

TITAN: inference of copy number architectures in clonal cell populations from tumor whole-genome sequence data

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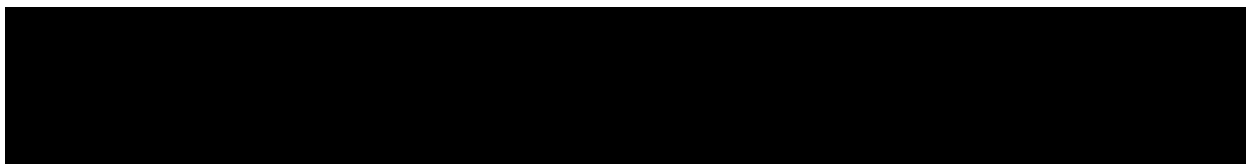
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