# Combining High Productivity and High Performance in Image Processing Using Single Assignment C on Multi-core CPUs and Many-core GPUs

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## ABSTRACT

In this paper the challenge of parallelization development of industrial high performance inspection systems is addressed concerning a conventional parallelization approach versus an auto-parallelized technique. Therefore, we introduce the functional array processing language Single Assignment C (SAC), which relies on a hardware virtualization concept for automated, parallel machine code generation for multicore CPUs and GPUs. Additional, software engineering aspects like programmability, productivity, understandability, maintainability and resulting achieved gain in performance are discussed from the point of view of a developer. With several illustrative benchmarking examples from the field of image processing and machine learning, the relationship between runtime performance and efficiency of development is analyzed.

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#### 1. INTRODUCTION

The landscape of parallel computing has substantially changed in the last years. It is not only obvious that "the future is parallel" but also current trends confirm that computing power through parallelism will be provided by many-core architectures.<sup>1</sup> General-purpose many-core architectures must conveniently support a wide range of programming styles and languages. If architectures prefer a particular model of parallel programming, they are not likely to become widely accepted, especially if such architectures require programming skills that probably overstrain the average programmer. Furthermore, the development of high performance applications on novel and ever-changing hardware environments like multi- and many-core systems, Graphics Processing Units (GPUs) or Field Programmable Gate Arrays (FPGAs) is cost- and time-intensive. Writing explicitly parallel code for each and any of these architectures for each and any relevant part of a software system in theory would yield the best possible performance, but is highly uneconomical.

What is needed is a convenient abstract language that supports automatic parallelization on different architectures without changing source code and robust performance benefits. Since the early days of computing programmers are used to work with high-level programming languages (e.g., Algol, APL, Pascal, Fortran, Lisp, C) to hide low-level details of the architecture. Software engineers neither should need to design and develop in unintuitive ways nor to deal with a variety of hardware and language details just to avoid design mistakes or bottlenecks or just to achieve an attractive speed-up. From the economical point of view, an approach that yields the desired performance with minimal effort will be preferred, particularly for real-time performance applications in industry as well as for less efficient hardware.

Such a high-level programming language is Single Assignment C (SAC) — a strict, purely functional programming language, offering the combination of high-level language constructs with the high performance of manually optimized low-level modules. SAC combines C/Matlab-style syntax; it is designed to support high-level multi-dimensional stateless array processing. The SAC compiler generates competitive code for homogeneous multi-core/multi-processor systems,<sup>2</sup> for many-core NVidia graphics accelerators<sup>3</sup> and for the MicroGrid chip multiprocessor architecture.<sup>4</sup>

In the field of quality inspection of textured surfaces, e.g., metal, foils, woven fabrics, we have to cope with high scanning speeds, a large amount of data to process, and a complex phenomenology of textures and defects. This requires the application of advanced cost-intensive algorithms of image processing as well as machine learning, the use of high-performance computational hardware like GPUs or multi-core systems and the exploitation of parallelization potentials. The analysis of the whole processing pipeline (image acquisition, preprocessing, feature extraction, registration, defect detection and classification) with standard languages regarding performance is a resource- and time-intensive challenge. Figure 1 gives an example of such a image processing pipeline.



Figure 1. Typical image processing pipeline

While the performance of the image acquisition part mainly depends on the selected hardware and communication interfaces, the performance of major parts of preprocessing and feature extraction can be computed in a well-predictable way, due to the fixed size of filter operations and the amount of data known in advance. Besides less computational intensive methods, e.g., thresholding, image arithmetic, etc., the application of more sophisticated preprocessing methods is indispensable, e.g., enhancement of faults with the anisotropic diffusion filter, however, it increases the execution time. Nevertheless, sophisticated preprocessing methods can reduce the complexity and hence the computational costs of high-level pattern recognition and classification methods. For example, at the beginning of the processing pipe, a typical low-level scenario uses local filter operations, acting, e.g., on 9x9 matrices, while afterwards global operations on the whole image data are applied, such as a registration with a reference model based on thousands of feature points. So far the processing steps are acting on a physical, appearance-based level which only depends on the image intensity values. Finally, defect candidates have to be identified, located and classified. This final high-level step heavily depends on parameters that are not coded within the image, e.g., the customer's judgment whether some product quality aspects can be accepted or have to be rejected. The complexity of the classification step correlates with the quality of the preprocessing on the one hand, i.e., the best possible enhancement of faults and elimination of noise, and with the complexity of the defect taxonomy on the other hand.

In this paper a comparison and benchmarking of different implementations of two major parts of the above introduced image processing pipeline is performed. One part is the investigation of the Perona-Malik Anisotropic Diffusion<sup>1</sup>) which has poor performance characteristics by default, and the other examined method is a classification by Support Vector Machines (SVMs).<sup>5</sup> After a short overview of related work in Section 2, we introduce the functional array language SAC (Section 3). Then a detailed explanation of the anisotropic diffusion and the the SVM (and the according parallelized versions) is given in Section 4. Afterwards, we compare the SAC optimization strategies (with and/or without GPU support) against those of the OPENCV2.3<sup>6</sup> in Section 5. Finally, in Section 6 a comparison of the different implementations concerning programmability, understandability, productivity, maintainability is given.

## 2. RELATED WORK

The industry standard for programming NVidia GPUs is CUDA.<sup>7</sup> CUDA is a vendor-specific, architecture-specific and, hence, very low-level API. It allows the experienced programmer to adapt a program to the architectural peculiarities of GPU processing and to achieve high performance, if programming effort is not a big concern. However, software engineering on this level of abstraction is both tedious and cumbersome. If CUDA marks one end of the spectrum of GPU programming, then SAC<sup>8</sup> marks the other. SAC programs are architectureagnostic – it is solely up to the compiler and runtime system to make efficient use of GPUs where and when they are present.<sup>3</sup> Our goal is to provide scientists whose areas of expertise lie elsewhere than in high-performance computing, with nearly the same program performance as if they had been written by a highly skilled computer programmer. Analysis of that trade-off between performance and productivity is the subject of this paper.

