

New Conservation Functions and a Partial Taxonomy for 1-D Cellular Automata

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Abstract: We present algorithms that permit increased efficiency in the calculation of conservation functions for cellular automata, and report results obtained from implementations of these algorithms to report conservation laws for 1-D cellular automata of higher order than any previously known. We introduce the notion of trivial and core conservation functions to distinguish truly new conservation functions from simple extensions of lower-order ones. We give new theorems related to these concepts, and show our use of them to derive more efficient algorithms for finding conservation functions. We then present the complete list of conservation functions up to order 16 for the 256 elementary 1-d binary cellular automata. These include CAs that were not previously known to have nontrivial conservation functions.

I. INTRODUCTION

Conservation functions indicate fundamental physical properties of systems, and are therefore eagerly sought after in science. Conservation of energy and conservation of angular momentum, for example, are two fundamental principles at different physical levels that offer profound insights into the nature of the world we live in.

Similarly, conservation functions of living systems, when they can be found, offer insights into the evolution of the system over time. The simplest conservation functions indicate rules on the number of organisms, but more complex ones such as conservation of “energy”, where energy is suitably defined, suggest something about the inherent nature of the automaton under study. Cellular automata with identical conservation functions are, in some sense, closely related to one another. Thus conservation functions can be used to classify cellular automata, and to identify connections between seemingly unrelated systems.

The classification scheme we present here permits finer distinctions between CAs than merely their lowest order of conservation function (as has been done in the past) by grouping functions together with identical sets of basis vectors

and by distinguishing *core* functions from *simple extensions*. Of the 88 equivalent 1-dimensional cellular automata, 8 have provably trivial conservation functions only, 47 have conservation functions of various kinds, and 33 remain unclassified. Those that we have classified include CAs with newly discovered conservation functions of order 8, 9, 12, 13 and 14. Our calculations also show there are no CAs with conservation functions of order 15 or 16. We can therefore present here the complete classification scheme for 1-dimensional cellular automata based on conservation functions up through order 16.

II. TERMINOLOGY

A one-dimensional *cellular automaton* is an array of finite size W of cells containing $\{0, 1, \dots, s-1\}$. The array is called the *universe*, and the number of cells it contains is the size of the universe. The universe is considered to have no boundaries, which means in one dimension the end cells are considered adjacent to one another.

The current set of values in each cell is the *state* of the universe. This state changes over time based on a *characteristic function*, parameterized by a neighborhood size n . To determine the next state of any cell, the state of the $(n-1)/2$ cells on each side of the current cell, along with the state of the current cell itself is examined, and rules applied that uniquely determine the new state. (Since we are only interested in symmetric neighborhoods, n will always be odd). The new state of the universe as determined by the parallel application of the characteristic function is called the *successor state*. A 1-dimensional CA with s states and neighborhood of size n can be configured in s^n ways, not all of which are unique.

Let \mathbf{x} be the state of the universe at a given time, and \mathbf{x}_i be the state of cell i . An energy function $E(\mathbf{x})$ is defined as:

$$E(\mathbf{x}) = \sum_{i=0}^{n-1} f(\mathbf{x}_i, \mathbf{x}_{i+1}, \dots, \mathbf{x}_{i+m-1})$$

where subscripts outside the universe wrap around. The energy function is a sum of the energy of n different regions,

each of which contains m cells. We refer to m as the *order* of the function. Let $\text{succ}(\mathbf{x})$ be the successor state of the universe reached on the next time step if it starts in state \mathbf{x} . We say that E is conserved, or that E is an energy conservation function, iff, for all finite universes:

$$\text{Conserved: } \forall \mathbf{x} \quad E(\mathbf{x}) = E(\text{succ}(\mathbf{x}))$$

This is equivalent to the following [1]:

$$\forall \mathbf{x}, \mathbf{y} \quad E(\mathbf{x}) - E(\text{succ}(\mathbf{x})) = E(\mathbf{y}) - E(\text{succ}(\mathbf{y}))$$

where \mathbf{x} and \mathbf{y} are states that differ by exactly one cell.

