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Author	Message
PIsaacson	D Posted: Thu Apr 23, 2009 9:05 pm Post subject:
	Denis,
Joined: 02 Jul 2008 Posts: 183 Location: Campbell, CA	I became involved in studying ALSs and Allan's set/link-set logic to the point that I neglected to complete my promised statistics for the sudogen0 collection. Mea culpa. I need to re-trace my steps and re-read this thread to get back in sync
	Cheers, Paul
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Allan Barker	Dested: Fri Apr 24, 2009 6:47 am Post subject:
	StrmCkr wrote:
Joined: 21 Feb 2008	nice to see you back at it again denis
Location: Bangkok	Ditto. Hi Denis, welcome back.
	Allan
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denis_berthier	D Posted: Fri Apr 24, 2009 8:49 am Post subject:
	Hi Paul, Allan, StrmCkr
Joined: 19 Jun 2007 Posts: 631 Location: Paris, France	Glad to see you're still here.
	As for me, I've been (and still am) completely involved in other topics and out of the sudoku world for several months - except for the brief incursion in the above posts.
	StrmCkr, as I still have no time for Sudoku, I think you should open a thread to

discuss your new ideas instead of sending them to me alone ; you'd certainly get
much more interesting comments.

Paul, if you complete your computations for whips/braids with ALS inserts on sudogen0, I'm still interested. Given my above results, it doesn't matter much if you do them with braids or whips. I'm not sure : does your solver have "pure" ALS-chains (with no z- or t- candidates) ? Does it have their supersymmetric counterpart (i.e. with AHS and super-hidden-ALS or "A-fish" - whatever you call them)?

Allan, as I'ven't been reading what has been published in the last few months, you may have already answered these old questions of mine : have you defined any measure of complexity for your patterns? If so, can your algorithm be tuned to provide resolution paths that grant the "simplest" solution according to this measure?

Indeed, I'm still interested in anything that deals with principled measures of complexity for well defined families of patterns and in any comparison between such measures - the topic of this thread. I'm aware of nothing of this sort for pure ALS-chains.

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ttt	Dested: Fri Apr 24, 2009 3:25 pm Post subject:
	Me too, nice to see you back after around half year 😌
Joined: 20 Oct 2006 Posts: 173 Location: vietnam	ttt
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denis_berthier	D Posted: Sun Apr 26, 2009 8:01 am Post subject:
	ttt wrote:
Joined: 19 Jun 2007 Posts: 631	Me too, nice to see you back after around half year Θ
Location: Paris, France	ttt
	HI, ttt , nice to see you.
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PIsaacson	D Posted: Mon Apr 27, 2009 3:07 am Post subject:
	Denis,
Joined: 02 Jul 2008 Posts: 183 Location: Campbell, CA	I don't have braids with ALS/AHS group extensions, so I can only execute using nrczt chains/whips.
	Your question on "pure" ALS chains raises a point that I've been considering lately

from a programming perspective. Currently, I have a separate engine for ALS chaining and a separate engine for nrczt chaining. They both use a common ALS generator that builds the map of all possible ALS/AHS pairs within RC/RN/CN spaces, so: Yes, both engines have all the supersymmetrics available for use.

However, they build chains with ALSs using very different algorithms. In the nrczt engine, preference is (usually) given to linking 3d nrc candidates and attempting z/t promotion when possible. ALS/AHS nodes are (usually) only considered after exhausting all other possible nrc children of a given parent node. The ALS engine always attempts to chain ALS/AHS nodes directly via RCC peer linkage first, and then attempts to chain via a limited number of techniques such as using bivalue/local cells, ERs, GSLs, AURs and X-cycles to "bridge" RCCs from a parent ALS to a potential child ALS. At some point, I would like to have these two techniques "merge" into a super-engine.

I qualified the nrczt engine with the phrase "usually" because for these tests, I opted to have the nrczt engine select an ALS whenever possible instead of as a last resort. It made the results for short chains entirely different and came close to emulating the ALS engine restricted to linking no more than 2 or 3 ALSs.

So with that out of the way, here are the results from running nrczt-whips with ALS/AHS group extensions:

pNRCZT1_0: 4247 pNRCZT1: 1135 pNRCZT2: 3899 pNRCZT3: 525 pNRCZT4: 109 pNRCZT5: 45 pNRCZT6: 25 pNRCZT6: 25 pNRCZT7: 6 pNRCZT7: 6 pNRCZT8: 5 pNRCZT9: 3 pNRCZT10: 1

This demonstrates that ALSs lead to somewhat shorter solution paths. I believe this is simply because an ALS typically includes more weak link paths than a standard nrc candidate. Also, given the resolution power of braids and with the advent of whips, I am uncertain that ALS groups offer additional solutions.