In between CUDA and SAC, a number of other approaches aim at facilitating GPU programming. OpenCL,<sup>9</sup> originally proposed by Apple, is now promoted by AMD (the only major manufacturer of both multi-core CPUs and GPUs); in particular, AMDs upcoming Fusion architecture will soon combine both worlds on a single chip. OpenCL is only marginally more abstract than CUDA. Programmers defines computational kernels, which can be executed on different kinds of GPUs and even on multi-core CPUs. Instead of providing access to concrete architectural features, OpenCL abstracts them into a machine model that captures essential properties of today's GPU-enhanced computing systems across individual manufacturers and models. Nonetheless, to obtain high performance, OpenCL programmers must concern themselves with a variety of machine-level details that lower their productivity.

OpenMP<sup>10</sup> has a track record of facilitating programming of symmetric shared memory systems (multi-core, multi-processor) through compiler directives. The OpenMPC<sup>11</sup> project aims at generating CUDA code from eligible standard OpenMP directives. This approach is particularly attractive if application code is already equipped with OpenMP directives. Still, OpenMP is on a much lower abstraction level than SAC. We want to mention a recent proposal to extend OpenMP by clauses for the explicit placement of computations on the host or an a GPGPU.<sup>12</sup>

Last, but not least, HiCuda<sup>13</sup> is another approach to programming NVidia GPUs. based on compiler directives; it essentially imitates the OpenMP approach for symmetric multicores and proposes a tailor-made directive language for CUDA-enabled GPUs. Technically, HiCuda does simplify GPU programming, but it nonetheless exposes the same variety of architectural features as CUDA. Programmers need to make all relevant design decisions in application engineering, but can express them much more concisely than when using vanilla CUDA.

## 3. SINGLE ASSIGNMENT C

SAC is a purely functional programming language that, as far as possible, adopts a C-like notation to ease transition of programmers with a background in imperative languages; the language core is a functional, side-effect-free, subset of ISO C; assignment sequences are treated as nested let-expressions, branches as conditional expressions and loops as tail-end recursive functions; details can be found in.<sup>8</sup> Despite the radically different underlying execution model (context-free substitution of expressions vs. step-wise manipulation of global state), all language constructs adopted from C exhibit exactly the operational behaviour expected by C programmers. This equivalence allows programmers to choose their favourite style of SAC code; meanwhile, the compiler exploits the benefits of SACs side-effect free semantics to provide advanced optimisations and automatic parallelisation.

On top of this language kernel SAC provides genuine support for processing truly multidimensional (see Figure 2) and truly stateless/functional arrays using a shape-generic style of programming. Any SAC expression evaluates to an array. Arrays may be passed between functions without restrictions. Array types include arrays of fixed shape, e.g. int[3,7], arrays of fixed rank, e.g. int[.,.], and arrays of any rank, e.g. int[\*]. The latter include scalars, which we, following APL, consider to be rank-0 arrays with an empty shape vector. For convenience and equivalence with C, we use int, rather than the equivalent int[], as a type notation for scalars. The hierarchy of array types induces a subtype relationship, and SAC supports function overloading with respect to subtyping.

SAC provides only a small set of built-in array operations. Essentially, there are primitives to retrieve data pertaining to the structure and contents of arrays, e.g. an array's rank (dim(array)) or its shape (shape(array)).

$\left(\begin{array}{ccc} 1 & 2 & 3 \end{array}\right)$	dim:	2
4 5 6	shape:	[3,3]
$\left(\begin{array}{ccc} 7 & 8 & 9 \end{array}\right)$	data:	[1,2,3,4,5,6,7,8,9]
	dim:	3
	shape:	[2,2,3]
4 5 6	data:	[1,2,3,4,5,6,7,8,9,10,11,12]
	dim:	1
$[\ 1,\ 2,\ 3,\ 4,\ 5,\ 6\ ]$	dim: shape:	1 [6]
$[\ 1,\ 2,\ 3,\ 4,\ 5,\ 6\ ]$	dim: shape: data:	1 [ 6 ] [1,2,3,4,5,6]
$[\ 1,\ 2,\ 3,\ 4,\ 5,\ 6\ ]$	dim: shape: data: dim:	1 [6] [1,2,3,4,5,6] 0
[1, 2, 3, 4, 5, 6]	dim: shape: data: dim: shape:	1 [ 6 ] [1,2,3,4,5,6] 0 [ ]

Figure 2. Truly multidimensional arrays in SAC and their representation by data vector, shape vector and rank scalar



Figure 3. Three-level hierarchy of array types: arrays of unknown dimensionality (AUD), arrays of known dimensionality (AKD) and arrays of known shape (AKS)

A selection facility provides access to individual elements or entire subarrays using a familiar square bracket notation: *array[idxvec*].

All aggregate array operations are specified using WITH-loop expressions, a SAC-specific array comprehension:

with {

( lower\_bound <= idxvec < upper\_bound) : expr;</pre>

}: genarray( shape, default)

Here, lower\_bound and upper\_bound denote expressions that must evaluate to integer vectors of equal length.

They define a rectangular index set of arbitrary dimension. The identifier *idxvec* represents elements of this set, similar to loop variables in FOR-loops. However, we deliberately do not define any order on these index sets. Hence, a WITH-loop essentially specifies a FORALL loop nest. We call the specification of such an index set a *generator* and associate it with some potentially complex SAC expression. Thus, we create a mapping between index vectors and values, in other words an array. As an example, consider the WITH-loop

1 with {
2 ([0,0] <= iv < [3,5]) : 42;
3 }: genarray([3,5], 0)</pre>

that defines a  $3 \times 5$  matrix with all elements set to 42. The scope of the index vector, *idxvec* (here named *iv*) is confined to the expression associated with the generator. The index vector can be used to access the current index location. For example, the WITH-loop

1	with {
2	$([0] \le iv \le [5]) : iv [0];$
3	}: genarray( [5], 0)

computes the vector [0,1,2,3,4]. Note that iv denotes a 1-element vector rather than a scalar. Therefore, we need to select the first (and only) element from iv to achieve the desired result. Actually, it is not the generator that defines the shape of the resulting array, but the first expression following the keyword genarray. So far, the two have always coincided, but for example

1 with {
2 ([1] <= iv < [4]) : 42;
3 }: genarray( [5], 0)</pre>

computes the vector [0,42,42,42,0]. SAC still creates a 5-element vector, but only the three inner elements are defined as 42; all others are set to the *default value*, which is given by the second expression following the key word genarray, in this case 0. Since the default expression is not within the scope of a generator, it has no access to the index. Hence, all array elements not covered by any generator are guaranteed to have the same value.

