

A SAT Attack on the Erdős Discrepancy Conjecture

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The Erdős discrepancy conjecture is interesting (even for $C = 2$).
EDP2 can be settled by reduction to SAT.

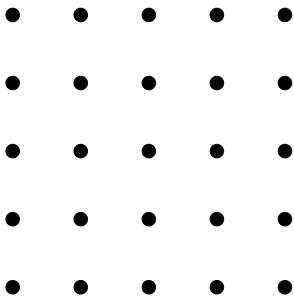
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EDP2 can be settled by reduction to SAT.

- Discrepancy Theory
- Erdős Discrepancy Conjecture
- SAT attack on the EDP
- Results and Perspectives

Discrepancy theory is a branch of mathematics dealing with
inevitable irregularities of distributions

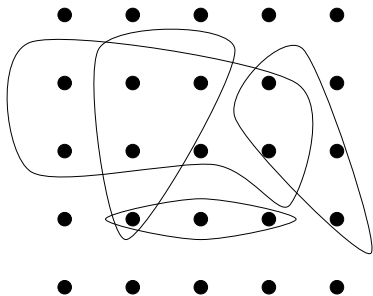
Combinatorial discrepancy



- A set U
- Set of subsets S

- A hypergraph $\mathcal{H} = (U, S)$
- Consider a colouring $c : U \rightarrow \{+1, -1\}$ of the elements of U in *blue* (+1) and *red* (-1) colours;
- **Question:** Is there a colouring such that in every element of S colours are distributed uniformly or a **discrepancy** of colours is always inevitable?

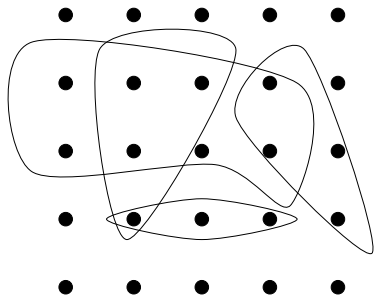
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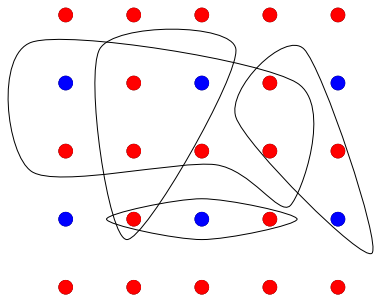
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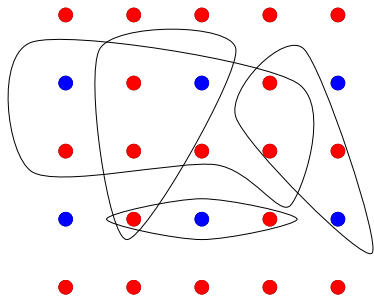
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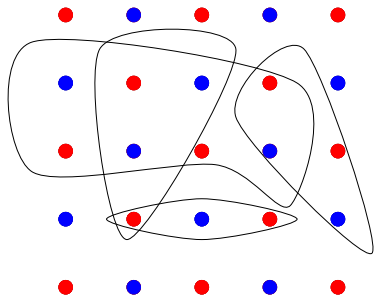
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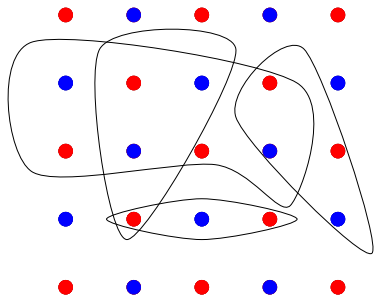
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$$d(\mathcal{H}) = 1$$

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Combinatorial number theory

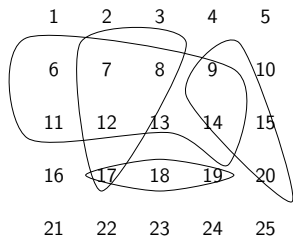
1	2	3	4	5
6	7	8	9	10
11	12	13	14	15
16	17	18	19	20
21	22	23	24	25

- $U \subseteq \mathbb{N}$
- S — 'arithmetically interesting' subsets of U

Theorem (Roth, 1964)

For $U_n = \{1, 2, \dots, n\}$ and $S_n = \{(a \cdot i + b)\}$
the discrepancy grows at least as $\frac{1}{20} n^{1/4}$

Combinatorial number theory



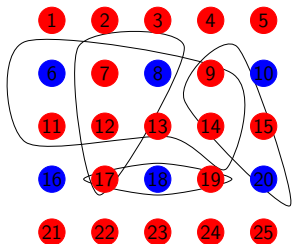
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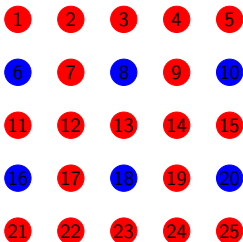


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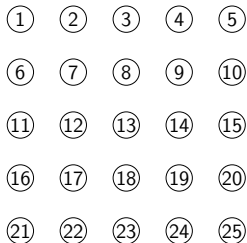


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Erdős Discrepancy Conjecture (EDP)



What about **homogeneous** arithmetic progressions?

