A new algebraic approach to

the graph isomorphism and clique problems

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As it follows from Gödel's incompleteness theorems, any consistent formal system of axioms and rules of inference should imply a true unprovable statement. Actually this fundamental principle can be efficiently applicable in Computational Mathematics and Complexity Theory concerning the computational complexity of problems from the class NP, particularly and especially the NP-complete ones. While there is a wide set of algorithms for these problems that we call heuristic, the correctness or/and complexity of each concrete algorithm (or the probability of its correct and polynomial-time work) on a class of instances is often too difficult to determine, although we may also assume the existence of a variety of algorithms for NPcomplete problems that are both correct and polynomial-time on all the instances from a given class (where the given problem remains NP-complete), but whose correctness or/and polynomial-time complexity on the class is impossible to prove as an example for Gödel's theorems. However, supposedly such algorithms should possess a certain complicatedness of processing the input data and treat it in a certain algebraically "entangled" manner. The same algorithmic analysis in fact concerns all the other significant problems and subclasses of NP, such as the graph isomorphism problem and its associated complexity class GI.

The following short article offers a couple of algebraically entangled polynomial-time algorithms for the graph isomorphism and clique problems whose correctness is yet to be determined either empirically or through attempting to find proofs.

An heuristic polynomial-time algorithm

for the graph isomorphism problem.

Denotation:

For a real-valued matrix B, by Sp(B) let's denote the set of values its entries are equal to, and by MSp(B) the multi-set of those values (including the multiplicity of each value). We'll call those set and multi-set the *entry spectrum* and the *entry multi-spectrum* of B correspondingly.

Given two simple undirected graphs G_1 and G_2 with n vertices whose adjacency matrices are A_1 and A_2 correspondingly, we're going to build two sequences of real-valued matrices $A_1^{(i)}$ and $A_2^{(i)}$ by the following recursive scheme:

$$A_1^{(0)} = A_1, \qquad A_2^{(0)} = A_2$$

Beginning with i = 1, at the step i of our recursive process, first of all we determine whether $MSp(A_1^{(i)}) = MSp(A_2^{(i)})$.

If it's not so then G_1 and G_2 are definitely non-isomorphic. If, otherwise, the equality holds then we create a random $|Sp(A_1^{(i)})|$ -vector y and in both matrices $A_1^{(i)}$ and $A_2^{(i)}$ we replace each entry whose value is the j-th element of $Sp(A_1^{(i)})$ by y_j . After that, we choose a random polynomial p(t) of degree n-1 and calculate $A_1^{(i+1)} = p(A_1^{(i)})$ and $A_2^{(i+1)} = p(A_2^{(i)})$.

We stop the whole process when either the entry multi-spectrums of the two current matrices are different or the cardinality of their common entry spectrum doesn't increase any more from one step to another, i.e. $|Sp(A_1^{(k)})| = |Sp(A_1^{(k-1)})|$ (accordingly, the overall number k of performed steps can't exceed n²).

We hence declare the initial graphs' non-isomorphism in the former case, and their isomorphism in the latter one.

When the process is stopped, with the overall number of steps equal to k and the final entry spectrum $\{\alpha_1, ..., \alpha_m\}$, there should exist symmetric 0,1-matrices $H_1^{(1)}, ..., H_1^{(m)}, H_2^{(1)}, ..., H_2^{(m)}$ such that

$$A_1^{(k)} = \sum_{u=1}^m \alpha_u H_1^{(u)}$$
, $A_2^{(k)} = \sum_{u=1}^m \alpha_u H_2^{(u)}$,