WITH-loops are not limited to a single generator. For example, the WITH-loop

```
2 ([1] \le iv < [4]) : 1;
3 ([3] \le iv < [5]) : 2;
4 }: genarray([6], 0)
```

defines the vector [0,1,1,2,2,0]. All elements of the resulting array still not covered by any of the generators are initialised with the value of the default expression, 0 in the example. Whenever the index sets defined by the various generators are not pairwise disjoint, the order of the generators matters: in the example the array's value at index location [3], which is covered by both generators is set to 2 rather than to 1, i.e., the last generator dominates.

SAC actually features several variants of WITH-loops. Let us assume we have named the array defined by the previous WITH-loop A. Then, the modarray-WITH-loop

1	with {
2	$([0] \le iv \le [3]) : 3;$
3	}: modarray( A)

computes the vector [3,3,3,2,2,0]. More precisely, it computes a new array that has the same shape as the existing array denoted by the expression following the key word modarray. The computation of those elements covered by one or more generators follows exactly the same pattern as in the case of genarray-WITH-loops, but the remaining elements are defined by the values of the corresponding elements in the referenced array rather than by a common default value. Further WITH-loop variants support the definition of reduction operations and strided index sets.

As a more complete example, consider the 2-dimensional, 5-point stencil relaxation shown in Figure 4. Here, a C-style FOR-loop implements iterative Jacobi-relaxation, while a SAC with-loop array comprehension defines a single relaxation step. The five nearest neighbour elements of the argument array **a** are selected using explicit index computations. The dots in the generator refer to the least and the greatest index vector of the argument array **a**, respectively. In conjunction with the less-than relational operator, the dots form a convenient way to define the set of all non-boundary indices of array **a**.

While with-loops can always be used to define application-specific array operations like the 5-point stencil relaxation in Figure 4, their primary purpose is to support the definition of rank- and shape-generic basic array processing building blocks, which we denote as the *principle of abstraction*. Those blocks, are then used to compose application-specific functions, following the *principle of composition* common in array and functional

```
double[.,.] relax (double[.,.] a, int iter)
1
2
   {
3
      for (i=0; i<iter; i++) {
       a = with {
4
              (. < iv < .) : 0.2 * (a[iv] + a[iv - [1,0]] + a[iv - [0,1]]
5
                                            + a[iv + [0,1]] + a[iv + [1,0]]);
6
7
            }: modarray( a);
8
     }
9
      return a;
10
  }
```

Figure 4. 2-dimensional 5-point stencil relaxation in SAC

```
double[*] step (double[*] a)
1
2 {
3
     b = a;
     for (d=0; d < dim(a); i++) {
4
5
       b += rotate(d, -1, a) + rotate(d, 1, a);
6
     }
7
     return b * (1.0 / tod(2*dim(a)+1));
8 }
9
   double[*] relax (double[*] a, int iter)
10
11
   {
     for (i=0; i<iter; i++) {
12
      a = step(a);
13
14
     }
15
     return a;
16
  }
```

Figure 5. Rank-generic nearest-neighbour stencil relaxation in  $\operatorname{SAC}$ 

langauges.

Figure 5 demonstrates how SAC code is engineered based on these principles. We first define a shape- and rank-invariant version of nearest-neighbour relaxation that, as before, is based on a sequential FOR-loop to implement a series of relaxation steps. The function **step** uses another FOR-loop over the rank (number of dimensions) of the argument array. In each dimension, we rotate the argument array by one element towards ascending indices and by one element towards descending indices and, eventually, add up all these arrays in an element-wise manner. Finally, we divide all elements of the resulting array by the number of additions, i.e. twice the rank of the argument array plus one (for the non-rotated argument array). The function **tod** implements conversion from integer to floating point numbers. It is worthwhile to note that all functions used in the definition of **step** (e.g. rotation and element-wise array arithmetic) are not built-in primitives of the SAC language, but are defined in the SAC standard library, and are based on WITH-loops.



Figure 6. Architecture of SAC compilation infrastructure

As the various examples demonstrate, SAC code is completely architecture-agnostic. The SAC compilation infrastructure **sac2c** exploits this fact to generate specific code for a variety of target hardware architectures from the same SAC source code, thus achieving *hardware virtualisation* from a software engineering perspective. Figure 6 illustrates this concept. Based on aggressively optimised sequential code<sup>14</sup> **sac2c** at the time of writing supports symmetric multicore multiprocessor systems <sup>15</sup>, the MicroGrid chip multiprocessor architecture<sup>4</sup> and NVidia GPGPUs.<sup>16</sup> Work is on-going to extend this list.

# 4. APPLICATIONS

We now demonstrate how SAC combines high productivity in software engineering with high performance in program execution, by means of two methods from the industrial inspection system introduced in Section 1. First, we present a brief theoretical introduction to the anisotropic diffusion filter of Perona-Malik,<sup>1</sup> and to the decision function of the single class support vector machine.<sup>17</sup> Then, we show our surprisingly simple and concise SAC implementations of these applications.

#### 4.1 Perona-Malik Anisotropic Diffusion

Essential factors for robust and reliable defect detection are the enhancement of defects, such as scratches or blowholes, and attenuation of environmental influences, e.g., irregular reflections, noise or dust. Defect enhancement is supported by the Perona-Malik anisotropic diffusion filter,<sup>1</sup> whose principal characteristic is to reduce noise and concurrently enhance higher contrast regions.

