



Figure 3. Averaged over 5 runs of each step computation time for both algorithms for the Ionosphere dataset. For  $C = 100$ , we have the largest computation time since in such case there is no warm-start.

Table 2. Averaged time computation (in seconds) of an approximated regularization path with 100 samples of  $C$  varying from 0.1 to 100. We compare the performance of the MKL SILP algorithm and our algorithm.

data	SILP	Adaptive	Ratio
Liver	$612 \pm 180$	$116 \pm 19$	5.3
Pima	$4496 \pm 730$	$754 \pm 44$	6.0
Credit	$4830 \pm 754$	$753 \pm 64$	6.4
Ionosphere	$5000 \pm 788$	$221 \pm 22$	22.6
Sonar	$11234 \pm 2928$	$170 \pm 25$	66.1

ing more kernels than the SILP MKL algorithm. However, we have empirically showed that our algorithm is more stable in the kernel weights and thus it needs fewer iterations to converge towards a reasonable solution, making it globally faster than the SILP algorithm.

Future works aim at improving both speed and sparsity in kernels of the algorithm (for instance by initializing the SILP framework with the results of our algorithm) and by extending this algorithm to other SVM algorithm such as the one-class or the regression SVM.

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