Knots, graphs and Khovanov homology II

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Turaev surface

Ribbon graphs

Quasi-trees

Homological width

Conclusion

Let *D* be a link diagram and let s_A and s_B be the all-*A* and all-*B* states of *D*.

Turaev constructed a cobordism between s_A and s_B :

Let $\Gamma \subset S^2$ be the 4-valent projection of D at height 0. Put s_A at height 1, and s_B at height -1, joined by saddles:



Turaev surface



Turaev surface F(D): Attach $|s_A| + |s_B|$ discs to all boundary circles above.

Turaev genus of D, $g_T(D) := g(F) = (c(D) + 2 - |s_A| - |s_B|)/2$.

Turaev genus of non-split link $L := g_T(L) = \min_D g_T(D)$.

Properties

- Non-split link L is alternating iff $g_T(L) = 0$.
- *D* is alternating on the Turaev surface.
- *g*_T(*L*) ≤ *dalt*(*L*) = min number of crossing changes to make *L* alternating.
- Measures "distance" from alternating.
- Turaev surface can be constructed for any two complementary states of *D*.

Conjecture (Tait) A reduced alternating diagram D has minimal crossing number among all diagrams for the alternating link L.

The proof follows from three claims:

- Although defined for diagrams, the Jones polynomial V_L(t) is a link invariant.
- s_A and s_B contribute the extreme terms $\pm t^{\alpha}$ and $\pm t^{\beta}$ of $V_L(t)$.
- span V_ℓ(t) = α − β ≤ c(ℓ) − g_T(ℓ), with equality if ℓ is alternating (generally, *adequate*).

Since D is alternating on the Turaev surface, we can generalize the Tait graph construction to get graphs on surfaces.

The Turaev surface F(D) can be checkerboard colored with $|s_A|$ white regions (height > 0), and $|s_B|$ black regions (height < 0).

Let G_A , $G_B \subset F(D)$ be the adjacency graphs for respective regions. G_A and G_B are embedded and

$$v(G_A) = |s_A|, \quad e(G_A) = c(D), \quad f(G_A) = |s_B|$$

 G_A (and G_B) give a cell decomposition of F(D).

If D is alternating, G_A and G_B are dual Tait graphs on $F(D) = S^2$.

Example: Pretzel links

Let $p_i, q_j \in \mathbb{N}$. The pretzel link $P(p_1, \ldots, p_n, -q_1, \ldots, -q_m)$, is a link with diagram of the form



• If m = 0, then the pretzel link is alternating and $g_T = 0$.

If m > 0 then the pretzel link is non-alternating and can be embedded on the torus. Hence g_T = 1. An (oriented) ribbon graph G is a multi-graph (loops and multiple edges allowed) that is embedded in an oriented surface F, such that its complement is a union of 2-cells. The genus g(G) := g(F).

Example





G can also be described by a triple of permutations $(\sigma_0, \sigma_1, \sigma_2)$ of the set $\{1, 2, \ldots, 2n\}$ such that

- σ_1 is a fixed-point-free involution.
- $\sigma_0 \circ \sigma_1 \circ \sigma_2 =$ Identity

This triple gives a cell complex structure for the surface of G such that

- Orbits of σ_0 are vertices.
- Orbits of σ₁ are edges.
- Orbits of σ_2 are faces.

The genus
$$g(G) = (2 - v(G) + e(G) - f(G))/2$$
.

Ribbon graph example





$$\sigma_0 = (1234)(56)$$

$$\sigma_1 = (14)(25)(36)$$

$$\sigma_2 = (1)(246)(35)$$

$$\sigma_0 = (1234)(56)$$

 $\sigma_1 = (13)(26)(45)$
 $\sigma_2 = (152364)$





Ribbon graph from any state of a link diagram

 G_A , G_B defined earlier as checkerboard graphs on Turaev surface F(D) were ribbon graphs. We can construct the ribbon graph G_s directly from any state *s* of *D*:

- 1. For each crossing of D, attach an edge between state circle(s).
- 2. Collapse each state circle of s to a vertex of G_s .



1. (2001) Bollobás and Riordan extended the Tutte polynomial to an invariant of oriented ribbon graphs, Bollobás–Riordan–Tutte polynomial.

2. (2006) Dasbach, Futer, Kalfagianni, Lin, and Stoltzfus showed that $V_L(t)$ can be recovered from BRT polynomial of G_A .