It can be shown that for neighborhood size of $n=3$, we need only analyze the energy of a $2m+3$ -cell array and the $2m+1$ -cell array of its successor state in order to determine if a conservation law exists, despite the fact that the universe itself may be arbitrarily large. This is what makes the computation of conservation functions possible.

It is a theorem of Hattori and Takesue [1], later restated in simpler form in [2], that if we zero the block of cells in \mathbf{x} and \mathbf{y} either to the left or the right of the non-identical cell, our energy calculations will be unaffected. Thus applying this theorem over all possible $(m+1)$ -cell states of the universe for a given CA gives a system of $s^{(m+1)}$ equations in s^m unknowns, with the right hand side equal to zero. If this system of equations has a solution (that is, if the resulting matrix has a non-empty null space), then an energy conservation function of order m has been found. If it does not, then no such function exists for that CA.

Conservation laws are fundamental properties eagerly sought in studies of physical phenomena. Conservation functions are equally important for mathematical objects because they say something fundamental about the entity under study. Cellular automata can be classified by the order of their conservation functions, and can be understood more deeply by the nature of the quantities conserved. CAs are simple objects that generate surprisingly complex behavior. Conservation functions can shed light on how such behavior might emerge.

III. TRIVIAL, NON-TRIVIAL, AND CORE ENERGY CONSERVATION FUNCTIONS

Clearly energy functions that assign the same value to all states are conserved. We call such functions *trivial*. Of greater interest are non-trivial energy conservation functions. Trivial functions can be eliminated from the solution space through the addition of a small number of additional constraints on the system, using the following two theorems [3].

Theorem 1: For a 1D, 2-state, neighborhood 3 cellular automaton, the following set of 2^{m-1} trivial energy functions form a basis set for all possible trivial energy functions over m bits, where $1 < S < 2^{m-1}$. and the expression $0S$ ($1S$) represents a 0 (1) followed by the bits of S in binary:

$$\begin{aligned} f_0(\mathbf{x}) &= 1 \\ f_{2^{m-1}}(\mathbf{x}) &= 1 \text{ if } \mathbf{x} = 000\dots0001 \\ &= -1 \text{ if } \mathbf{x} = 1000\dots000 \\ &= 0 \text{ otherwise} \\ f_S(\mathbf{x}) &= 1 \text{ if } \mathbf{x} = 0S \text{ or } \mathbf{x} = 1S \\ &= -1 \text{ if } \mathbf{x} = S0 \text{ or } \mathbf{x} = S1 \\ &= 0 \text{ otherwise} \end{aligned}$$

Theorem 2: For a 1D, 2-state, neighborhood 3 cellular automaton, a basis set for the set of all conserved linear functions over m bits can consist solely of energy functions satisfying the constraint $f(0S)=0$ for all $m-1$ element strings S . In other words, the energy of an m -cell region can be defined to be zero for any region whose leftmost cell is zero.

Once a non-trivial conservation function of order m has been found, it can be extended in a very simple way to produce a function of order $m+1$ by ignoring the newly added cell on either side. The conservation functions reported in [1], however, do not distinguish between new functions derived in this way, since a conservation function of order m is by definition a conservation function of order $n \geq m$ for all n . We believe, however, that it is useful to distinguish between conservation functions derived from existing ones by ignoring newly added cells and conservation functions that are completely different.

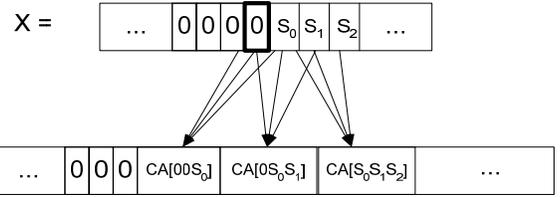
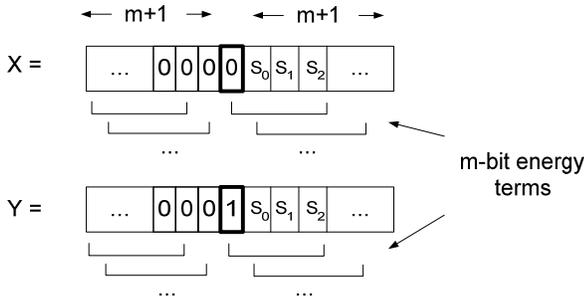
We refer to a non-trivial conservation function for a given CA that cannot be derived from non-trivial conservation of lower order as a *core* function. The core function of lowest order for a given CA is the *primary core* function for that CA. We call functions derived from a core by ignoring adjacent cells *simple extensions*. Simple extensions can be eliminated from the solution space through the addition of the constraint equations given above.

