

@INPROCEEDINGS{6012857, author={de O Sandes, E. F. and de Melo, A. C. M. A.},
booktitle={Parallel Distributed Processing Symposium (IPDPS), 2011 IEEE International},
title={Smith-Waterman Alignment of Huge Sequences with GPU in Linear Space}, year={2011},
month=may, pages={1199-1211}, keywords={bioinformatics;cellular
biophysics;coprocessors;parallel algorithms;GPU;GTX 285 Board;Myers-Miller algorithm;Smith-
Waterman alignment;ancestral relationships;bioinformatics;cross-species chromosome
alignments;high performance computing platform;linear space complexity;parallel algorithm;species
peculiarity identification;Bioinformatics;Computer architecture;Graphics processing unit;Heuristic
algorithms;Instruction sets;Mathematical model;Microprocessors}, doi={10.1109/IPDPS.2011.114},
ISSN={1530-2075},}

From:

<https://www.cs.colostate.edu/AlphaZ/wiki/> - **AlphaZ**

Permanent link:

<https://www.cs.colostate.edu/AlphaZ/wiki/doku.php?id=melange:papers:spring2017&rev=1484759865>

Last update: **2017/01/18 10:17**

