1 Description

The first we just calculated several Barycentric Cubical Tori. The distribution of calculated toruses we can see in the Figure 1a.

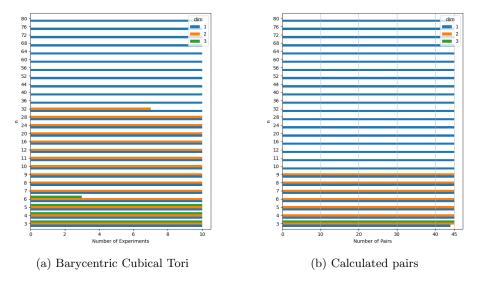


Figure 1: Size/dimension distribution

Then for each pair of filtrations with similar dimension and n $f_0, f_1 : \mathbb{T}_n^d \to \mathbb{R}$ we can define a linear homotopy $h : [0,1] \times \mathbb{T}_n^d \to \mathbb{R}$:

$$h(t,\sigma) = (1-t) \cdot f_0(\sigma) + t \cdot f_1(\sigma)$$

The moment of time $t \in [0,1]$ such there exist a pair of cells $\sigma_0, \sigma_1 \in \mathbb{T}_n^d$ such that $h(t,\sigma_0) = h(t,\sigma_1)$ is a moment of transposition of these cells during the homotopy (the probability, that this will be full segment is 0 for pair of independent barycentric cubical tori).

We have found all transposing pairs like this, classified them and calculated how these transpositions cannges the Depth Poset. The distribution of calculated pairs of filtration we can see in the Figure 1b.

2 Transpositions

There could be 3 types of cell in the terms of birth-death pairs:

- Births The cells which are births of some cycles
- Deaths The cells which are deaths of some cycles
- Unpaired The cells which are not birth or deaths of any cycle

So we have defined 5 types of transpositions:

- Birth-Birth The switch transpositions of 2 brith cells
- Birth-Death The switch transpositions of brith and death cells
- Death-Death The switch transpositions of 2 death cells
- Undefined The transpositions where at least 1 cell is unpaired
- No switch No switch transposition, where all cells are paired

In the Figure 2 we can see the mean count of transpositions of each type for barycentric cubical tori of different dimensions d and size n. We can see that no switch transpositions are majority of cases. So we also can look at Figure 3 to get how other types are distributed.

The distribution of Transposition Types by the Size of the Torus n0-dim transpositions in \mathbb{T}_n^1 birth-birth death-death 1500 1500 no switch no switch undefined undefined 1000 1000 500 500 50 10 40 0-dim transpositions in \mathbb{T}_n^2 1-dim transpositions in T_n^2 2-dim transpositions in T_n^2 birth-birth birth-birth death-death 6000 1500 1500 birth-death undefined undefined death-death 4000 no switch 1000 1000 2000 500 500 0-dim transpositions in T_n^3 1-dim transpositions in T_n^3 2-dim transpositions in T_n^3 3-dim transpositions in T_n^3 200 1500 1250 150 150 birth-birth birth-birth 1000 birth-birth birth-death birth-death - death-death 100 no switch death-death death-death 100 no switch 750 undefined no switch no switch undefined 500 500 undefined undefined 50 50 250 2.6 2.8 3.2 3.4 2.8 3.0 3.2 2.8 3.0 3.2 2.6 2.8 3.0 3.2 3.4

Figure 2: The mean value of number of transpositions of each type.