We can consider, for u = 1,...,m, $H_1^{(u)}$ and $H_2^{(u)}$ as the adjacency matrices of some graphs $G_1^{(u)}$ and $G_2^{(u)}$. In case if the initial graphs G_1 and G_2 are really isomorphic, those two graphs should be isomorphic as well. Moreover, due to the entry spectrum's non-growing at the end of the above-described process, for any pair v, w $\in \{1, ..., m\}$, $v \neq w$, $G_1^{(v)}$ and $G_1^{(w)}$ should possess no common edges and $G_2^{(v)}$ and $G_2^{(w)}$ should too; also there should exist a sequence of coefficients $d_1^{(v,w)}$, ..., $d_m^{(v,w)}$ such that $H_1^{(v)}H_1^{(w)} + H_1^{(w)}H_1^{(v)} = \sum_{u=1}^m d_u^{(v,w)}H_1^{(u)}$ and $H_2^{(v)}H_2^{(w)} +$ $H_2^{(w)}H_2^{(v)} = \sum_{u=1}^m d_u^{(v,w)}H_2^{(u)}$ (the latter condition also concerns the case v = w). Verifying these additional relations is the proposed algorithm's final action when declaring the isomorphism of G_1 and G_2 . Thus we get a pair of identical commutative algebras that are two linear spaces with the bases $H_1^{(1)}$, ..., $H_1^{(m)}$ and $H_2^{(1)}$, ..., $H_2^{(m)}$ whose elements are $X = \sum_{u=1}^m x_u H_1^{(u)}$ and X = $\sum_{u=1}^m x_u H_2^{(u)}$ correspondingly and whose algebra product is defined as X * Y = XY + YX.

However, if, after some global steps, in each of the above linear spaces we take "standard" matrix products of various space elements (related by having the same coefficients for each space) then we'll receive, upon the entry-spectrum's ceasing to grow with further performed steps, a pair of corresponding identical non-commutative algebras whose algebra product is just the standard matrix product (instead of the above-introduced symmetric one) and whose product closure condition is, accordingly, $H_1^{(v)}H_1^{(w)} = \sum_{u=1}^m d_u^{(v,w)}H_1^{(u)}$ and $H_2^{(v)}H_2^{(w)} = \sum_{u=1}^m d_u^{(v,w)}H_2^{(u)}$ for any $v, w \in \{1, ..., m\}$ (instead of the above symmetric one).

The above-formulated algorithm can be naturally adjusted for digraphs with absolutely the same computational circuit (while, though strange, generating, in the commutative algebra case, commutative algebras at the very end as well). Moreover, it can be generalized for an arbitrary field (instead of \mathbb{R}) with all the computations performed over that field. In case if the chosen field is of characteristic 2, we'll receive Lie algebras.

Another type of generalization this algorithm might be subjected to is as follows.

Definition:

Given an n×n- matrix B, let's define a product $B^{r_1}1_{n\times n}B^{r_2}1_{n\times n} \dots B^{r_{s-1}}1_{n\times n}B^{r_s}$ (where $1_{n\times n}$ is an n×n-matrix all whose entries equal unity and r_1, \dots, r_s are non-negative integers unexceeding n-1) as its *meta-power* of *meta-degree* (r_1, r_2, \dots, r_s) and a linear combination of a set of its meta-powers with coefficients taken from a chosen field as a *meta-polynomial* in B over the field.

If an n×n-matrix can be turned into another one via permuting its rows and columns by a permutation π (i.e. if we have a pair of isomorphic matrices) then an arbitrary meta-polynomial computed in both of them should give us a pair of isomorphic matrices as well (with the same transitional permutation π). Hence the idea of replacing, in the proposed algorithmic approach, random polynomials p(t) of degree n-1 by random meta-polynomials may look yet perspective, even though it's still difficult to figure out how the meta-polynomials' sets of utilized meta-degrees could be restricted in such a case. Nevertheless, the principle of stopping the recursive process upon getting either different entry multi-spectrums of the two current matrices or their common entry spectrum's cardinality ceasing to grow from one step to another remains intact (while the matrices $A_1^{(i)}$ and $A_2^{(i)}$ generically cease to be symmetric for i > 0 even in the case of undirected initial graphs). The final splittings of the two matrices $A_1^{(k)}$ and $A_2^{(k)}$ (where k is the overall number of performed steps) apparently will have nearly the same structure as in the case of random polynomials p(t), but with the only additional condition of $H_1^{(u)} 1_{n \times n}$ and $1_{n \times n} H_1^{(u)}$ belonging to the first final algebra and $H_2^{(u)} 1_{n \times n}$ and $1_{n \times n} H_2^{(u)}$ to the second one for u = 1,...,m.

