

[Epigenomic analysis detects aberrant super-enhancer DNA methylation in human cancer](#)

URL: <http://genomebiology.com/2016/17/1/11http://link.springer.com/content/pdf/10.1186/s13059-016-0879-2>

Authors: [Heyn, Holger](#) / [Vidal, Enrique](#) / [Ferreira, Humberto](#) / [Vizoso, Miguel](#) / [Sayols, Sergi](#) / [Gomez, Antonio](#) / [Morán, Sebastián](#) / [Boque-Sastre, Raquel](#) / [Guil, Sonia](#) / [Martinez-Cardus, Anna](#) / [Lin, Charles](#) / [Royo, Romina](#) / [Sanchez-Mut, Jose](#) / [Martinez, Ramon](#) / [Gut, Marta](#) / [Torrents, David](#) / [Orozco, Modesto](#) / [Gut, Ivo](#) / [Young, Richard](#) / [Esteller, Manel](#)

Publication: Genome Biology

Barcelona Supercomputing Center - Centro Nacional de Supercomputación

Source URL (retrieved on 3 Jul 2024 - 23:19): <https://www.bsc.es/es/research-and-development/publications/epigenomic-analysis-detects-aberrant-super-enhancer-dna>