

遺伝子診療研究部 論文(2018年)

【英語論文】

原著論文

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2. Hatakeyama K, Nagashima T, Urakami K, Ohshima K, Serizawa M, Ohnami S, Shimoda Y, Ohnami S, Maruyama K, Naruoka A, Akiyama Y, Kusuhara M, Mochizuki T, Yamaguchi K. tumor mutational burden analysis of 2,000 Japanese cancer genomes using whole exome and targeted gene panel sequencing. *Biomed Res.* 2018;39(3):159-167.