Such a final algebra doesn't depend on the chosen random polynomials or, generally, metapolynomials within the algorithmic circuit (as well as the random substitution vectors we use for entry spectrums' replacements). A proof of this fact can be received via using, instead of random elements of the basic field, independent indeterminates as the entries of substitution vectors in each spectrum replacement and the coefficients of utilized polynomials at each global step and getting, accordingly, polynomials in those indeterminates as the entries of transformed matrices (in such a case the algorithm ceases to be polynomial-time, though). We'll call it a *splitting algebra* of a graph that is a system of its invariants, while an associative splitting algebra will be also called a *splitting ring*.

Hence we can reduce, via the proposed method, determining whether two graphs are isomorphic to determining whether a related (by having identical sequences of coefficients at

 ${\rm H}_1^{(u)}$ and ${\rm H}_2^{(u)})$ random pair of elements of their splitting algebras represented by the corresponding matrices $\sum_{u=1}^m x_u H_1^{(u)}$ and $\sum_{u=1}^m x_u H_2^{(u)}$ is isomorphic as a pair of weighted graphs (or digraphs), with an application of all the well-known approaches for weighted graphs/digraphs including comparing their eigenvalues and eigenvectors where, nevertheless, the principle of entry spectrum random replacement can be applied again, hence providing new splitting options for possibly building some extensions of the splitting algebras. For this purpose, for an n×n-matrix $M = PJP^{-1}$ (where J is its Jordan form) and an |Sp(P)|-vector y, let's define its generalized eigenvector entry-spectrum replacement by y as the matrix $\widehat{P}J\widehat{P}^{-1}$ where \widehat{P} is the matrix received from P via replacing P's entry spectrum by y (when all the Jordan cells are of size 1×1, i.e. M has its eigendecomposition, we just can call it an eigenvector entryspectrum replacement – and it's the case for any symmetric M). Therefore the above-defined random polynomial/meta-polynomial and entry-spectrum replacement transformations of a matrix might be completed by a random generalized eigenvector entry-spectrum replacement modification performed after each global step (in this regard, actually, let's notice that a polynomial transformation is equivalent to a replacement of the matrix's eigenvalues, unlike a meta-polynomial one). We accordingly can define the final algebra appearing in such a process upon the matrix's entry spectrum's ceasing to grow as an extended splitting algebra of a graph/digraph (or a splitting algebra extended by generalized eigenvector entry-spectrum replacements) which may be conjectured to be a complete system of graph invariants.

It would be also worth noting that another types of splitting algebras' extensions might be based on other compatible (with any isomorphism) matrix decompositions such as the polar decomposition. In this regard, further we're going to formalize this principle, as well to give some set-theoretical basis to all the above-stated algorithmic schemes.

Denotation:

Given a permutation $\pi \in S_n$, by I_{π} we'll denote the matrix received from I_n via subjecting its rows to the permutation π .

Definition:

Given a ring K, a transformation ω : $K^{n \times n} \rightarrow K^{n \times n}$ (applicable for any natural n) will be called isomorphism-commutative if $\forall \pi \in S_n$, $\forall X \in K^{n \times n} \omega(I_{\pi}XI_{\pi}^T) = I_{\pi}\omega(X)I_{\pi}^T$.

Lemma 1:

The function composition of two isomorphism-commutative transformations is an isomorphism-commutable transformation, i.e. the set of isomorphism-commutative transformations is a semigroup under the function composition operation.

Lemma 2:

Given a square matrix A over a ring K, the set of its isomorphism-commutative transformations is a ring (that is an extension of K) under the matrix arithmetic operations.

Hence the set of isomorphism-commutative transformations has the two important properties that it's closed under the function composition and the matrix arithmetic operations. However, the same two properties are possessed by the set of isomorphism-commutative transformations computable in polynomial time. Let's denote the two sets by ICT and ICTP correspondingly.

Definition:

Given a ring K, a function f: $K^{n \times n} \times ... \times K^{n \times n} \rightarrow K^{n \times n}$ in m n×n-matrix variables $X_1, ..., X_m$ (applicable for any natural n) will be called isomorphism-commutative if $\forall \pi \in S_n$, $\forall X_1, ..., X_m \in K \quad f(I_{\pi}X_1I_{\pi}^T, ..., I_{\pi}X_mI_{\pi}^T) = I_{\pi}f(X_1, ..., X_m)I_{\pi}^T$.

We hence can consider an isomorphism-commutative transformation as an isomorphismcommutative function in one variable. Besides, the set of isomorphism-commutative functions is also closed under function composition and the matrix arithmetic operations.