Eliminating simple extensions allows for the possibility of a CA having a completely different conservation function that would otherwise not be detected. Notice that through the judicious choice of basis vectors, we are able to reduce the size of the state space matrix by a factor of 2.

IV. AN ALGORITHM FOR CALCULATING CONSERVATION FUNCTIONS OF ORDER M FOR A GIVEN CA

By examining in detail how each of the energy terms are formed, it is possible to generate the state space matrix without doing explicit lookups and generation of all states and next state vectors. We do this by breaking the state space matrix up into three distinct matrices of different sizes, combining them only at the end when the null space must be calculated.

To see how this is done, consider first the terms in $E(X) - E(Y)$, the energy terms in the X and Y initial states, and what happens to them as the order m increases. We have for any m :



where $S_0, S_1 \dots$ are state counting bits (we have the leftmost bit change the fastest, for reasons that will become clear shortly). The energy expressions $E(X)$ and $E(Y)$ are just the sums of the m -bit energy terms for the X and Y states respectively. It is clear by inspection that all terms in $E(X) - E(Y)$ cancel except for those involving the center bit. Because we may also assume that $f(0, \dots) = 0$ for our energy function f , the matrix for $E(X) - E(Y)$ becomes

$$-f(1 \ S_0 \ S_1 \ \dots \ S_{m-2})$$

where the S bits are as previously described. In other words, each row has a -1 in the column indicated by bits $1S_0S_1 \dots S_{m-2}$. Making the leftmost 1 implicit in the column numbering, and reversing the column subscripts, we have

$$\text{For } m=1, E(X) - E(Y) = -1$$

$$\text{For } m=2, E(X) - E(Y) =$$

S_0	<u>10</u>	<u>11</u>
0	-1	0
1	0	-1

$$\text{For } m=3, E(X) - E(Y) =$$

S_0S_1	<u>100</u>	<u>110</u>	<u>101</u>	<u>111</u>
00	-1	0	0	0
10	0	-1	0	0
01	0	0	-1	0
11	0	0	0	-1

and in general $E(X) - E(Y)$ is given by a negative identity matrix of 2^{m-1} rows and 2^{m-1} columns.

Now consider the energy terms of the successor states, $E(X')$ and $E(Y')$. It is useful to first consider just those CAs that map the neighborhood region 000 to the 0 state. We shall refer to these CAs as zero-preserving automata.

For zero-preserving automata, we have the following relationship between X and its successor state X' :

since zero-preserving automata map 000 to 0 . (Here $CA[xyz]$ denotes the appropriate bit of the number of the CA written in binary). A similar analysis holds for states Y and Y' .

For $m = 1$, we can compute the energy difference of the successor states $E(X') - E(Y')$ as

$$\{E(X') - E(Y')\}_1 = CA[00S_0] + CA[0S_0S_1] - CA[001] - CA[01S_0] - CA[1S_0S_1]$$

since all other terms cancel, and we assume the energy function f maps all terms with an MSB of 0 to 0 . Let us rewrite the conservation law as

$$E(X) - E(Y) + E(Y') - E(X') = 0$$

Plugging our formula for $m=1$ above gives

$$-I + CA[001] + CA[01S_0] + CA[1S_0S_1] - CA[00S_0] - CA[0S_0S_1] = 0$$

where I is the identity matrix. Ignoring signs, we will refer to these terms as $E_x - E_y$, D_1 , D_2 , D_3 , R_1 and R_2 for reasons that will become apparent.