Example: From diagram to Tait graph and ribbon graph



From spanning trees to quasi-trees

For a planar graph, a spanning tree is a spanning subgraph whose regular neighbourhood has one boundary component.

Example





A quasi-tree of a ribbon graph is a spanning ribbon subgraph with one face. The genus of a quasi-tree is its genus as a ribbon graph.

Example: Genus one quasi-tree of a genus two ribbon graph.



Quasi-trees and Chord diagrams

Every quasi-tree corresponds to an ordered chord diagram,



Let *D* be a connected link diagram, *G* its Tait graph, G_A its all-*A* ribbon graph.

Theorem. (C-Kofman-Stoltzfus) Quasi-trees of G_A are in one-one correspondence with spanning trees of G:

 $\mathbb{Q}_j \leftrightarrow T_v$ where $v + j = (V(G) + E_+(G) - V(G_A))/2$

 \mathbb{Q}_j is quasi-tree of genus j, and \mathcal{T}_v is spanning tree with v positive edges.

Graphs and Ribbon graphs

Graphs	\longleftrightarrow	Ribbon graphs
Spanning trees	\longleftrightarrow	Quasi-trees
Tutte polynomial	\longleftrightarrow	BRT polynomial
Activity w.r.t. spanning trees	\longleftrightarrow	Activity w.r.t. quasi-trees
Spanning tree expansion of the Tutte polynomial	\longleftrightarrow	Quasi-tree expansion of the BRT polynomial

Quasi-trees and Khovanov homology

Theorem (C-Kofman-Stoltzfus) For a knot diagram D, there exists a quasi-tree complex $\mathbb{C}(G_A) = \{\mathbb{C}_v^u(G_A), \partial\}$ that is a deformation retract of the reduced Khovanov complex, where

$$\mathbb{C}_{v}^{u}(G_{A}) = \mathbb{Z}\langle \mathbb{Q} \subset G_{A} | u(\mathbb{Q}) = u, -g(\mathbb{Q}) = v \rangle$$

From above, if \mathbb{Q}_j is quasi-tree of genus j, and T_v is spanning tree with v positive edges,

$$v + j = (V(G) + E_+(G) - V(G_A))/2$$

Corollary (C-Kofman-Stoltzfus) For any knot K, the width of its reduced Khovanov homology $w_{KH}(K) \leq 1 + g_T(K)$.

Proof. For any ribbon graph G, $g(G) = \max_{\mathbb{Q} \subset G} g(\mathbb{Q})$. Therefore, the quasi-tree complex $\mathbb{C}(G_A)$ has at most $1 + g(G_A)$ rows.



Turaev genus and homological width

- Using w_{KH}(K), we get lower bounds for g_T(K). In particular, g_T(T(3,q)) → ∞.
- For an adequate knot K with an adequate diagram D, T. Abe showed g_T(K) = g_T(D) = w_{KH}(K) − 1 = c(K) − spanV_K(t).
- Dasbach and Lowrance also proved bounds in terms of g_T(K) for the Ozsváth-Szabó τ invariant and the Rasmussen's s invariant.
- ► Similar bounds for homological width of knot Floer homology in terms of g_T(K) were obtained by Adam Lowrance.

Related open problems

- Find families of homologically thin knots with g_T(K) > 1? Generally, are there any lower bounds independent of knot homology?
- Which operations on knots preserve or increase Turaev genus? For e.g. for adequate knots g_T(K#K') = g_T(K) + g_T(K') and g_T is preserved under mutation. How about non-adequate knots?
- 3. How is the Turaev genus related to the topology and hyperbolic geometry of knot complements ?

Computer programs to study Knots & Links

- SnapPy (study hyperbolic knots, links and 3-manifolds) by Weeks, Culler and Dunfield.
- ▶ Knotscape (old program to study knots) by Thistlethwaite.
- knot by Kodama.
- LinKnot by Jablan and Sazdanovic.
- KnotTheory by Dror Bar-Natan.
- KhoHo by Shumakovitch to compute Khovanov Homology.
- KnotAtlas (database) by Dror Bar-Natan.
- Table of Knot Invariants (database) by Livingston.

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Questions

Thank You

Slides available from : http://www.math.csi.cuny.edu/abhijit/