Definition:

Given a set $f = \{f_0(X_1, ..., X_m), f_1(X_1, ..., X_m), ..., f_r(X_1, ..., X_m)\}$ of r+1 n×n-matrix functions in m n×n-matrices and an n×n-matrix A over a ring K such that the matrix equation system

$$\begin{split} &f_0(X_1,\ldots,X_m)=Y\\ &f_q(X_1,\ldots,X_m)=0_{n\times n} \ \text{ for } q=1,\ldots,r \end{split}$$

has a unique solution $\text{Decom}_1(Y, f), \dots, \text{Decom}_m(Y, f)$ for any $Y \in K^{n \times n}$. Then $\text{Decom}_1(A, f), \dots, \text{Decom}_m(A, f)$ will be called the f-decomposition of A over K.

Lemma 3:

Given a set $f = \{f_0(X_1, ..., X_m), f_1(X_1, ..., X_m), ..., f_r(X_1, ..., X_m)\}$ of isomorphism-commutative functions over a ring K, $Decom_1(A, f), ..., Decom_m(A, f)$ are isomorphism-commutative transformations of a square matrix A over K.

Conjecture 1:

For any ring K, any two square matrices A, B are isomorphic if and only if $MSp(\omega(A)) = MSp(\omega(B))$ for any isomorphism-commutative transformation.

Conjecture 2:

For any ring K, there exists a polynomial p(n) such that, for any two $n \times n$ -matrices A, B over a ring K, they are isomorphic if and only if $MSp(\omega(A)) = MSp(\omega(B))$ for any isomorphism-commutative transformation computable in p(n) arithmetic operations over K.

If the latter conjecture is true then the question whether two given matrices are isomorphic can be answered, by an algorithm from the complexity class RP, via subjecting both matrices to a

random isomorphism-commutative transformation computable in p(n) arithmetic operations over the basic ring (i.e. a random element of ICTP generating, accordingly, the algorithm's polynomial-time complexity) and determining whether the entry multi-spectrums of the two transformed matrices are equal. In such a case we'll also get a randomized complete system of graph invariants, although, as it was already mentioned earlier for splitting algebras received via the use of eigenvectors, there might exist a fixed set of ITCP's elements that is the complete invariant system.

A special case of the graph isomorphism problem that is a system of group equations.

Let's consider a special case of the isomorphism problem for two colored graphs (where all the vertices and edges are given colors) $G_1 = (V, E_1)$ and $G_2 = (V, E_2)$ such that

$$\begin{split} &V=\cup_{i=1}^m V^{[i]}, \ \forall v\in V^{[i]} \ \text{color}(v)=c_i, \\ &V^{[j]}\cap V^{[k]}=\emptyset \ \text{and} \ c_j\neq c_k \ \text{if} \ j\neq k \end{split}$$

In such a case we get the following system of equations:

$$\psi_{k}\psi_{j}G(V^{[j]} \cup V^{[k]}, E_{1}(V^{[j]}, V^{[k]})) = G(V^{[j]} \cup V^{[k]}, E_{2}(V^{[j]}, V^{[k]}))$$

where $G(V^{[j]} \cup V^{[k]}, E_1(V^{[j]}, V^{[k]}))$, $G(V^{[j]} \cup V^{[k]}, E_2(V^{[j]}, V^{[k]}))$ are the bipartite graphs generated by G_1, G_2 correspondingly on the set $V^{[j]} \cup V^{[k]}$ as the induced graphs where all the internal edges inside the sets $V^{[j]}$, $V^{[k]}$ were eliminated, and ψ_i is a permutation of $V^{[i]}$ (considered as a transformation of any graph whose vertex set contains $V^{[i]}$) belonging to the set of isomorphisms between $G(V^{[i]}, E_1(V^{[i]}))$ and $G(V^{[i]}, E_2(V^{[i]}))$ where $G(V^{[i]}, E_1(V^{[i]}))$, $G(V^{[i]}, E_2(V^{[i]}))$ are the graphs induced by $V^{[i]}$ in G_1, G_2 correspondingly. As we can represent any ψ_i as $\rho_i \phi_i$ where ρ_i is a fixed isomorphism between $G(V^{[i]}, E_1(V^{[i]}))$ and $G(V^{[i]}, E_2(V^{[i]}))$ and $\phi_i \in Aut(G(V^{[i]}, E_1(V^{[i]}))$, we'll hence receive the system of group equations for $\phi_1, ..., \phi_m$

$$\rho_{j}\phi_{j}\rho_{k}\phi_{k}G(V^{[j]} \cup V^{[k]}, E_{1}(V^{[j]}, V^{[k]})) = G(V^{[j]} \cup V^{[k]}, E_{2}(V^{[j]}, V^{[k]})) \quad j, k = 1, ..., m, j \neq k$$