Cheers, Paul

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denis_berthier

Dested: Mon Apr 27, 2009 7:41 am Post subject:

(quote K edit)

PIsaacson wrote:

Joined: 19 Jun 2007
Posts: 631
Location: Paris, FranceThis demonstrates that ALSs lead to somewhat shorter solution paths. I
believe this is simply because an ALS typically includes more weak link
paths than a standard nrc candidate.

	Maybe, the length of a chain should be increased by 2/3/4 instead of 1 when a Pair/Triplet/Quad is inserted? That'd be consistent with the fact that Pairs are chains of length 2 and (most) Triplets/Quads are chains of length 3/4.
	Anyway, I think it's worth giving a full example of a "zt promoted" naked, hidden or super-hidden subset within a whip.
	PIsaacson wrote:
	Also, given the resolution power of braids and with the advent of whips, I am uncertain that ALS groups offer additional solutions.
	It is interesting that nrczt-whips and supersymmetric ALS-chains have separately almost the same solving power (see my work with Mike, here http://www.sudoku.com/boards/viewtopic.php? t=5591&postdays=0&postorder=asc&start=90 and next page) but their combination through the zt-ing principle (z/t promotion in your vocabulary) is much stronger, as shown by the tables at the end of this page : http://www.sudoku.com/boards/viewtopic.php? t=6390&postdays=0&postorder=asc&start=30. And your super-engine would be
	able to solve very hard puzzles.
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denis_berthier	D Posted: Sat Jun 20, 2009 5:40 am Post subject:
Joined: 19 Jun 2007	NEW STATISTICAL RESULTS
Location: Paris, France	sudogen0_1M is a random sample of 1,000,000 different independent minimal puzzles generated with the suexg program (with seed 0). More on it in the next post.
	Using this extended sudogen0_1M collection, I computed the following variables for all the puzzles in it: the NRCZT-ratings, the SER ratings, the number of clues, the number of partial nrczt-chains.
	1) Here are first the correlation coefficients:
	NRCZT vs SER : 0.895 This confirms what I had already shown with the first 10,000 puzzles in the collection: although the two ratings are based on very different sets of rules, they are very strongly correlated.
	NRCZT vs log(#chains) : 0.946 This confirms that the length of chains is statistically a good indicator for the complexity of a puzzle: the more useless partial chains there are, the harder it is to find the useful ones. The number of useless partial chains increases exponentially with the length of the longer chain necessary to solve the puzzle (in the range of validity of these results : 1 to 10).

NRCZT vs #clues : 0.116 SER vs #clues : 0.121

This gives a precise meaning to what was already intuitively known concerning the number of clues: it has no impact on the complexity of minimal puzzles.

2) Here is now the number of puzzles per number of clues: (all the puzzles in the collection have between 20 and 30 clues).

Code:			
nb-clues	nb-puzzles		
20	44		
21	2,428		
22	34,548		
23	172,512		
24	342,335		
25	297,838		
26	122,116		
27	25,315		
28	2,686		
29	168		
30	10		

This shows that puzzles with less than 20 or more than 30 clues are extremely rare - in frequency. (Notice however that, in numbers, as there are billions of billions of minimal puzzles, "extremely rare" may still mean thousands or millions - we already know more than 64,000 17-clue puzzles). Most of the puzzles have between 23 and 26 clues.

3) The SER in the collection varies upto 9.2

4) Here is finally the number of puzzles per NRCZT-level (already posted in the "supersymmetric chains" thread)

Code:		
Level	Number	Total
1_0	417,624	417,624
1	120,618	538,242
2	138,371	676,613
3	168,355	844,968
4	123,153	968,121
5	24,187	992,308
6	5,511	997,819
7	1,514	999,333
8	473	999,806
9	130	999,936
10	38	999,974
11	15	999,989
12	9	999,998
13	2	1,000,000
14	0	

file:///Users/berthier/Desktop/DB-SPF-pdf/TODO/Rating%20Rules%2...g%20rules%20:%20Puzzles.%20Ordering%20the%20rules-10.webarchive Page 5 sur 11