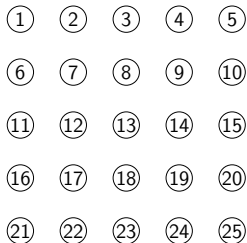
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Conjecture (Erdős, circa 1930)

For any $C > 0$ in any infinite ± 1 sequence (x_n) there exists a subsequence $x_d, x_{2d}, x_{3d}, \dots, x_{kd}$ such that $|\sum_{i=1}^k x_{i \cdot d}| > C$.

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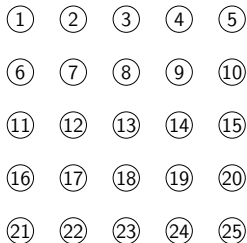
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Known results on the discrepancy of ± 1 sequences

- For random ± 1 sequences the discrepancy grows as $n^{1/2+o(1)}$ (folklore?);
- An explicit constructions of a sequence with slowly growing discrepancy $\log_3 n$ [Borwein, Choi, Coons, 2010];
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| | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|----|----|----|---------|
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|---|---|---|---|---|---|---|---|---|----|----|----|---------|
| + | - | - | + | - | - | - | - | - | - | - | - | $d = 2$ |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | |

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$d = 3$

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$d = 4$

Example: $C = 1$

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| | | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|----|----|----|--|---------|
| + | - | - | + | - | + | + | - | - | - | - | - | | $d = 1$ |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | | |

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| | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|----|----|----|---------|
| + | - | - | + | - | + | + | - | - | + | - | - | $d = 5$ |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | |

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| | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|----|----|----|
| + | - | - | + | - | + | + | - | - | + | - | - |
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$d = 6$

Example: $C = 1$

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| | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|----|----|----|---------|
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|---|---|---|---|---|---|---|---|---|----|----|----|---------|
| + | - | - | + | - | + | + | - | - | + | + | - | $d = 3$ |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | |

But then $x_3 + x_6 + x_9 + x_{12} = -1 + 1 - 1 - 1 = -2$

- A 2009-2010 topic of Polymath — ‘massively collaborative maths’ project started and coordinated by T. Gowers
 - A computer attack
 - Discrepancy 2 sequences of length 1124 (backtracking search)

“... given how long a finite sequence can be, it seems unlikely that we could answer this question just by a clever search of all possibilities on a computer...”

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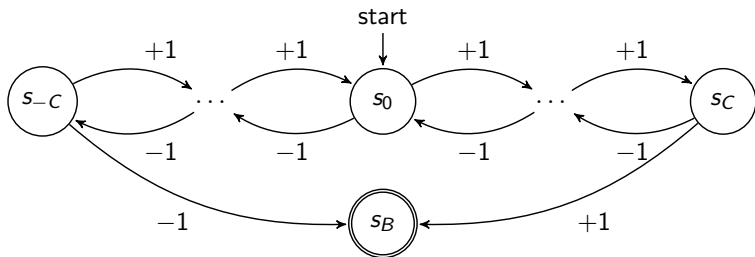
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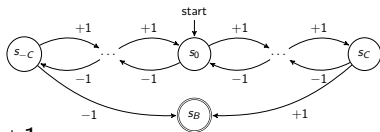
- There are ± 1 sequences of length 1160 and discrepancy 2,
- There are no ± 1 sequences of length 1161 (or more) and discrepancy 2.