The latter equation for a given pair j, k is a requirement for $\rho_j \phi_j$, $\rho_k \phi_k$ to transform the $|V^{[j]}| \times |V^{[k]}|$ -matrix $M_1^{[j,k]}$ generated by $G(V^{[j]} \cup V^{[k]}, E_1(V^{[j]}, V^{[k]}))$ as the labeled (according to the given colors) adjacency matrix between its parts $V^{[j]}$, $V^{[k]}$ into the corresponding matrix $M_2^{[j,k]}$ generated by $G(V^{[j]} \cup V^{[k]}, E_2(V^{[j]}, V^{[k]}))$ – while $\rho_j \phi_j$ acts as a permutation of $M_1^{[j,k]}$, s rows and $\rho_k \phi_k$ as a permutation of $M_1^{[j,k]}$, s columns.

We can also consider the compacted weighted graphs $Com(G_1)$, $Com(G_2)$ with the common vertex set $\{V^{[1]}, ..., V^{[m]}\}$ where the weight-graph of an existing edge $(V^{[j]}, V^{[k]})$ is defined, for $j \neq k$, as $G(V^{[j]} \cup V^{[k]}, E_1(V^{[j]}, V^{[k]}))$ for $Com(G_1)$ and as $G(V^{[j]} \cup V^{[k]}, E_1(V^{[j]}, V^{[k]}))$ for $Com(G_2)$, and the loop $(V^{[i]}, V^{[i]})$'s weight-graph (or the loop-graph of $V^{[i]})$ is defined as $G(V^{[i]}, E_1(V^{[i]}))$ for $Com(G_1)$ and as $G(V^{[i]}, E_2(V^{[i]}))$ for $Com(G_2)$.

One interesting partial case of such a construction is a pair of bipartite weighted graphs $Com(G_1)$, $Com(G_2)$ where the two connected parts the vertex set $\{V^{[1]}, \dots, V^{[m]}\}$ is partitioned into are interpreted as the set of "variables" and the set of "equations", while for each "variable" its loop-graph is an empty graph, its set of initial vertices is partitioned, in both graphs, into two q-subsets (the same for $Com(G_1)$, $Com(G_2)$) interpreted as "0" and "1", and the weight-graph of the edge between it and an "equation" it's connected with is two disconnected copies of $K_{q,q}$ whose parts in the "variable" are its sets "0" and "1" and whose parts in the "equation" are two disjoint q-subsets (generically different for $Com(G_1)$, $Com(G_2)$) interpreted as the "0-impact" and "1-impact" of this "variable" in the "equation" correspondingly, possibly with some non-incident (to the edge's weight-graph's initial edges) initial vertices in the "equation". In case if no "impacts" of two "variables" intersect in any "equation", we'll get the following problem: is it possible to assign to every "variable" a Boolean value so that for each "equation" there exists an isomorphism from its loop-graph in $Com(G_1)$ to its loop-graph in $Com(G_2)$ mapping, for each "variable" it's connected with, the "variable's" "impacts" in the equation into themselves if its value is 0 and to each other if it's 1. We can accordingly formulate the problem of *impacted isomorphism*: whether there is a couple of graphs with a common vertex set partitioned into pairs of subsets, with at least three pairs, such that for any permutation of the set of those subsets mapping exactly one pair to each other and all the other subsets to themselves there exist an isomorphism of the first graph to the second one compatible with that permutation, but there is no such isomorphism compatible with any other permutation of the set of those subsets mapping each subset either to itself or to the other subset of its pair. If it's true then we can polynomial-time reduce the NP-complete 3-dimensional matching problem to partition the set $X \cup Y \cup Z$, where X, Y, Z are pair-wise disjoint sets of the same cardinality, into triples from a given set $T \subseteq X \times Y \times Z$ (this problem remains NP-complete when each element of $X \cup Y \cup Z$ is incident to no more than 3 triples) to a graph isomorphism problem via associating "variables" with the triples and "equations" with the elements of $X \cup Y \cup Z$, hence proving the graph isomorphism problem's NP-completeness.