	 This confirms that: more than 99% of the minimal puzzles can be solved with whips of length 5 or less; more than 99.9% of the minimal puzzles can be solved with whips of length 7 or less; "almost all" the puzzles can be solved with nrczt-whips of length 13 or less. Of course, such statistical results can't say anything about extreme cases wih very low probability (e.g. cases with very special symmetries, such as the EasterMonster.
	family).
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denis_berthier	D Posted: Sat Jun 20, 2009 7:08 am Post subject:
Joined: 19 Jun 2007	THE SUDOGEN0_1M COLLECTION
Location: Paris, France	sudogen0_1M is a random sample of 1,000,000 different independent minimal puzzles generated with the suexg program (with seed 0).
	suexg is a free program developed by Guenter Stertinbrink aka dukuso at Magictour (thanks Coloin for the exact name). It produces about 20 minimal puzzles per minute even on an old cheap micro-computer (1 GHz). There are several versions of it. For definiteness, a copy of the exact version I use is available on my web pages: http://www.carva.org/denis.berthier/HLS/Results- Classif/sudoku_gen.c
	Ever since I started being interested in Sudoku, I've used the "small" sudogen0 collection with 10,000 puzzles, but I've now extended it to 1,000,000. The reason is that hard puzzles are very rare (in frequency) and the higher we want to go into the complexity hierarchy (as reflected, say, by the NRCZT-level) the more puzzles we need to analyse.
	Now, the question that naturally arises is: how random is the "large" sudogen0_1M collection? This is a major question, because randomness is what allows to extend the results in the above posts from the sudogen0_1M collection to the full set of minimal puzzles.
	"Real randomness" can rarely be fully proven, but good indicators of randomness can be checked.
	The first thing that can be checked is that all the puzzles are different. Another thing one might want to check is that no two of them are essentially equivalent. I'ven't done this, first because it would have been one more long computation and secondly because it isn't really relevant here: we are dealing with minimal puzzles and not with equivalence classes of such puzzles. Anyway, in terms of proportions, even if a few puzzles were essentially equivalent, this wouldn't change the results. If anyone has a fast checker of essential-equivalence,

	he can do it. Also, as Coloin pointed out in a PM after this post, the probability of two puzzles being essentially equivalent in a 1,000,000 random sequence is very low).
	The second thing one is expecting from a random collection is the independence of the puzzles in it. As usual, independence of a sequence of non-numerical variables (the puzzles) is tested indirectly through the (normalised) auto-correlation function of one (or more) numerical function(s) of this variable. The numerical functions of the puzzles I'll consider here are: - the number of clues
	- the NRCZT rating - the SER (Sudoku Explainer Rating)
	What is the (normalised) auto-correlation function? Its k-th value (which is a number between -1 and +1) measures the degree of correlation between any element in the sequence and the element k places after it. (Look any textbook on statistics of sequences for a precise definition). How did I compute it? I used Matlab, with the exact function: xcorr(x, maxlag,'unbiased'); here "maxlag" is a parameter allowing to specify that we don't take into account values of k (beyond maxlag) that would rely on too few puzzles; I choose maxlag = 990,000.
	Now for the results: the maximum absolute values for the correlation function were: - for the number of clues: 0,041 - for the SER: 0,037 - for the NRCZT: 0,039 All these values are very small. All these sequences of variables are white noise.
	Conclusion: wrt to all the above criteria, the sudogen0_1M sequence has null auto-correlation.
	Soon, I'll put the whole series of results (lists of puzzles, SER, NRCZT, #clues) on my web pages, so that anyone can check these results and add his own.
	Edited 06/23/2009 to take into account two remarks from Coloin .
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denis_berthier	Denosted: Tue Jun 23, 2009 6:38 am Post subject:
Joined: 19 Jun 2007 Posts: 631 Location: Paris, France	The average number of clues in a minimal puzzle One result I had forgotten to give is the average number of clues in a minimal puzzle (it can easily be computed from the distribution of clues though). As Coloin evoked this number in a PM, here is the result for the sudogen0_1M collection: mean number = 24.38 standard deviation = 1.120
	For all the results reported in the above posts, what is important is that the