The formal definition of the Perona-Malik anisotropic diffusion filter is defined by introducing D(.,.) from Equation (1) with the boundary condition in Equation (2), where D(.,.) depends on the local derivative in Equation (3) and Equation (4).

$$\frac{\partial}{\partial t}\varphi = div(D\nabla\varphi),\tag{1}$$

with boundary condition

$$\varphi(.,.,0) = \varphi, \tag{2}$$

where D depends on the local derivatives. Perona-Malik propose two different derivates

$$D = \frac{1}{1 + \left(\frac{\|\nabla\varphi\|}{K}\right)^2} \tag{3}$$

and

$$D = \exp^{-\left(\|\nabla\varphi\|/K\right)^2} \tag{4}$$

where Equation (3) acts as a smoothing filter that suppresses fine (noisy) structures, while Equation (4) strengthens high contrast edges. For an illustration, see Figure 7 to Figure 9, where we can see that only the connected wide regions are left, whereas noise structure is largely removed. The use of the deviation in Equation (4) in Figure 9 shows us that beside the big deep scratch in the middle also fine, noisy, high contrast edges are left. Suppose that the parameters of the illustrated results in Figure 8 and Figure 9 are defined as follows: NITER is the number of iterations, which means how many times the filter should applied to the image, delta



Figure 7. stainless steel plate with

noise surface and scratch

----

Figure 8. Application of Equation (3)

on image of Figure 7; NITER=5;

lambda=1/3; kappa=10;



Figure 9. Application of Equation (4) on image of Figure 7; NITER=5; lambda=1/3; kappa=10;

defines the stepsize of iteration and kappa K is the gradient modulus that controls the sensitivity to the edges.

The data-independent characteristic of the anisotropic diffusion filter allows an objective performance analysis of manually coded, as well as automatically SAC generated, GPU code. We present benchmarking results in section 5; subsection 4.2 outlines the implementation details of Perona-Malik anisotropic diffusion in Single Assignment C.

## 4.2 Implementation Single Assignment C versus Matlab

We show an abridgment of our SAC as well as Matlab implementation of the anisotropic diffusion filter. The following comparison should give an impression of the similarity between SAC and Matlab syntax.
SAC Matlab

•	• Line(3): assign image dimension
• Line(3): apply the filter niter times	• Line(5): apply the filter niter times
•	• Line(8-9): zero padding around image
• Line(6-9): apply stencil operation	• Line(12-15): apply stencil operation
• Line(12-15): calculate conduction	• Line(18-21): calculate conduction
• Line(17-20): assemble image	• Line(23-27): assemble image
• Line(23): return result	• Line(30): return result

Beside of the assign of the image dimension (Matlab - Line(3)) and the zero padding around the image for stencil operation<sup>18</sup> (Matlab - Line(8-9)) the differences of the code syntax are not significant.

Listing 1. SAC implementation of Perona-Malik Anisotropic filter

```
1 float [.,.] executeAnisotropicFilterSAC (float [.,.] image, float kappa, float delta, int niter)
2 {
3
     for (i=0; i<niter; i++)
4
     {
       /* stencil operation */
\mathbf{5}
       deltaNorth = shift(0, -1, 0f, image) - image;
\mathbf{6}
       deltaSouth = shift(0, 1, 0f, image) - image;
7
       deltaEast = shift(1, -1, 0f, image) - image;
8
       deltaWest = shift(1, 1, 0f, image) - image;
9
10
11
       /* Conduction */
12
       condNorth = 1f / (1f + pow (deltaNorth / kappa, 2f));
       condSouth = 1f / (1f + pow (deltaSouth / kappa, 2f));
13
14
       condEast = 1f / (1f + pow (deltaEast / kappa, 2f));
       condWest = 1 f / (1 f + pow (deltaWest / kappa, 2 f));
15
16
       image += delta * ( condNorth * deltaNorth
17
18
                           + condSouth * deltaSouth
                           + condEast * deltaEast
19
20
                           + condWest * deltaWest)
21
     }
22
23
     return image:
24 }
```

Listing 2. Matlab implementation of Perona-Malik Anisotropic filter

```
1 function image = executeAnisotropicFilterMATLAB(image, niter, kappa, delta, option)
2 {
     [rows, cols] = size(im);
3
4
\mathbf{5}
     for i = 1:niter
6
\overline{7}
       % zero padding
       diffl = zeros(rows+2, cols+2);
8
       diffl(2:rows+1, 2:cols+1) = image;
9
10
11
       \% North, South, East and West differences
       deltaNorth = diffl(1:rows , 2:cols+1) - image;
12
       deltaSouth = diffl(3:rows+2, 2:cols+1) - image;
13
14
       deltaEast = diffl(2:rows+1, 3:cols+2) - image;
       deltaWest = diffl(2:rows+1, 1:cols) - image;
15
16
       % Conduction
17
       \texttt{condNorth} = 1./(1 + (\texttt{deltaN}/\texttt{kappa}).^2);
18
       condSouth = 1./(1 + (deltaS/kappa).^2);
19
       condEast = 1./(1 + (deltaE/kappa).^2);
20
```

```
21
       condWest = 1./(1 + (deltaW/kappa).^2);
22
23
       image = image + delta * ( condNorth .* deltaNorth
24
                                + condSouth .* deltaSouth
25
                                + condEast .* deltaEast
26
                                 + condWest
                                            .* deltaWest
27
                                );
28
      end
29
30
      return image:
```

# 4.3 Classification with One-Class Support Vector Machine

Support vector machines (SVM) are based on the concept of separating data of different classes by determining the optimal separating hyperplanes.<sup>19</sup> The main idea behind support vector machines - and their distinctness to other learning algorithms - is the method of *structural risk minimization*. Instead of optimizing the training error (which often leads to the problem of over-fitting), attention focuses on minimization of an *estimate of the test error*.<sup>5</sup> Due to that underlying generalization,SVMs have become widely used learning methods which provide state-of-the art solutions for various application areas, e.g. text categorization, texture analysis, and gene classification.

