# Exact Algorithms for Finding Well-Connected 2-Clubs in Sparse Real-World Graphs: Theory and Experiments 

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## 2-Club Problem

Input: An undirected graph $G=(V, E)$.
Task: Find the maximum size 2-club (= diameter-two subgraph) in $G$.


- proposed as clique relaxation in social network analysis
[Mokken; Quality and Quantity, 1979]
- NP-hard [Balasundaram, Butenko, Trukhanov; Journal of Combinatorial Optimization, 2005]
- NP-hard to approximate within a factor $|V|^{1 / 2-\epsilon}$
[Asahiro, Doi, Miyano, Samizo, Shimizu; Algorithmica, 2018]


## 2-Club is hard? Not in practice!

Existing implementation: [Hartung, Komusiewicz, N.; Journal of Graph Algorithms and Applications, 2015] Data set: Graphs from clustering and coauthor category of the 10th DIMACS challenge Implementation: Written in Java Machine: CPU 3.60 GHz (Xeon); 64 GB main memory



## Analysis: 2-Club size



## $\rightsquigarrow$ Almost optimal algorithm:

Return a maximum degree vertex with its neighbors

## 2-Club extensions

Definition ([Veremyev, Boginski; Eur J Oper Res, 2012])
$t$-robust 2-club $G$ :
Any pair of vertices is connected by $t$ internally vertex-disjoint paths of length at most two.

Definition ([Pattillo, Youssef, Butenko; Eur J Oper Res, 2013]) $t$-hereditary 2-club $G$ :
$G-U$ is a 2-club for all $U \subset V(G)$ where $|U| \leq t$. $\Longleftrightarrow$ any pair of nonadjacent vertices has $t+1$
 common neighbors.

Definition ([Pattillo, Youssef, Butenko; Eur J Oper Res, 2013]) $t$-connected 2-club $G$ :
$G$ is a 2-club and $t$-vertex-connected.

Example: A $K_{3,3}$ is a

- 1-robust 2-club
- 2-hereditary 2 -club
- 3-connected 2-club


## Our results

## Goal:

transfer algorithmic work (theoretical \& practical) from 2-Club to $t$-robust / $t$-hereditary / $t$-connected 2-Club

## Results:

- "unifying view" on all three considered models
- FPT algorithms
- competitive implementation


## Simple search tree

Example: Find largest 2-hereditary 2-club (deleting any 2 vertices yields a 2 -club)

Observation: At most one red vertex in a solution.

Generic Search tree:

## FindSolution ( $G$ )



1. If $G$ is a solution then return $G$
2. $u, v \leftarrow$ two "incompatible" vertices

## 3. Return $\max \{$ FindSolution $(G-v)$, FindSolution $(G-u)\}$

$\rightsquigarrow$ running time $O\left(2^{\ell} n m\right) \quad \ell \ldots$ number of vertices not in a solution
Note: no $2^{(1-\varepsilon) \ell} n^{O(1)}$ algorithm for any $\varepsilon>0$, unless SETH fails
[Hartung, Komusiewicz, N.; Journal of Graph Algorithms and Applications, 2015]

## Compatible vertices - unifying view

## Definition

Two vertices $v$ and $w$ in a graph are called compatible

- for $t$-robust 2-clubs if they are adjacent and have at least $t-1$ common neighbors, or if they have at least $t$ common neighbors,
- for $t$-hereditary 2-clubs if they are adjacent or if they have at least $t+1$ common neighbors,
- for $t$-connected 2-clubs if they are at distance at most two and are connected by at least $t$ internally vertex-disjoint paths.


## Turing Kernelization



1. sol $\leftarrow \emptyset$
2. foreach $v \in V$ do
3. $\quad T \leftarrow$ all vertices at distance $\geq 2$ from $v$
4. $\quad S \leftarrow$ largest solution in $T$ that contains $v$
5. if $S$ is larger than sol then sol $\leftarrow S$
6. delete $v$
7. return sol

## Turing kernelization - practical effect



$$
\begin{aligned}
& -*-n \\
& -\infty \\
& -\infty-\text { maximum degree } \\
& -*-\text { average degree } \\
& -\infty \text { maximum 2-neighborhood } \\
& -*-\text { average 2-neighborhood } \\
& \rightarrow-h_{2} \text {-index }
\end{aligned}
$$

## Advantage:

Turing kernelization allows to store data for each pair of vertices (e.g. number of common neighbors)

## Data reduction \& lower bounds

## Reduction Rule

Remove vertices whose degree is too low.
Incompatibility graph:
Two vertices are adjacent in the the incompatibility graph iff they are not compatible.

input graph

incompatibility graph

## Observation:

The size of a maximum independent set in the incompatibility graph is an upper bound on the solution size in the input graph.
$\rightsquigarrow$ upper bound worse than best previously found solution $\Rightarrow$ discard current Turing kernel

## Experiments I

Data set: Graphs from clustering and coauthor category of the 10th DIMACS challenge Implementation: Written in Java Machine: CPU 3.60 GHz (Xeon); 64 GB main memory


## Experiments II

Data set: Graphs from clustering and coauthor category of the 10th DIMACS challenge Implementation: Written in Java Machine: CPU 3.60 GHz (Xeon); 64 GB main memory


## Experiments III

$$
\begin{aligned}
& \rightarrow(t-1) \text {-hereditary } * t \text {-robust } \\
& -t \text {-connected }
\end{aligned}
$$




Graph: coPapersCiteseer

## Experiments IV

$$
\begin{aligned}
& \rightarrow(t-1) \text {-hereditary } * t \text {-robust } \\
& \rightarrow-t \text {-connected }
\end{aligned}
$$




Graph: coAuthorsCiteseer

## Summary \& Outlook

## Key results:

- Unifying approach for several 2-club variants.
- Efficient implementation (= data reduction + Turing kernelization + search tree).

Work in progress: $\gamma$-relative robust 2-club $S$ :
$0<\gamma \leq 1$ : Any pair of vertices connected by at least $\gamma \cdot|S|$ paths of length at most two.
Example: $\gamma=0.5$

input graph

incompatibility graph

input graph
$\square$
incompatibility graph

Open Question: Is $t$-robust / $t$-hereditary / $t$-connected 2-club fixed parameter tractable with respect to the solution size?

## Thank you!

