

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.

da <del>ta</del> –	SH	P	Adaptive	Retic
Liver	612	180	116# 19	<b>5</b> .3
Pilna	4496	730	7.1	6.0
Credit	4830	754	73% 84	<del>6</del> .4
Ionosphere	5000	789	261 22	-22.6
Sonai	11234 J	2928	170 1 25	<del>66</del> .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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