Taking into account the fact that the proposed graph with "variables" and "equations" is just one of the simplest cases of the above-stated compacted graph construction, we accordingly receive an apparatus for formulating a variety of conjectures for graph isomorphism (like the statement that such a couple of graphs doesn't exist) whose incorrectness therefore implies the equality GI = NP.

A polynomial-time heuristic approach

to the clique problem

The general types of techniques and notions that we applied when dealing with the graph isomorphism problem could also be applicable in resolving the much more important problems

of determining a graph's clique number and finding its maximum clique, i.e. for the clique problem which is well-known to be NP-complete, unlike the graph isomorphism one.

Given a simple undirected graph G with n vertices whose adjacency matrix is A, we're going to build a sequence of real-valued n×n-matrices $X^{(q)} = X^{(q)}(G)$ (whose columns we'll consider as the coordinate-vectors of n points in \mathbb{R}^n , while the j-th point we'll associate with the vertex j of G) by the following recursive scheme:

$$\mathbf{X}^{(0)} = \mathbf{I}_{\mathbf{n}}$$

Beginning with q = 1, at the step q of our recursive process, we define for j =1,...,n

$$(*) \quad x_{j}^{(q+1)} = x_{j}^{(q)} + g \sum_{k \in \{1, \dots, n\} \setminus \{j\}} \frac{\epsilon_{a_{jk}}}{d^{s}(x_{j}^{(q)}, x_{k}^{(q)})} (x_{k}^{(q)} - x_{j}^{(q)})$$

where $x_j^{(q)}$ is the j-th column of $X^{(q)}$, h, s, ε are the proposed algorithm's parameters (h, s, $\varepsilon > 0$), and for two n-vectors y, z **d**(y,z) denotes the Euclidean distance between them in \mathbb{R}^n .

The above recursive relation is, in fact, a computational circuit for numerically solving the autonomous system of differential equations

$$\dot{x}_{j}(t) = g \sum_{k \in \{1, \dots, n\} \setminus \{j\}} \frac{a_{jk}}{\boldsymbol{d}^{s} \left(x_{j}(t), x_{k}(t) \right)} (x_{k}(t) - x_{j}(t))$$

and the chief idea standing behind it is the conjecture that after a certain period of time since the beginning of the initial point system's contraction under the influence of the introduced "gravitational forces" between pairs of points connected, as vertices, by G's edges the smallest distance will always appear between a pair of vertices belonging to a maximum clique of G.

A special interest this conjecture presents for the case of regular and semi-regular graphs, particularly for the graph generated by a CNF so that its vertex set is the CNF's set of literals and a pair of literals isn't to be connected by an edge if and only if either both of them belong to one disjunctive clause or they're the opposite powers of one variable (this graph's clique number equals the CNF's number of disjunctive clauses if and only if the CNF is satisfiable and is smaller otherwise, and the graph becomes close to regular upon bounding by constants the CNF's quantity of literals in a disjunctive clause and the number of times a variable can occur in literals, while it's regular when the two numbers are just constants, i.e. same for all the clauses and all the variables correspondingly). It's actually known that the clique problem is NP-complete for regular graphs, even when restricted to the case of graphs complimentary to cubic planar ones.

Hence such an algorithm is supposed to perform a certain polynomial (in n) number of steps (*) (considered as the algorithm's functional parameter), determine a pair of vertices (u,v) of the smallest distance, and construct the graph $G_{N_G(u,v)}$ received from G as the graph induced by the set $N_G(u,v)$ of common neighbors of u and v in G. (We additionally conjecture that, after a certain period of time since the gravitational contraction's beginning, any pair of vertices of the

smallest distance is to belong to a maximum clique, although we may also suppose that graphs with no automorphism should get just one such pair at each moment). After that the whole process is to be repeated for $G_{N_G(u,v)}$ etc. until we get either an empty graph or a graph with one vertex.

