



Qing Zhong

Kontakt

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Publikationen (4)

Zhu Y, Christiansen A, Fritz C, Rupp N, Poyet C, Rushing E, Weller M, Roth P, Haralambieva E, Hofer S, Chen C, Jochum W, Gao X, Teng X, Chen L, Zhong Q, Wild P, Aebersold R, Ljubicic J, Rutishauser D, Schmid M, Weiss T, Zhang Q, Sun R, Wang B, Yi X, Wu Z, Gao H, Cai X, Ruan G, Zhu T, Xu C, Lou S, Yu X, Gillet L, Blattmann P, Saba K, Fankhauser C, Guo T. High-throughput proteomic analysis of FFPE tissue samples facilitates tumor stratification. *Mol Oncol* 2019; 13:2305–2328.

Guo T, Aebersold R, Wild P, Poyet C, Saba K, Fankhauser C, Jochum W, Rueschoff J, Wagner U, Wong C, Charmpi K, Rupp N, Zhong Q, Li L, Beyer A. Multi-region proteome analysis quantifies spatial heterogeneity of prostate tissue biomarkers. *Life Sci Alliance* 2018; 1

Zhong Q, Diebold J, McKee T, Jochum W, Kashofer K, Hofman P, Zischka M, Moch H, Rechsteiner M, Rogel U, Vassella E, Wagner U, Kurt H, Molinari F, Cathomas G, Komminoth P, Barman-Aksözen J, Schneider-Yin X, Rey J, Wild P. Multi-laboratory proficiency testing of clinical cancer genomic profiling by next-generation sequencing. *Pathol Res Pract* 2018; 214:957–963.

Zhong Q, Poyet C, Blattner M, Soldini D, Moch H, Rubin M, Noske A, Rüschoff J, Haffner M, Jochum W, Perner S, Buhmann J, Rüschoff J, Guo T, Gabrani M, Schöffler P, Rechsteiner M, Liu Y, Fuchs T, Rupp N, Fankhauser C, Wild P. Image-based computational quantification and visualization of genetic alterations and tumour heterogeneity. *Sci Rep* 2016; 6:24146.

Projekte (0)

Keine Resultate gefunden.

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