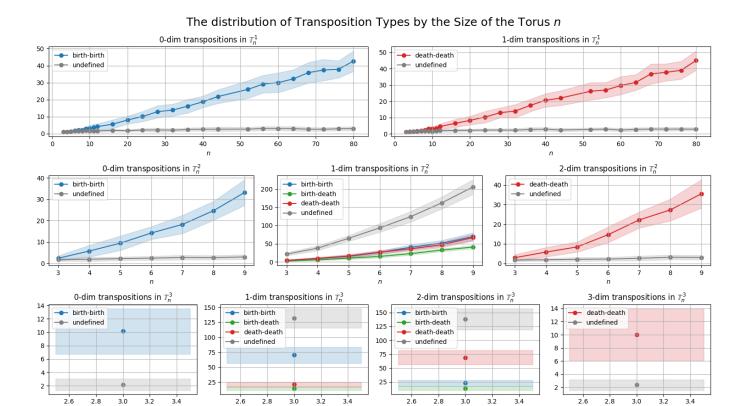


Figure 3: The mean value of number of transpositions of each type (without no switch transpositions).

3 Similarity Scores

We have 10 scores to measure the similarity of 2 depth posets:

- birth_relation_cell_similarity The Jacard index of arcs (edges) in the birth relation (given by row left to right reduction algorithm). Consider 2 birth-death pairs are similar if they corespond the similar cells. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 4
- birth_relation_index_similarity The Jacard index of arcs (edges) in the birth relation (given by row left to right reduction algorithm). Consider 2 birth-death pairs are similar if ther indices in the filtration pairs are similar. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical

tori dimension d and size n in Figure 5

- death_relation_cell_similarity The Jacard index of arcs (edges) in the death relation (given by column bottom to top reduction algorithm). Consider 2 birth-death pairs are similar if they corespond the similar cells. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 6
- death_relation_index_similarity The Jacard index of arcs (edges) in the death relation (given by column bottom to top reduction algorithm). Consider 2 birth-death pairs are similar if ther indices in the filtration pairs are similar. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 7
- poset_arc_cell_similarity The Jacard index of arcs (edges) in the Depth Poset. Consider 2 birth-death pairs are similar if they corespond the similar cells. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 8
- poset_arc_index_similarity The Jacard index of arcs (edges) in the Depth Poset. Consider 2 birth-death pairs are similar if ther indices in the filtration pairs are similar. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 9
- poset_node_cell_similarity The Jacard index of nodes (elements) in the Depth Poset. Consider 2 birth-death pairs are similar if they corespond the similar cells. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 10
- poset_node_index_similarity The Jacard index of nodes (elements) in the Depth Poset. Consider 2 birth-death pairs are similar if ther indices in the filtration pairs are similar. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 11
- relation_cell_similarity The Jacard index of arcs (edges) in the relation (given by reduction algorithms). Consider 2 birth-death pairs are similar if they corespond the similar cells. We can see the values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 12
- relation_index_similarity The Jacard index of arcs (edges) in the relation (given by reduction algorithms). Consider 2 birth-death pairs are similar if ther indices in the filtration pairs are similar. We can see the

values of this score applying to depth posets before and after transpositions of different types for barycentric cubical tori dimension d and size n in Figure 13