Typically, the SVM is a supervised learning algorithm working on two classes (binary classification, see also<sup>5</sup>). But for industrial quality inspection, where mostly large amounts of good samples are available and just a small fraction of possible defects are known, the application of an outlier-detection version has been proposed (one-class or single-class SVM, see<sup>17</sup> and<sup>20</sup>). The training of the one-class SVM (OC-SVM) relies only on one data class (positive samples) and tries to construct a hyperplane that separates the surface region containing data from the region containing no data. This is done by determining the hyperplane with maximal distance from the point of origin with all (or almost all) data points lying on the opposite side of the origin. For an illustration see figure 10.



Figure 10. One-Class SVM: Separation of data points and origin

During training, a decision function f is determined, which is positive on all (or most) given positive samples

and negative on the complement:

$$f(\mathbf{x}) = \operatorname{sgn}(\sum_{i} \alpha_{i} k(\mathbf{x}_{i}, \mathbf{x}) - \rho),$$
(5)

where  $\mathbf{x}$  is a new sample that needs to be classified. The kernel function k(.,.) can be seen as similarity measure between the new sample point and the support vectors  $\mathbf{x}_i$  (a sub-set of the good samples from training, describing the outer sphere of the data cluster). The parameter  $\rho$  (decision boundary) and the non-zero weights  $\alpha_i$  (of the corresponding support vector  $\mathbf{x}_i$ ) are determined during the training phase.

For further details on the determination of the parameters and support vectors, and on possible kernel choices (polynomial, Gaussian radial basis function, etc.)  $\sec^5$  and.<sup>21</sup>

For the following implementation in SAC and the benchmark tests (see Section 5), we use the Gaussian kernel

$$k(\mathbf{x}, \mathbf{x}') = e^{-\gamma ||\mathbf{x} - \mathbf{x}'||^2}$$

with  $\gamma = \frac{1}{\sigma^2}$ , where  $\sigma > 0$  is the spread of the radial basis function, is used.

Often image processing applications are time-critical systems, e.g. in-line process control, where speed can be a limiting factor for usability. So the most essential part is the speed-up of the classification step, therefore a parallelization of the above mentioned decision function (see Equation 5) was considered.

#### 4.4 Implementation in Single Assignment C

In this section, we implement the decision function of a single class support vector machine in SAC, see Listing 3. In the Gaussian kernel function (kernel\_rbf) the parameter x contains the candidates to be classified, and the parameter sv contains the trained support vectors. The second function, (decision\_oneclass\_rbf computes the classification for one data point, where alpha is the weight of the according support vector. The 3rd (overloaded) function (predict\_oneclass\_rbf), maps the 2nd function onto the whole data set, as demanded by equation 5. Both overloaded instances of the decision\_oneclass\_rbf function make use of SAC's *axis control notation*.<sup>22</sup> Abstracting from some complexities of WITH-loops, this notation maps an index variable (in both cases iv) to an index space that is derived from the shape of an array into which the index variable indices within the right hand side expression (e.g. data in the 2nd instance). That expression is evaluated for each legal index value and the resulting values laminated to form a new array of the same shape as the one that is indexed into.

Listing 3. SAC implementation of the decision function of support vector machine

<sup>1</sup> inline double kernel\_rbf(double[.] x, double[.] sv, double gamma)

```
2 {
3
    return exp(-gamma * sum((x-sv)*(x-sv)));
4 }
5
6 inline double decision_oneclass_rbf(double[.] data, double[.,.] svs, double[.] alpha,
7
                                        double rho, double gamma)
8
  {
     return sum({ iv \rightarrow alpha[iv] * kernel rbf(data, svs[iv], gamma)}) > rho ? 1.0 : 0.0;
9
10 }
11
12 double[.] predict_oneclass_rbf(double[.,.] data, double[.,.] svs, double[.] alpha,
13
                                   double rho, double gamma)
14 {
15
    return { iv -> decision_oneclass_rbf(data[iv], svs, alpha, rho, gamma) };
16 }
```

#### 5. BENCHMARKING

Runtime benchmarking depends heavily on which hardware specification is used; also, the selection of hardware is problem specific. Hence, for our test scenario, we use two different hardware environments, i.e., a DELL Precision<sup>™</sup>690 and a SONY VAIO<sup>™</sup>PCG-81112M laptop. The DELL Precision<sup>™</sup>690 has two separate Intel<sup>®</sup>Xeon<sup>®</sup>5060 with 3.2GHz, giving 8 cores in total, and 2GB full buffered DDR2 memory, and a NVIDIA GeForce 8800 Ultra graphic card. The SONY VAIO<sup>™</sup>PCG-81112M laptop has an Intel<sup>®</sup>Core<sup>™</sup>i7-740QM Processor, 8GB RAM, and an NVIDIA GeForce GT 425M graphic card. The NVIDIA GeForce 8800 Ultra has 128 streaming processors with a core frequency of 612 MHZ, memory frequency of 1080MHz, 786MB memory and a memory bandwidth of 103.7 GB/sec where the NVIDIA GeForce GTX 425M has 96 streaming processors with a core frequency of 1120 MHZ, memory frequency of 800MHz, up to 1024 MB memory and a memory bandwidth of 25.6 GB/sec.

## 5.1 Benchmarking Anisotropic Diffusion

The dimension of the input data for the anisotropic filter ranges from  $256 \times 256$  pixels to  $4096 \times 4096$  pixels, with pseudo-randomly generated 8-bit values between 0 and 255 since, in this example, only the dimension of the data affects execution time. Therefore, in Table 1 we present five different input sizes and propagate for each of them Equation (3) ten times. Furthermore, we implemented the filter in SAC with auto generated CPU-and CUDA-code; the CUDA version is manually optimized; the OpenCV2.3 framework is measured with and without GPU and Intel TBB support. Finally, a Matlab implementation is benchmarked as well.