An additional idea for enhancing the above approach to the clique problem may be introducing "repelling forces" (aka "anti-gravitational") between pairs of points that aren't connected, as vertices, in G. In such a case we'll receive the equation

$$\begin{split} \dot{x}_{j}(t) &= g \sum_{k \in \{1, \dots, n\} \setminus \{j\}} \frac{a_{jk}}{\mathbf{d}^{s} \left(x_{j}(t), x_{k}(t) \right)} (x_{k}(t) - x_{j}(t)) - \\ &- g_{1} \sum_{k \in \{1, \dots, n\} \setminus \{j\}} \frac{1 - a_{jk}}{\mathbf{d}^{s_{1}} \left(x_{j}(t), x_{k}(t) \right)} (x_{k}(t) - x_{j}(t)) \end{split}$$

with the corresponding computational circuit for numerical solving

 $(**) \quad x_{j}^{(q+1)} = x_{j}^{(q)} + g \sum_{k \in \{1, \dots, n\} \setminus \{j\}} \frac{\epsilon_{a_{jk}}}{d^{s}(x_{j}^{(q)}, x_{k}^{(q)})} (x_{k}^{(q)} - x_{j}^{(q)}) -$

$$-g_1 \sum_{k \in \{1, \dots, n\} \setminus \{j\}} \frac{\epsilon(1-a_{jk})}{d^{s_1}(x_j^{(q)}, x_k^{(q)})} (x_k^{(q)} - x_j^{(q)})$$

Let's call g and g_1 the gravitation and anti-gravitation coefficients correspondingly.

Besides, we can notice that the matrices $X^{(q)}(G)$ are, of course, invariant under any of G's automorphisms and, given two graphs G_1 and G_2 , we accordingly may also use $X^{(q)}(G_1)$ and $X^{(q)}(G_2)$ for determining if they're isomorphic (via comparing $MSp(X^{(q)}(G_1))$ and $MSp(X^{(q)}(G_2))$ and eventually obtaining a pair of final algebras upon their entry spectrums' ceasing to grow), but, nevertheless, in such a case we're actually supposed to figure out whether it's not a partial case of the initial graph's adjacency matrix's modification via a series of meta-polynomial and entry spectrum replacement transformations, though, in any case, $X^{(q)}(G)$ is, of course, an isomorphism-commutative transformation of G's adjacency matrix.

The proposed algorithm for the clique problem had been tested (via computer modeling) on graphs received (as described above) from random CNF samples with several hundred Boolean variables whose maximal number of literals in a clause and maximal number of a variable's occurrence was 3 and showed correctness and polynomial-time performance in finding, in case of the CNF's satisfiability, a satisfying Boolean vector.

And, at last, it would be worth noting that any kind of approach to the clique problem can be further enhanced with a randomization parameter via embedding the given graph G with n vertices whose clique number we need to determine into the graph received from G through *bipartitely gluing* it with a random graph G_1 with n_1 vertices whose clique number we know, -- while we define the *bipartite gluing* of two graphs G = (V, E), $G_1 = (V_1, E_1)$ for disjoint V, V_1 as the graph $G \oslash G_1 = (V \cup V_1, E \cup E_1 \cup K_{V,V_1})$ where K_{V,V_1} is the complete bipartite graph on

the parts V, V_1 . The clique number of $G \oslash G_1$ obviously equals the sum of the clique numbers of G and G_1 and this gluing is $(k + n_1)$ -regular when G is k-regular, G_1 is k_1 -regular and $n + k_1 = n_1 + k$. Hence, in case if our algorithm works out on a certain sufficiently large fraction of q-regular graphs with m vertices for a certain set of values of the ratio q:m (containing NPcomplete cases), we can also conjecture that such a random gluing may be quite capable of resolving, via the proposed "gravitational" algorithm, the most hard cases of instances with a sufficiently high (for being a polynomial-time randomized computational circuit) probability of success. However, the general direction of research regarding the above-stated gravitation contraction model may, of course, be rather related to attempting to understand the behavior of its differential equation's solution and even trying, on the basis of such understanding, to reduce the algorithm's polynomial-time complexity through modifying its vertex selection criterion for to take, at each global step, not just one pair, but a much bigger set of vertices as supposedly belonging to a maximum clique.

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