

Computational tools used for telomeric content

TelomereHunter

Methodology article | [Open Access](#) | [Published: 28 May 2019](#)

TelomereHunter – in silico estimation of telomere content and composition from cancer genomes

[Lars Feuerbach](#) , [Lina Sieverling](#), [Katharina I. Deeg](#), [Philip Ginsbach](#), [Barbara Hutter](#), [Ivo Buchhalter](#), [Paul A. Northcott](#), [Sadaf S. Mughal](#), [Priya Chudasama](#), [Hanno Glimm](#), [Claudia Scholl](#), [Peter Lichter](#), [Stefan Fröhling](#), [Stefan M. Pfister](#), [David T. W. Jones](#), [Karsten Rippe](#) & [Benedikt Brors](#)

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[Lina Sieverling](#), [Chen Hong](#), [Sandra D. Koser](#), [Philip Ginsbach](#), [Kortine Kleinheinz](#), [Barbara Hutter](#), [Delia M. Braun](#), [Isidro Cortés-Ciriano](#), [Ruibin Xi](#), [Rolf Kabbe](#), [Peter J. Park](#), [Roland Eils](#), [Matthias Schlesner](#), [PCAWG-Structural Variation Working Group](#), [Benedikt Brors](#), [Karsten Rippe](#), [David T. W. Jones](#), [Lars Feuerbach](#)  & [PCAWG Consortium](#)

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qMotif

- qMotif is an unpublished tool

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Nucleic Acids Research, Volume 46, Issue 10, 1 June 2018, Pages 4903–4918,
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