Table 1 gives the benchmarking results achieved on the DELL Precision<sup>™</sup>690; Table 2 gives the same for the SONY VAIO<sup>™</sup>PCG-81112M. The first two rows compare CPU performance, where SAC-MT 1T denotes sequential execution and SAC-MT 8T denotes automatic parallelization to 8 cores. For all non-trivial problem sizes, we observe a speedup of about 5. This reduction in runtime is realized with no more development effort, but solely recompilation of the SAC source code. In industrial practice, this substantial performance improvement can be a time buffer for using more complex algorithms or giving a significant competitive advantage against other applications. Furthermore, the execution time of the auto-parallelized SAC-CUDA code compares favorably with manually optimized CUDA-code. It can be generally observed that GPU and TBB support of the OpenCV implementation has, in this use case, no impact to the overall performance. In general, the runtime of the Matlab implementation is not as bad as expected but, for larger input sizes (e.g., 4096x4096), it runs out of memory, resulting in disastrous execution times.

	image dimension				
	$px256 \times px256$	$px256 \times px256 \qquad px512 \times px512 \qquad px1024$		$px2048 \times px2048$	$px4096 \times px4096$
SAC-MT 1T	0.199 sec	0.788 sec	3.165 sec	12.55 sec	50.18 sec
SAC-MT 8T	0.067 sec	0.156 sec	0.635 sec	2.562 sec	10.18 sec
SAC-CUDA	0.003 sec 0.006 sec		0.015 sec	0.036 sec	0.181 sec
CUDA-manual	0.005 sec 0.007 sec		0.016 sec 0.050 sec		0.190 sec
OpenCV(1)	0.119 sec	0.540 sec	2.374 sec	9.571 sec	54.45 sec
OpenCV(2)	0.097 sec	0.550 sec	2.386 sec	9.507 sec	53.21 sec
OpenCV(3)	0.121 sec	0.512 sec	2.298 sec	9.532 sec	53.89 sec
OpenCV(4)	0.098 sec	0.498 sec	2.321 sec	9.532 sec	54.10 sec
Matlab	0.164 sec	0.608 sec	2.518 sec	10.06 sec	438.6 sec

Table 1. Runtime results of anisotropic filter benchmarked on a DELL Precision<sup> $^{14}$ </sup>690, SAC-MT 1T/8T = SAC on CPU executed with 1 and 8 threads, SAC-CUDA = SAC implementation, CUDA-manual = CUDA implementation, OpenCV(1) = OpenCV2.3v without CUDA and TBB support, OpenCV(2) = OpenCV2.3v with CUDA support, OpenCV(3) = OpenCV2.3v with TBB support, OpenCV(4) = OpenCV2.3v with CUDA and TBB support, Matlab = Matlab version 2011b.

But why do we not achieve a speedup of 8 with the CPU code? In fact, both experimental systems only feature 4 real cores which are twice hyperthreaded, but hyperthreading is not effective for this kind of workload. In this sense, the four-fold speedup is close to optimal. The functional programming paradigm results in a low memory usage of about 350 MB on average. However, if we need more performance for the application scenario, we have the possibility either to re-implement the whole algorithm with NVIDIAs CUDA framework or automatically generate executable GPU code with the SAC-CUDA backend. A CUDA re-implementation definitely requires higher development costs and programming know-how from experts, whereas SAC-CUDA allows flexible time and cost-efficient development.

In Table 2 are the benchmarking results achieved on the SONY VAIO<sup>™</sup>PCG-81112M, where the general performance on CPU is slightly better than on DELL Precision<sup>™</sup>690. Although the SONY VAIO<sup>™</sup>PCG-81112M

	image dimension				
	$px256 \times px256$	$px256 \times px256 \qquad px512 \times px512 \qquad px1024 \times px1024 \qquad px1024 \ px1024 \times px1024 \qquad px1024 \ px102$		$px2048 \times px2048$	$px4096 \times px4096$
SAC-MT 1T	0.094 sec	0.334 sec	1.334 sec	5.388 sec	20.08 sec
SAC-MT 8T	0.085 sec	0.183 sec	0.692 sec	2.805 sec	11.06 sec
SAC-CUDA	0.019 sec	0.046 sec	0.134 sec	0.456 sec	1.690 sec
CUDA-manual	0.039 sec	0.047 sec	0.077 sec	0.077 sec 0.186 sec	
OpenCV(1)	0.047 sec	0.171 sec	0.763 sec	3.011 sec	12.15 sec
OpenCV(2)	0.041 sec	0.178 sec	0.787 sec	3.321 sec	12.53 sec
OpenCV(3)	0.045 sec	0.175 sec	0.752 sec	3.211 sec	12.21 sec
OpenCV(4)	0.035 sec	0.172 sec	0.770 sec	3.078 sec	12.42 sec
Matlab	0.103 sec	0.369 sec	1.315 sec	4.034 sec	15.657 sec

Table 2. Runtime results of anisotropic filter benchmarked on a SONY VAIO<sup>TM</sup>PCG-81112M, SAC-MT 1T/8T = SAC on CPU executed with 1 and 8 threads, SAC-CUDA = SAC implementation, CUDA-manual = CUDA implementation, OpenCV(1) = OpenCV2.3v without CUDA and TBB support, OpenCV(2) = OpenCV2.3v with CUDA support, OpenCV(3) = OpenCV2.3v with TBB support, OpenCV(4) = OpenCV2.3v with CUDA and TBB support, Matlab = Matlab version 2011b.

has the newer graphic card, the benchmarks are significantly slower than on the DELL Precision<sup> $^{14}$ </sup>690, because of the lower hardware performance characteristic in the laptop.

