[14] PANDA A C, DE S, GRAMMATIKAKIS I, MUNK R, YANG X, PIAO Y, et al. High-purity circular RNA isolation method (RPAD) reveals vast collection of intronic circRNAs[J/OL]. Nucleic Acids Res, 2017, 45: e116. doi: 10.1093/nar/gkx297.

[15] MEMCZAK S, JENS M, ELEFSINIOTI A, TORTI F, KRUEGER J, RYBAK A, et al. Circular RNAs are a large class of animal RNAs with regulatory potency[J]. Nature, 2013, 495: 333-338.

[16] HANSEN T B, JENSEN T I, CLAUSEN B H, BRAMSEN J B, FINSEN B, DAMGAARD C K, et al. Natural RNA circles function as efficient microRNA sponges[J]. Nature, 2013, 495: 384-388.

[17] SCHNEIDER T, HUNG L H, SCHREINER S, STARKE S, ECKHOF H, ROSSBACH O, et al. CircRNA-protein complexes: IMP3 protein component defines subfamily of !! circRNPs[J/OL]. Sci Rep, 2016, 6: 31313. doi:10.1038/ !! srep 31313. ! !!! ! !!!

[18] ASHWAL-FLUSS R, MEYER M, PAMUDURTI N, IVANOV A, BARTOK O, HANAN M, et al. CircRNA biogenesis competes with pre-mRNA splicing[J]. Mol Cell, 2014, 56: 55-66.

[19] ZHU X, WANG X, WEI S, CHEN Y, CHEN Y, FAN X, et al. Hsa_circ_0013958: a circular RNA and potential novel biomarker for lung adenocarcinoma[J]. FEBS J, 2017, 284: 2170-2182.

[20] ZHAO J, LI L, WANG Q, HAN H, ZHAN Q, XU M. CircRNA expression profile in early stage lung adenocarcinoma patients[J]. Cell Physiol Biochem, 2017, 44: 2138-2146.

[21] JIANG M M, MAI Z T, WAN S Z, CHI Y M, ZHANG X, SUN B H, et al. Microarray profiles reveal that circular RNA hsa_circ_0007385 functions as an oncogene in non-small cell lung cancer tumorigenesis[J]. J Cancer Res Clin Oncol, 2018, 144: 667-674.

[22] ZHANG S, ZENG X, DING T, GUO L, LI Y, OU S, et al. Microarray profile of circular RNAs identifies hsa_circ_0014130 as a new circular RNA biomarker in non-small cell lung cancer[J/OL]. Sci Rep, 2018, 8: 2878. doi: 10.1038/S41598-21300-5.

[23] YAO J T, ZHAO S H, LIU Q P, LV M Q, ZHOU D X, LIAO Z J, et al. Over-expression of circRNA_100876 in non-small cell lung cancer and its prognostic value[J]. Pathol Res Pract, 2017, 213: 453-456.

[24] LUO Y H, ZHU X Z, HUANG K W, ZHANG Q, FAN Y X, YAN P W, et al. Emerging roles of circular RNA hsa_circ_0000064 in the proliferation and metastasis of lung cancer[J]. Biomed Pharmacother, 2017, 96: 892-898.

[25] HUANG Z K, YAO F Y, XU J Q, DENG Z, SU R G, PENG Y P, et al. Microarray expression profile of circular RNAs in peripheral blood mononuclear cells from active tuberculosis patients[J]. Cell Physiol Biochem, 2018, 45: 1230-1240.

[26] Sð7 •, Sð#1!`,4ƒ#1, 7ß8 , ™ Ò Ÿ,¼ &L~ˌ0u. =l"r(Û 'â RNA hsa_circ_0009024 =”Dê T5æ3ÿ d@ö*Ã Y+° À*T # h[J]. Y z ìO¸ g ’ n ,2018,41:399-404.

[27] HIRAI J, SAKANASHI D, HAGIHARA M, HARANAGA S, UECHI K, KATO H, et al. Bacteremia due to *Streptococcus tigurinus*: a case report and literature review[J]. J Infect Chemother, 2016, 22: 762-766.

[28] WAN T W, KHOKHLOVA O E, IWAO Y, HIGUCHI W, HUNG W C, REVA I V, et al. Complete circular genome sequence of successful ST8/SCC*mec*Ⅳ community-associated methicillin-resistant *Staphylococcus aureus* (OC8) in Russia: one-megabase genomic inversion, IS256’s spread, and evolution of Russia ST8-Ⅳ[J/OL]. PLoS One, 2016, 11: e0164168. doi: 10.1371/journal.pone.0164168.

[29] ZEHR E S, BAYLES D O, BOATWRIGHT W D, TABATABAI L B, REGISTER K B. Complete genome sequence of *Ornithobacterium rhinotracheale* strain ORT-UMN 88[J/OL]. Stand Genomic Sci, 2014, 9: 16. doi: 10.1186/1944-3277-9-16.

[30] BLASI B, TAFER H, TESEI D, STERFLINGER K. From glacier to sauna: RNA-Seq of the human pathogen black fungus *Exophiala dermatitidis* under varying temperature conditions exhibits common and novel fungal response[J/OL]. PLoS One, 2015, 10: e0127103. doi: 10.1371/journal.pone.0127103.

[31] ; BFý ,EÜ t o , z%ß7û w' ,'ð ’ , DL^ ± ,0u z!S S5æ Ô 65P · ;TL5æ+°(Û'â RNA =”DêA]-@/¢[J/CD]. Y z5æF *ê*ñ n (*a|t),2015,8:34-38.

[32] WANG J, ZHU M C, KALIONIS B, WU J Z, WANG L L, GE H Y, et al. Characteristics of circular RNA expression in lung tissues from mice with hypoxia-induced pulmonary hypertension[J]. Int J Mol Med, 2018, 42: 1353-1366.

[33] ™ ³ Ã,Cs I, DB±,‹!d6 ,] ,, ½%F,0u. Ö(Û(Û'â RNA hsa_circ_0029642 : < æ t U S ï6;*ñ5æ Ô65P ·+° Ÿ2'[J]. 0X 5 Ç g S ’ ’ Ñ,2017,39:1005-1008.

[本文编辑] e 8b