We reorder this expression to group terms with identical numbers of state bits together, giving

$$-I + CA[001] + CA[01S_0] - CA[00S_0] + CA[1S_0S_1] - CA[0S_0S_1] = 0$$

or equivalently

$$E_x - E_y + D_1 + D_2 - R_1 + D_3 - R_2 = 0$$

These six terms, with some bookkeeping, can be represented with three matrices of different sizes that grow in similar ways as m increases. Their size differences do not matter until the calculation of a null space is required, at which time they can all be made of uniform size and summed.

To see how this process works, first note that as m increases the energy terms grow to the right. Using the convention of having the leftmost state bit increase first, if we read the CA bit index from right to left, then each increment of m corresponds to simply copying the existing matrix both above itself and horizontally, and then shifting the individual rows of the new matrix by exactly half the columns if a particular bit of the CA is 1 .

For example, consider the D1 term for CA = 50 = 00110010 in binary. Recall that we read CA bit indices from right to left. For m=1, we have

$$D1_1 = CA[001] = 1$$

For m = 2, we have

$$D1_2 = CA[001] CA[01S_0] = CA[001] CA[010] = 1 \ 0$$

$$CA[001] CA[011] = 1 \ 0$$

$$= \begin{array}{cc} S_0 & \underline{10} & \underline{11} \\ 0 & 1 & 0 \\ 1 & 1 & 0 \end{array}$$

meaning for the state with $S_0 = 0$ this matrix has an energy term of $f(10)$, just as it does for the state with $S_0 = 1$ (recalling again that the MSB of 1 in the energy term is implicit in the column numbering). Note that we obtained $D1_2$ from $D1_1$ by copying the rows, doubling the columns, and moving the individual columns over to the right half of the matrix based on the values of the vector $CA[01S_0] = (CA[010] \ CA[011]) = \{0 \ 0\}$. We refer to this doubling and shifting operation as *expansion via the vector* $\{0 \ 0\}$, and refer to $\{0 \ 0\}$ as the *expansion vector*. In this case both values in the expansion vector are zero so no shifting of the columns occurs.

For m = 3, we have

$$D1_3 = CA[001] CA[01S_0] CA[1S_0S_1]$$

$$= \begin{array}{ccc} CA[001] CA[010] CA[100] & 101 \\ CA[001] CA[011] CA[110] & 100 \\ CA[001] CA[010] CA[101] & 101 \\ CA[001] CA[011] CA[111] & 100 \end{array}$$

$$= \begin{array}{cccc} S_0S_1 & \underline{100} & \underline{110} & \underline{101} & \underline{111} \\ 00 & 0 & 0 & 1 & 0 \\ 10 & 1 & 0 & 0 & 0 \\ 01 & 0 & 0 & 1 & 0 \\ 11 & 1 & 0 & 0 & 0 \end{array}$$

meaning for the state with $S_0S_1 = 00$ this matrix contributes an energy term of $f(101)$, and so forth. This time the expansion vector is $CA[1S_0S_1] = \{CA[100] \ CA[110] \ CA[101] \ CA[111]\} = \{1 \ 0 \ 1 \ 0\}$, corresponding to the third non-zero block in the previous figure.

After m = 3, the expansion vector always $CA[S_{m-4}S_{m-3}S_{m-2}]$. So $D1$ is now completely determined by:

$$D1_1 = CA[001]$$

$$D1_2 = \text{expansion_via}(D1_1, CA[01S_0])$$

$$D1_3 = \text{expansion_via}(D1_2, CA[1S_0S_1])$$

$$D1_m = \text{expansion_via}(D1_{m-1}, CA[S_{m-4}S_{m-3}S_{m-2}]) \text{ for } m > 3$$

It can be shown that after m=1, D3 and R2 have identical expansion vectors and may therefore be combined into a single matrix $C = D3-R2$. Similarly, after m=2, D2 and R1 may be combined into a single matrix $B = D2 - R1$. $Ex-Ey$ may be combined immediately with $D1$ into $A = Ex-Ey+D1$. Thus after some preliminary bookkeeping up through m=3, we need only maintain three energy term matrices A, B and C using only expansion operations with vectors derived from the CA under test. When the desired m has been reached, we replicate A 4 times and B twice, and then solve the $2^{m+1} \times 2^{m-1}$ system of equations $A+B+C = 0$.