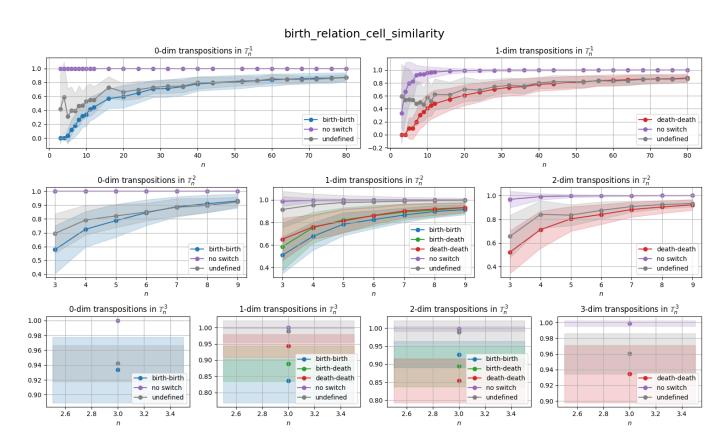


Figure 4: birth_relation_cell_similarity values

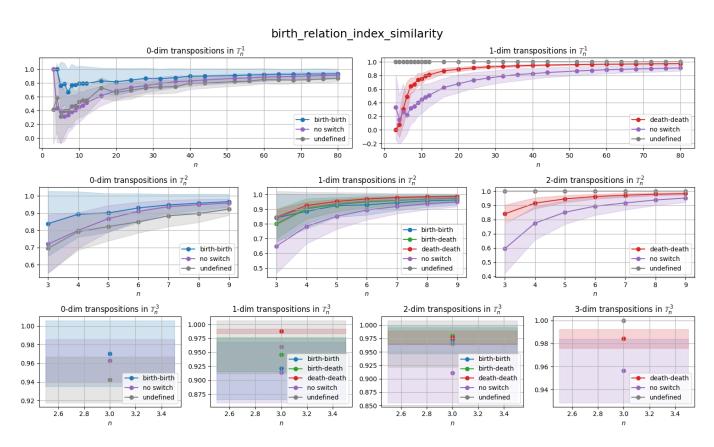


Figure 5: $birth_relation_index_similarity$ values

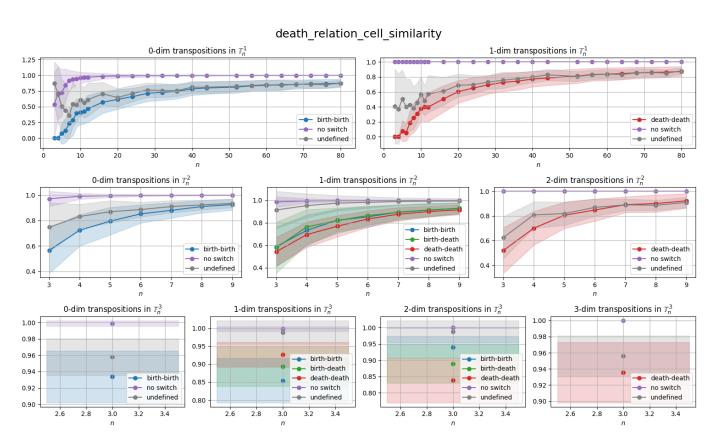


Figure 6: death_relation_cell_similarity values

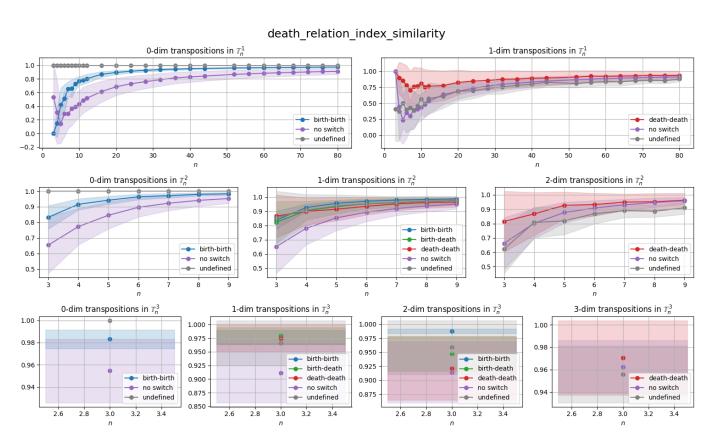


Figure 7: $death_relation_index_similarity$ values

