> . 2018 11 22; 379(21):1991-2001. |
| <i>PLoS Med</i> | Enhancing the prediction of acute kidney injury risk after percutaneous coronary intervention using machine learning techniques: A retrospective cohort study. | 30481186 | Huang C, Murugiah K, Mahajan S, Li SX, Dhruva SS, Haimovich JS, Wang Y, Schulz WL, Testani JM, Wilson FP, Mena CI, Masoudi FA, Rumsfeld JS, Spertus JA, Mortazavi BJ, Krumholz HM. Enhancing the prediction of acute kidney injury risk after percutaneous coronary intervention using machine learning techniques: A retrospective cohort study. <i>PLoS Med</i> . 2018 Nov; 15(11):e1002703. |

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| <i>PLoS Pathog</i> | Gut and blood differ in constitutive blocks to HIV transcription, suggesting tissue-specific differences in the mechanisms that govern HIV latency. | 30440043 | Telwatte S, Lee S, Somsouk M, Hatano H, Baker C, Kaiser P, Kim P, Chen TH, Milush J, Hunt PW, Deeks SG, Wong JK, Yukl SA. Gut and blood differ in constitutive blocks to HIV transcription, suggesting tissue-specific differences in the mechanisms that govern HIV latency. <i>PLoS Pathog.</i> 2018 Nov; 14(11):e1007357. |
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