	sudogen0_1M collection is unbiased. This is enough to extend these results to the full set of minimal puzzles.
	But another interesting point is that suexg itself is unbiased, whatever the seed we use. This can be inferred from the algorithm or this can be checked experimentally on several seeds. I therefore tried another seed, 173, on a smaller sequence of 100,000 puzzles, and I checked only the number of clues. max number = 29 min number = 20 mean number = 24.387 standard deviation = 1.125 This is fully compatible with the results for the sudogen0. 1M collection
	Anyone can do these elementary computations starting with differents seeds. In order to avoid random fluctuations of the mean, a minimum of 100,000 puzzles seems reasonable.
	Last edited by denis_berthier on Fri Jul 03, 2009 5:28 am; edited 1 time in total
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denis_berthier	Dested: Wed Jun 24, 2009 8:08 am Post subject:
Joined: 19 Jun 2007 Posts: 631 Location: Paris, France	As I anounced in a previous post, I've now put all the above rating results online on the following Web page: http://www.carva.org/denis.berthier/HLS/Ratings.html From the middle of this page, you can download the sudogen0_1M collection, the ratings (SER, NRCZT) used above, the numbers of clues and the solution grids. These are all simple .txt files, with 1 puzzle or value per line.
	Also, I have updated some of my Web pages related to Sudoku http://www.carva.org/denis.berthier/HLS-Supplements. These are mainly a synthetic compilation of my posts in this forum.
	Last edited by denis_berthier on Sat Jul 04, 2009 5:16 pm; edited 1 time in total
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m_b_metcalf	D Posted: Wed Jun 24, 2009 11:29 am Post subject:
	denis_berthier wrote:
Joined: 15 May 2006 Posts: 2150 Location: Berlin	suexg is a free program developed by Guenter Stertinbrink aka dukuso at Magictour (thanks Coloin for the exact name). It produces about 20 minimal puzzles per minute even on an old cheap micro-computer (1 GHz).
	I was sufficiently intrigued by this result that I fired up my own program and let it run for one hour real time (@ 2GHz). The summary is:
	Code:

	Number: Average #	of clues:	51492 (~860/min) 24.36532	
	20	2	21.00002	
	21	142		
	22	1853		
	24 1	7656		
	25 1	5306		
	26	6127 1257		
	28	149		
	29	4		
	Looks comparable who'd like it. Whe another question.	. The zipped f ther I'd want	file is 1.3MB and I'd be happy to send to go for another 19 hours to reach	d it anyone the million is
	Regalus,			
	Mike Metcalf			
	P.S. As an exercis zipped file is 24.8	e, I once gen MB.	erated 1 million solution grids (in 10	0secs). The
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denis_berthier	Dested: Wed Jur	1 24, 2009 12	:56 pm Post subject: 🔍	juote) 🖧 edit
	m_b_metca	If wrote:		
Posts: 631 Location: Paris, France	I was sufficie	ently intrigued	by this result that	
	Which result exact	tly did intrigue	e you? Do you know why you were ir	trigued by it?
	BTW, which seed	did you use in	suexg?	
	Yes, your findings random fluctuation	are compara ns due to sam	ble. Minor discrepancies can be explanple size.	ained by normal
	Average # of clue	s: 24.36532 v	rs 24,38	
	Code:			
	#Clues # results	Puzzles	#Puzzles*1,000,000/51492	vs my
	20	2	39 vs 44	
	21	142 1853	2,470 vs 2,428 35.986 vs 34.548	
	22	1000		
	23	8996	174,797 vs 172,512	
	23 24 1	8996 7656	174,797 vs 172,512 342,888 vs 342,335	

	26 6127 118,989 vs 122,116 27 1257 24,412 vs 25,315 28 149 2,894 vs 2,686 29 4 78 vs 168 30 0 0 vs 10
Back to top	Regards
m b metcalf	D Posted: Wed Jun 24, 2009 1:17 pm Post subject:
	donie horthior wroto:
Joined: 15 May 2006 Posts: 2150 Location: Berlin	Which result exactly did intrigue you? Do you know why you were intrigued by it? BTW, which seed did you use in suexg?
	I was intrigued to know whether my <i>own</i> program (<i>not</i> suexg) would yield the same results, and by what factor my program might be faster (~20, as it turned out). Regards, Mike Metcalf P.S. I doubled the sample, yielding:
	Code: Number: 110840 Average: 24.36463 20 10 21 324 22 4081 23 19378 24 37965 25 32685 26 13271 27 2778 28 333 29 15 30 0
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denis_berthier	DPosted: Wed Jun 24, 2009 1:40 pm Post subject: (2) quote
Joined: 19 Jun 2007 Posts: 631 Location: Paris, France	<pre>m_b_metcalf wrote: I was intrigued to know whether my own program (not suexg) would yield the same results, and by what factor my program might be faster (~20, as it turned out).</pre>

	As your reslts are based on a different generator, they are still more interesting, especially if the generation algorithm is significantly different from suexg.
	Is your generator available? Or is there any place where the basic principles it uses are defined? Or can you state them in a few words?
	In any case, I'm interested by your file of 110840 puzzles. It'd be a good way to check my results about the nrczt and SER ratings and to confirm that they don't depend on the generator.
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