#### 5.2 Benchmarking One-Class SVM

We now present a comparison of the parallelized versions of the decision function (see Equation (5)). First, the SAC-CUDA implementation, shown in Figure 3, is benchmarked. In addition to SAC runtime performance, we present results of the  $GPUSVM^{23}$  implementation and an OpenCV implementation compiled with GPU support, as well as Intel TBB. The manually optimized implementation of the GPU-based OC-SVM Classifier is based on a third-party *C-Support Vector Classification* implementation called  $GPUSVM^{23}$  For processing SVM data in parallel on GPU-devices, the applied classification algorithm employs Map Reduce<sup>24</sup> techniques proposed by Google as well as a GPU-vendor supplied *Basic Linear Algebra Subroutines (CUBLAS)*. The developed GPU-based OC-SVM classifier is able to read LIBSVM data format, hence, LIBSVM can be used for the training of the SVM models (and providing support vectors for it). Furthermore, all implementations can handle sparse matrices representation.

For the presented test results (shown in Table 3 and Table 4), some publicly available data sets were used from the LIBSVM data sets repository.<sup>25</sup> Since this data repository does not contain data sets for OC-SVMs, we took binary sets and generated training data sets with a certain size (300 samples), consisting of data belonging only to one class. A simplified training with the standard settings of LIBSVM was performed, using the Gaussian RBF kernel with  $\gamma = 1/n$  (where *n* is the number of features of the input vectors) and  $\nu = 0.5$ . For an explanation of these parameters see .<sup>5</sup>

	data sets $  \#$ of data points $  \#$ of features				
	a1a   30956   123   a9a   32561   123   australian   690   14   w8a   49				
SAC-MT 1T	9.484 sec	10.16 sec	$0.479  \sec$	$15.72  \sec$	
SAC-MT 8T	$1.709  \sec$	1.885  sec	$0.096  \sec$	$2.850  \sec$	
SAC-CUDA	0.921 sec 0.951 sec 0.051 sec 1.356				
CUDA-manual	$0.249  \sec$	0.253  sec	$0.187  \sec$	$0.295 \ sec$	
OpenCV(1)	6.241  sec	$6.611 \ sec$	$0.055  \sec$	21.92  sec	
OpenCV(2)	6.211  sec	6.211 sec 6.598 sec		21.76 sec	
OpenCV(3)	1.410 sec	1.488 sec	$0.017  \sec$	4.941 sec	
OpenCV(4)	1.417 sec 1.494 sec 0.017 sec 4.939 sec				

Table 3. Runtime results of decision function of single class support vector machine benchmarked on a DELL Precision  $^{16}690$ , SAC-MT 1T/8T = SAC on CPU executed with 1 and 8 threads, SAC-CUDA = SAC implementation, CUDA-manual = CUDA implementation, OpenCV(1) = OpenCV2.3v without CUDA and TBB support, OpenCV(2) = OpenCV2.3v with CUDA support, OpenCV(3) = OpenCV2.3v with TBB support, OpenCV(4) = OpenCV2.3v with CUDA and TBB support.

	data sets $  \# \text{ of data points }   \# \text{ of features}$				
	a1a   30956   123	a9a   32561   123	australian   690   14	w8a  49749   300	
SAC-MT 1T	6.379 sec	6.834 sec	0.248 sec	9.620 sec	
SAC-MT 8T	1.759 sec	1.865 sec	0.071 sec	3.097 sec	
SAC-CUDA	2.766 sec	2.949 sec	$0.051   {\rm sec}$	3.882 sec	
CUDA-manual	0.569 sec	0.613 sec	0.324 sec	0.794 sec	
OpenCV(1)	3.066 sec	3.281 sec	$0.037  \sec$	11.80 sec	
OpenCV(2)	3.045 sec	3.265 sec	0.035 sec	11.83 sec	
OpenCV(3)	1.284 sec	1.342 sec	0.014 sec	4.811 sec	
OpenCV(4)	1.274 sec	1.322 sec	0.015 sec	4.812 sec	

Table 4. Runtime results of decision function of single class support vector machine benchmarked on a SONY VAIO<sup>TM</sup>PCG-81112M, SAC-MT 1T/8T = SAC on CPU executed with 1 and 8 threads, SAC-CUDA = SAC implementation, CUDA-manual = CUDA implementation, OpenCV(1) = OpenCV2.3v without CUDA and TBB support, OpenCV(2) = OpenCV2.3v with CUDA support, OpenCV(3) = OpenCV2.3v with TBB support, OpenCV(4) = OpenCV2.3v with CUDA and TBB support.

In general, we can observe for the one-class SVM use case in Table 3 and Table 4 that we achieve a speedup for all multi-core implementations (i.e., SAC-MT 8T, OpenCV(3), OpenCV(4)) in opposite to single threaded execution (i.e., SAC-MT 1T, OpenCV(1), OpenCV(2)). Especially for the test data a1a, a9a, and australian the OpenCV implementation performs better than the SAC one. For the test data w8a the SAC implementation

on CPU showed the best runtime performance. This can be explained due to the high sparseness of the data matrix.

	sparseness of				
	data set features				
ala	88.72%	88.69%			
a9a	88.72%	88.69%			
australian	20.04%	13.59%			
w8a	96.11%	94.58%			

Table 5. Sparseness of the used data sets and features

In the OpenCV implementation the GPU support can be neglected because the used functions in the implementation provides no GPU support. The manual coded Cuda application can outperform the SAC-CUDA approximately 4.8 times, which mainly depends on the sparseness of the input matrices.

Concerning optimization strategies, we can say that OpenCV offers an optimized Streaming SIMD Extensions 2 (SSE2) code. SSE2 is a processor supplementary instruction set for modern 32-bit x86 and 64-bit x64 Single-Instruction, Multiple-Data (SIMD) architectures, where many of the basic arithmetic functions can run significantly faster. OpenCV also contains Intel®Threading Building Blocks (TBB)<sup>26</sup> support for several functions. TBB is a C++ template library which offers a complete threading mechanism on modern multi-core processors. The advantages of this library are easy and efficient handling ( application engineers do not need to be threading experts), scalable performance and a higher-level, task-based parallelism. In our example TBB is irrelevant as we do not use OpenCV functions that support this library. OpenCV applies the TBB only to OpenCV applications, e.g., haartraining, traincascade, and not to basic arithmetic/filter operations.