We refer to the matrix $A+B+C$ as the *state space matrix N*. Finding a conservation function of order m for a given CA requires calculating the null space of N.

We see from the method of construction that A, B and C are sparse: Each one is the sum of matrices with only one non-zero term per row. This means that N can have at most six non-zero terms per row, out of rows that contain 2^{m-1} entries. In practice, N also contains a large number of redundant rows. Empirical analysis shows that removing all duplicate rows from N reduces its size by about a factor of two.

The authors have implemented the calculation of the null space of N for a given CA and order m in MATLAB. This program has been run at various DoD High Performance Research Center sites. We now turn to a description of the results we have obtained.

V. A TAXONOMY FOR 1-D CELLULAR AUTOMATA BASED ON CONSERVATION LAWS

We now present a complete taxonomy of 1-d CAs based on conservation functions of order $m \leq 16$. We use Wolfram's numbering scheme [4] to denote particular CAs, in which the successive state b of a cell appears as bit number i in an 8-bit binary number that designates the CA, where i is itself a 3-bit binary number determined by the left, center and right cell contents in a CA's next state rules.

When examining all s -state CAs, symmetry laws can be seen to divide the set into equivalence classes. Viewing the CA through a mirror produces a CA with identical properties, as does replacing state number j with state symbol $s-j$. For binary CAs, repeated application of these laws divides the 256 automata into 88 equivalence classes. For purposes of this discussion, if a single CA is mentioned, it is understood to refer to all CAs in its class. We will normally use the lowest numbered CA of a class to represent it.

Some CAs can be shown to have no non-trivial energy conservation functions. In particular, all CAs with next state functions of $x0x0x000$ have only trivial energy conservation functions, where x denotes 0 or 1. For a proof of this, see [3].

A. Classification By Lowest Order Conservation Function

We first present a table of CAs grouped by their lowest order conservation function. Table 1 lists all CAs with conservation functions up through order 16, grouped by the

order at which their first conservation function emerges. With one exception, this table is identical at lower orders to that in [1], but extends it to significantly higher values of m .

We note that [1] reports the existence of a CA with a conservation function at $m=7$. Wolfram [4] disagrees, and our results support his conclusions. We believe the entry in Table 1 in [1] for CA 19h and its equivalence class at $m=7$ should be 0.

There are 3 CA equivalence classes with conservation functions of order 1: The shifter (170 decimal/aa hex), the identity (204/cc), and 184/b8. 184/b8 is discrete asymmetric exclusion, and has been extensively studied [5].

At $m=2$ there are 11 equivalence classes with second order conservation properties, 8 at $m=3$, and so forth. The results for $m \geq 8$ are, to our knowledge, new ([6] reports higher order results for reversible CAs only), as are the basis functions identified in the next section.

B. Classification By Conservation Function Basis

We can gain more insight into the structure of CA conservation functions by examining the basis of the associated null space, noting both the number of dimensions and the basis vectors themselves. The tables in this section show the number of dimensions in the null space (which is equivalent to the dimensionality of the basis of the conservation function), and groups the CAs together for which the basis functions are identical. Such CAs should be regarded as equivalent in some natural sense. We can also examine what non-simple core functions emerge after the first one, as m increases.

Table 2 shows the non-zero half of the conservation function basis vectors for CAs with primary cores at $m=2$ through 5. (The three CAs with primary cores at $m=1$ conserve all functions on m bits and are therefore ignored). Entries that appear to the right of the first non-empty column represent functions that are also conserved and are not simple extensions of any cores to the left. For example, the CA of 15/f conserves the $m=2$ -bit function $f([10 \ 11]) = [1 \ 0]$, meaning that it conserves all "10" bit pairs. It also conserves the 3-bit function $f([100 \ 101 \ 110 \ 111]) = [-1 \ 1 \ -2 \ 0]$, and so forth. CAs with identical basis vectors for a given m are grouped by color. This table and those that follow have the property that all m -bit conservation functions for the indicated CA are some linear combination of trivial functions and the basis vectors shown.