Automata encoding of discrepancy conditions



- If for every $d : 1 \leq d \leq \lfloor \frac{n}{C+1} \rfloor$ the automaton \mathcal{A}_C does not accept the subsequence $x_d, x_{2d}, \dots, x_{kd}$, where $k = \lfloor \frac{n}{d} \rfloor$ then the discrepancy of the sequence \bar{x} does not exceed C

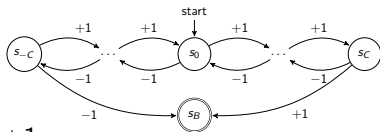
SAT representation



- p_i is true if \iff i -th letter is $+1$.
- $s_j^{(i,d)}$ is true \iff \mathcal{A}_C is in s_j having read first $(i - 1)$ letters.

$$\phi_{(n,C,d)} = s_0^{(1,d)} \bigwedge_{i=1}^{\lfloor \frac{n}{d} \rfloor} \left[\begin{aligned} & \bigwedge_{-C \leq j < C} \left(s_j^{(i,d)} \wedge p_{i \cdot d} \rightarrow s_{j+1}^{(i+1,d)} \right) \wedge \\ & \bigwedge_{-C < j \leq C} \left(s_j^{(i,d)} \wedge \neg p_{i \cdot d} \rightarrow s_{j-1}^{(i+1,d)} \right) \wedge \\ & \left(s_C^{(i,d)} \wedge p_{i \cdot d} \rightarrow B \right) \wedge \\ & \left(s_{-C}^{(i,d)} \wedge \neg p_{i \cdot d} \rightarrow B \right) \end{aligned} \right]$$

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Let

$$\phi_{(n,C)} = \neg B \wedge \bigwedge_{d=1}^{\lfloor \frac{n}{C+1} \rfloor} \phi_{(n,C,d)} \wedge \text{frame}_{(n,C)},$$

where $\text{frame}_{(n,C)}$ is a Boolean formula encoding that the automaton state is correctly defined.

Proposition

The formula $\phi_{(n,C)}$ is satisfiable if, and only if, there exists a ± 1 sequence $\bar{x} = x_1, \dots, x_n$ of length n of discrepancy C . Moreover, if $\phi_{(n,C)}$ is satisfiable, the sequence $\bar{x} = x_1, \dots, x_n$ of discrepancy C is uniquely identified by the assignment of truth values to propositions p_1, \dots, p_n .

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In fact we have used more “economical” encoding, where a state s_i of automaton is encoded not by a separate propositional variable p_i , but by the propositional variables encoding i in binary

In our experiments we used

- the Lingeling SAT solver, the winner of the *SAT-UNSAT* category of the SAT'13 competition, and
- the Glucose solver version, the winner of the *certified UNSAT* category of the SAT'13 competition.

All experiments were conducted on PCs equipped with an Intel Core i5-2500K CPU running at 3.30GHz and 16GB of RAM.

Experiments and Results

- By iteratively increasing the length of the sequence, we establish precisely that the maximal length of a ± 1 sequence of discrepancy 2 is 1160.
- On our system it took Plingeling, the parallel version of the Lingeling solver, about 800 seconds to generate a sequence of discrepancy 2 and length 1160.
- On the other hand, when we increased the length of the sequence to 1161, Plingeling reported unsatisfiability.
- We also used Glucose: It took the solver about 21 500 seconds to compute a Delete Reverse Unit Propagation (DRUP) certificate of unsatisfiability ($\sim 13\text{Gb}$).
- The certificate has been independently verified by the drup-trim tool

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- We also used Glucose: It took the solver about 21 500 seconds to compute a Delete Reverse Unit Propagation (DRUP) certificate of unsatisfiability ($\sim 13\text{Gb}$).
- The certificate has been independently verified by the drup-trim tool

Experiments and Results

- By iteratively increasing the length of the sequence, we establish precisely that the maximal length of a ± 1 sequence of discrepancy 2 is 1160.
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 - 13GB!
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We have applied the same methodology to the case $C=3$.

Proposition

There exists a sequence of length $\approx 14,000$ of discrepancy 3.

Life after SAT:

- Better SAT encodings
- Tuned search strategy
- 850Mb RUP certificate for EDP2
- Multiplicative and completely multiplicative EDP2 sequences (for EDP2)
- Longest completely multiplicative EDP2 sequence contains 127 elements
- Longest multiplicative EDP3 sequence also contains 127 elements
- Longest multiplicative EDP3 sequence is 334,000

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- Multiplicative and completely multiplicative sequences for EDP3

$$x_{m \cdot n} = x_m \cdot x_n$$

- Longest completely multiplicative EDP3 sequences contains 127 645 elements
 - Previously reported by La Tour, Lomer and Suman, <https://arxiv.org/abs/1405.3097>
- Longest multiplicative EDP3 sequences also contains 127 645 elements!
 - Valid only for $C=1$ and $C=2$
- New lower bound for EDP3 of **130 000**

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