The optimization strategy of SAC is different. One of the major design principles of SAC is the WITH-loop construct, which supports the specification of shape-invariant array operations. All primitive array operations of SAC can be defined as WITH-loops within a standard library rather than being implemented as part of the compiler. The basic idea is to use WITH-loops as a universal representation for array operations and to develop a general transformation scheme that allows the concentration of individual with-loops into complex ones that exposes a more favorable computation to memory load/store ratio and reduces the need for synchronization and communication in parallel execution. Together, WITH-LOOP-FOLDING, WITH-LOOP-FUSION and WITH-loops<sup>27</sup> stepwise transform any nesting of primitive array operations into a single loop construct that contains an element-wise specification of the resulting array. During the compilation process various conventional optimization techniques,<sup>28,29</sup> such as function inlining, constant folding, constant propagation, loop unrolling, and dead code removal, are applied to produce efficiently executable code. This code is fully automatically parallelized using

POSIX<sup>™</sup> threads<sup>2,30</sup> or CUDA.<sup>3</sup> The net result is that application programmers can concentrate on the science of their problem area, rather than being forced to become experts in parallel programming or GPU programming.

#### 6. EXPERIENCES

This section evaluates various key values, i.e., programmability, understandability, productivity, maintainability, IDE support and CPU/GPU execution time, based on developer statements. Future development concerns make the choice of application implementation language a crucial decision. We investigated these issues by implementing our mentioned applications in C++/OpenCV, Matlab, CUDA and SAC taking care in software engineering aspects during the whole application development life cycle. As a starting point, C programming skills are rated as neutral to allow comparison of language characteristics with other tools, languages and frameworks.

			n no de oticites	maintaina hilita	IDE summent	execution time	
	programmability	understandability	7 productivity maintainability IDE support		CPU	GPU	
C++/OpenCV	0	0	0	0	0	0	0
Matlab	+	0	0	+	++		0
CUDA	-	-		0	+	0	++
SAC	0	0	++	++		0	+

Table 6. Pros and cons of applied tools, languages and frameworks regarding various application development aspects

Researcher, developer and application engineers have different needs and expectations for languages and tools, hence each language has more or less a similar *programmability* and *understandability*, because of existing assets and drawbacks in specific application fields, e.g., Matlab is a simple to use programming language, especially for rapid prototyping, but normally the developer has no knowledge about internal optimization strategies. For GPU development with CUDA, the developer needs special expertise in hardware architecture and parallelization techniques, and has to cope with a fast growing and changing technology. If the developer has good programming skills in C++/Matlab, SAC is easy to learn and provides the programming comfort of Matlab, e.g., no pointer arithmetic. Furthermore, SAC provides auto-parallelization and optimization over the whole application.

For rapid prototyping development, Matlab offers high *productivity*, our experience is that in several cases a re-design/re-implementation, using a more efficient language/framework in terms of runtime, is needed. This is often intensive work because of the unknown optimization strategies within Matlab: results vary and are not comparable to the other languages. By using SAC it is possible to auto-generate code for the mentioned platforms; this is especially useful if the performance criteria of a project have changed. SAC-code can fully automatically be compiled to multicore CPUs, manycore GPUs and the MicroGrid chip multiprocessor architecture. This offers high flexibility during a project's life cycle and it brings great advantages in *maintainability*. For example, upgrading from an NVIDIA GeForce 8800 to the new NVIDIA Fermi architecture only required re-compilation of the exact same source code. For the other languages, maintainability has basically a similar complexity.

A drawback of SAC is the lack of an integrated development environment (*IDE support*), which means that debugging, code analysis, benchmarking, etc., can only be done via the command line, whereas the other tools, languages and frameworks offer consistently well-engineered tool support, e.g., on GPU the profiling and debugging can be done via external tools and integrated MS Visual Studio plugins.

The execution time on CPU and GPU is influenced by several factors, e.g., hardware environment, parallelization and benchmarking strategies or concurrent production processes that primarily occur in industry. However, in general, SAC performance on CPUs is as good as the performance of C if no optimization framework is used (e.g., OpenCV, Intel-IPP, Intel-MKL, etc.). Typically, the highest performance can be achieved with manually written CUDA code (there are some exceptions) but in some cases SAC is able to surpass manually written CUDA code due to whole program optimization and consistently optimized parallelization strategies, especially for array-based algorithms. Furthermore, the design of complex parallel algorithms in SAC is easier than with CUDA; this often results in a bug-free and runtime-optimized application development.

## 7. CONCLUSION

In this paper, we showed the advantage of the functional array language Single Assignment C (SAC) in the field of image processing, particularly for the anisotropic diffusion filter and for the decision function of a single class support vector machine. Such a sophisticated filter operation can enhance faults and eliminate noise in multiiteration steps. This is computationally intensive, but indispensable to reduce the complexity, and consequently, the computational costs of high-level pattern recognition and classification methods. A single class support vector machine provides a robust and reliable classification for defect candidates which algorithmic characteristic allows a fine-grained parallelization and hence an optimal performance gain.

Due to industrial needs for balance between scalable and high-performance applications on the one hand and the demand for constant or lower development costs on the other hand, we conducted a benchmarking experiment in which we compared the development effort using SAC and the common image library OpenCV2.3v.

In terms of language syntax, SAC is similar to Matlab because of the definition of various Matlab-like operations. Furthermore, with SAC development time can be reduced by the well-known C/C++ semantics, yet offering side-effect free semantics, most notably due to the absence of pointers and hardware virtualization. The hardware virtualization allows flexible and fast development on architectures corresponding to CPUs, GPUs or FPGAs using the same language and the same implementation. Moreover, on multi/many-core architectures, as

well as on GPUs, SAC with auto-parallelization is often able to obtain higher performance than with the other languages. Additionally, from the economic point of view, SAC provides us with an extremely good balance between time of development and performance.

Although SAC is well-suited for image processing as well as array based algorithms because of data-parallelism and n-dimensional array support, it provides limited support for development and debugging tools. This will be changed in the future by an intensive enhancement of SAC and community building.

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