The second-order conservation functions are exactly those identified in [2]. We believe all save 200/c8 belong together since they have identical basis vectors. Fuks notes that all CAs with second-order conservation functions save 200/c8 have identical invariants. Our results are equivalent.

Tables 3 and 4 show the dimensionality of null spaces and equivalence classes of CAs for all cores currently identified by our code.

It is surprising that, given how well-studied these simple automata are, new conservation functions remain to be discovered. A simple taxonomy for them, analogous to taxonomies for living systems, has yet to be found.

Classifying existing systems by the basis vectors of their conservations may point the way to a more exhaustive taxonomy. We would argue that CAs with identical basis vectors for a conservation function of a given order are related to one another in some important sense.

A total of 33 equivalence classes of the 1-D binary cellular automata remain to be classified. For these CAs, neither impossibility proofs nor existence proofs for conservation functions are currently available. Our classification scheme is thus incomplete; we have only determined by exhaustive search that no conservations of order 16 or below exist beyond those described.

The existence of an upper bound on the order of conservation functions for one-dimensional CAs of a given neighborhood size remains an open problem. The computational techniques described here may also be applied to higher dimensional binary automata, such as Conway's Game of Life, to determine the presence of conservation functions, but at present the computational requirements are still prohibitively large for even functions of very low order.

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TABLE 2
 CONSERVATION FUNCTIONS, BASIS VECTORS AND EQUIVALENCE CLASSES FOR CAS WITH FIRST CORES FROM M=2 THROUGH M=5

	m=2	m=3	m=4	m=5
12/c	[1 0]		[-1 -1 1 1 0 -2 0 0]	[-5 -5 3 3 1 1 1 1 0 -8 2 2 0 -8 0 0]
14/e	[1 0]			
15/f	[1 0]	[-1 1 -2 0]	[1 0 -1 0 -1 0 1 0] [2 2 -5 1 -5 1 0 0]	[-20 4 8 0 4 -4 8 0 4 -4 8 0 12 4 -24 0] [11 0 -7 4 -5 4 -7 0 -3 4 -9 0 -11 0 11 0] [-5 4 1 0 -1 4 -3 0 -3 0 -1 4 1 0 -9 0] [-1 4 -3 0 -1 0 -3 4 -3 4 -1 0 -3 0 -5 0]
34/22	[1 0]		[-1 -1 1 1 0 -2 0 0]	[-5 -5 3 3 1 1 1 1 0 -8 2 2 0 -8 0 0]
35/23	[1 0]			
42/2a	[1 0]	[0 0 1 0]	[-1 -1 1 1 1 -1 0 0]	[-3 -3 3 3 0 0 0 0 3 -3 0 0 0 0 0 0] [-2 -2 0 0 2 2 0 0 -1 -3 1 3 0 -6 0 0] [-2 -2 0 0 -1 -1 3 3 -1 -3 4 0 0 -6 0 0]
43/2b	[1 0]			
51/33	[1 0]	[-1 1 -2 0]	same as f	same as f
140/8c	[1 0]		[-1 -1 1 1 0 -2 0 0]	[-5 -5 3 3 1 1 1 1 0 -8 2 2 0 -8 0 0]
142/8e	[1 0]			
136/c8	[0 1]		[0 0 0 0 0 0 -1 1]	[0 0 0 0 0 0 0 -1 -1 -1 3 0 0 0 0] [0 0 0 0 0 0 0 0 0 0 0 0 0 0 -3 3]
2		[1 0 0 0]		
3		[1 0 0 0]		[-1 -1 1 1 -2 0 0 0 -2 -2 0 0 0 0 0 0]
4		[1 1 -1 0]		[1 1 1 1 -3 -3 1 1 -1 -1 3 -1 0 0 0 0]
10/a		[1 0 1 0]	[1 1 0 2 -1 -1 0 0]	
56/38		[1 1 1 0]		
76/4c		[1 1 1 0]	[1 1 1 1 -2 -2 3 0]	[3 3 3 3 0 0 -12 0 0 0 0 0 0 0 3 0] [0 0 0 0 0 0 0 -3 -3 3 3 3 -3 0 0] [0 0 0 0 3 3 -13 3 2 2 0 0 1 3 -3 0]
138/8a		[1 0 1 1]	[-2 -2 0 -3 1 1 1 1]	[0 0 0 0 0 0 1 -1 -1 -1 -1 -1 1 1 1]
172/ac		[1 0 0 0]		
1			[1 0 0 0 0 0 0 0]	
11/b			[1 1 0 1 0 0 0 0]	
27/1b			[1 1 0 1 0 0 0 0]	
29/1d			[1 1 1 1 1 0 0 0]	[-7 -7 -7 -7 3 3 3 3 3 13 -10 -10 -10 0 0 0]
38/26			[1 1 0 0 0 1 0 0]	
46/2e			[1 1 0 0 0 1 0 0]	
72/48			[0 0 0 0 1 1 -1 0]	
5				[1 0 1 0 0 0 0 0 1 0 1 0 0 0 0 0] [2 2 2 1 1 1 1 1 0 -1 0 -1 0 0 0 0] [0 0 0 0 1 0 0 0 1 1 0 0 0 0 0 0] [1 1 1 0 0 0 0 0 0 0 1 1 1 0 0 0] [1 1 1 1 -1 0 0 0 -1 -1 0 1 0 0 0 0] [1 1 1 1 1 0 0 1 1 1 0 0 0 1 0 0] [0 0 0 0 1 1 0 0 0 0 -1 0 0 0 0 0]
19/13				
24/18				
36/24				
108/6c				
132/84				

