

References

- [1] Markus J. Bauer, Anthony J. Cox, and Giovanna Rosone. Lightweight algorithms for constructing and inverting the bwt of string collections. *Theor. Comput. Sci.*, 483:134–148, 2013.
- [2] Dany Breslauer, Roberto Grossi, and Filippo Mignosi. Simple real-time constant-space string matching. *Theor. Comput. Sci.*, 483:2–9, 2013.
- [3] Daniel G. Brown and Jakub Truszkowski. Fast error-tolerant quartet phylogeny algorithms. *Theor. Comput. Sci.*, 483:104–114, 2013.
- [4] Domenico Cantone, Salvatore Cristofaro, and Simone Faro. Efficient string-matching allowing for non-overlapping inversions. *Theor. Comput. Sci.*, 483:85–95, 2013.
- [5] M. Christou, M. Crochemore, C.S. Iliopoulos, M. Kubica, S.P. Pissis, J. Radoszewski, W. Rytter, B. Szreder, and T. Waleń. Efficient seed computation revisited. *Theor. Comput. Sci.*, 483:171–181, 2013.
- [6] Raphaël Clifford, Markus Jalsenius, Ely Porat, and Benjamin Sach. Space lower bounds for online pattern matching. *Theor. Comput. Sci.*, 483:68–74, 2013.
- [7] David E. Daykin, Jacqueline W. Daykin, and W.F. Smyth. A linear partitioning algorithm for hybrid lyndons using v -order. *Theor. Comput. Sci.*, 483:149–161, 2013.
- [8] Riccardo Dondi, Guillaume Fertin, and Stéphane Vialette. Finding approximate and constrained motifs in graphs. *Theor. Comput. Sci.*, 483:10–21, 2013.
- [9] Michael Elberfeld, Danny Segev, Colin R. Davidson, Dana Silverbush, and Roded Sharan. Approximation algorithms for orienting mixed graphs. *Theor. Comput. Sci.*, 483:96–103, 2013.
- [10] Paolo Ferragina. On the weak prefix-search problem. *Theor. Comput. Sci.*, 483:75–84, 2013.
- [11] Travis Gagie, Juha Kärkkäinen, Gonzalo Navarro, and Simon J. Puglisi. Colored range queries and document retrieval. *Theor. Comput. Sci.*, 483:36–50, 2013.

- [12] Tomohiro I, Shunsuke Inenaga, and Masayuki Takeda. Palindrome pattern matching. *Theor. Comput. Sci.*, 483:162–170, 2013.
- [13] Sebastian Kreft and Gonzalo Navarro. On compressing and indexing repetitive sequences. *Theor. Comput. Sci.*, 483:115–133, 2013.
- [14] Stefanie Schirmer and Robert Giegerich. Forest alignment with affine gaps and anchors, applied in rna structure comparison. *Theor. Comput. Sci.*, 483:51–67, 2013.
- [15] Chris Thachuk. Compressed indexes for text with wildcards. *Theor. Comput. Sci.*, 483:22–35, 2013.