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## **Comprehensive Pan-Genomic Characterization of Adrenocortical Carcinoma**

Zheng, Siyuan ; Cherniack, Andrew D ; Dewal, Ninad ; Moffitt, Richard A ; Danilova, Ludmila ; et al ; Beuschlein, Felix

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## Comprehensive Pan-Genomic Characterization of Adrenocortical Carcinoma

Siyuan Zheng, Andrew D. Cherniack, Ninad Dewal, Richard A. Moffitt, Ludmila Danilova, Bradley A. Murray, Antonio M. Lerario, Tobias Else, Theo A. Knijnenburg, Giovanni Ciriello, Seungchan Kim, Guillaume Assie, Olena Morozova, Rehan Akbani, Juliann Shih, Katherine A. Hoadley, Toni K. Choueiri, Jens Waldmann, Ozgur Mete, A. Gordon Robertson, Hsin-Ta Wu, Benjamin J. Raphael, Lina Shao, Matthew Meyerson, Michael J. Demeure, Felix Beuschlein, Anthony J. Gill, Stan B. Sidhu, Madson Q. Almeida, Maria C.B.V. Fragoso, Leslie M. Cope, Electron Kebebew, Mouhammed A. Habra, Timothy G. Whitsett, Kimberly J. Bussey, William E. Rainey, Sylvia L. Asa, Jérôme Bertherat, Martin Fassnacht, David A. Wheeler, The Cancer Genome Atlas Research Network, Gary D. Hammer, Thomas J. Giordano,\* and Roel G.W. Verhaak\*

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Due to a miscommunication during the process of transferring this manuscript from our editorial team to Production, the TCGA network authors were not properly indexed in PubMed. This has now been corrected online. We apologize for any confusion or inconvenience that this oversight might have caused.

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## DNMT3A Loss Drives Enhancer Hypomethylation in FLT3-ITD-Associated Leukemias

Liubin Yang, Benjamin Rodriguez, Allison Mayle, Hyun Jung Park, Xueqiu Lin, Min Luo, Mira Jeong, Choladda V. Curry, Sang-Bae Kim, David Ruau, Xiaotian Zhang, Ting Zhou, Michael Zhou, Vivienne I. Rebel, Grant A. Challen, Berthold Göttgens, Ju-Seog Lee, Rachel Rau, Wei Li,\* and Margaret A. Goodell\*

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During figure preparation for this paper, the authors inadvertently duplicated the flow plot of Figure 5D in Figure 5B and duplicated the legend of Figure 5B in Figure 5D. Figure 5 has been corrected online and is shown here, along with the corrected panel (B) legend. The authors regret this error and apologize for any confusion that it may have caused to the readers.

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