TABLE 3
EQUIVALENCE CLASSES AND DIMENSIONALITY OF CONSERVATION FUNCTIONS FOR CAS WITH FIRST CORES AT M=2 THROUGH M=6

CA	m=2	m=3	m=4	m=5	m=6	m=7	m=8	m=9	m=10	m=11
12/c	1		1	1	2	3	5	8	13	21
14/e	1									
15/f	1	1	2	4	8	16	32	64	128	256
34/22	1		1	1	2	3	5	8	13	21
35/23	1									
42/2a	1	1	1	3	5	9	17	31	57	105
43/2b	1									
51/33	1	1	2	4	8	16	32	64	128	256
140/8c	1		1	1	2	3	5	8	13	20
142/8e	1									
136/c8	1	1	1	2	4	7	12	21	37	65
2		1			1	1	1	2	3	4
3		1		1	2	3	5	9	16	28
4		1		1	1	2	3	5	8	13
10/a		1	1		1	3	4	5	9	16
56/38		1								
76/4c		1	1	3	5	8	14	25	45	82
138/8a		1	1	1	2	4	7	12	21	37
172/ac		1			1	1	1	2	3	3
1			1			1	2	3	4	6
11/b			1							
27/1b			1			2	3	5	9	16
29/1d			1	1		2	4	7	13	24
38/26			1			1	2	2	4	8
46/2e			1			1	1	1	2	3
72/48			1			1	1	1	2	3
5				2	2		2	7	8	21
19/13				1			1	2	3	4
24/18				1			1	1	1	2
36/24				1			1	1	1	2
108/6c				1	1	3	5	7	10	18
132/84				1		2		4	1	8
23/17					1		2		4	1
50/32					1		2		4	1
77/4d					2		4		8	2
178/b2					1		2		4	1
248/e8					2		4		8	2

TABLE 4
EQUIVALENCE CLASSES AND DIMENSIONALITY OF CONSERVATION FUNCTIONS FOR CAS WITH FIRST CORE AT M=7 THROUGH M=16

CA	m=2	m=3	m=4	m=5	m=6	m=7	m=8	m=9	m=10	m=11	m=12	m=13	m=14	m=15
No CAs with primary cores at m=7														
44/2c							1		1					
73/49							1							
7								1		2				
No CAs with primary cores at m=10														
No CAs with primary cores at m=11														
33/21											1			
164/a4												1		
94/5e													1	
104/68													2	
No CAs with primary cores at m=15														
No CAs with primary cores at m=16														