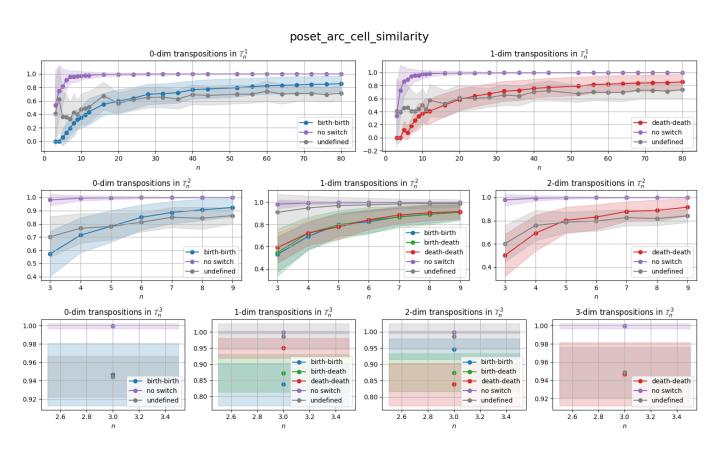


Figure 8: poset_arc_cell_similarity values

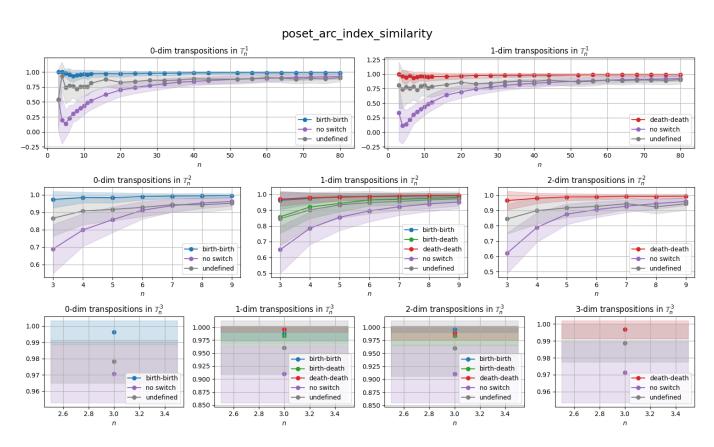


Figure 9: poset_arc_index_similarity values

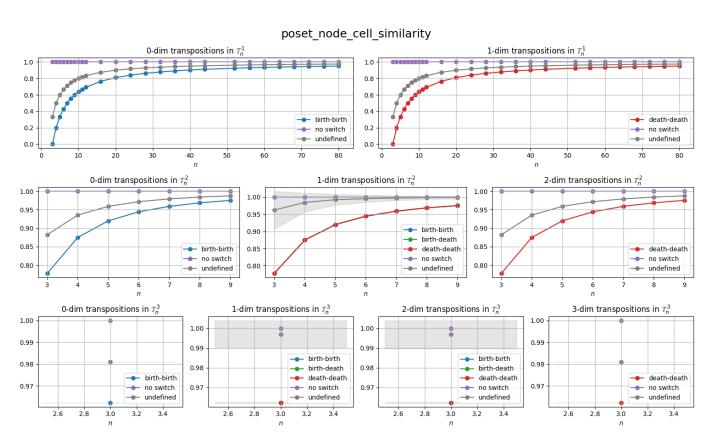


Figure 10: poset_node_cell_similarity values

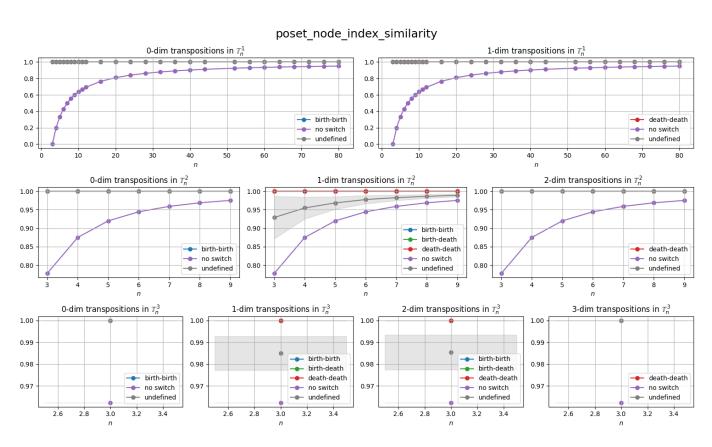


Figure 11: poset_node_index_similarity values

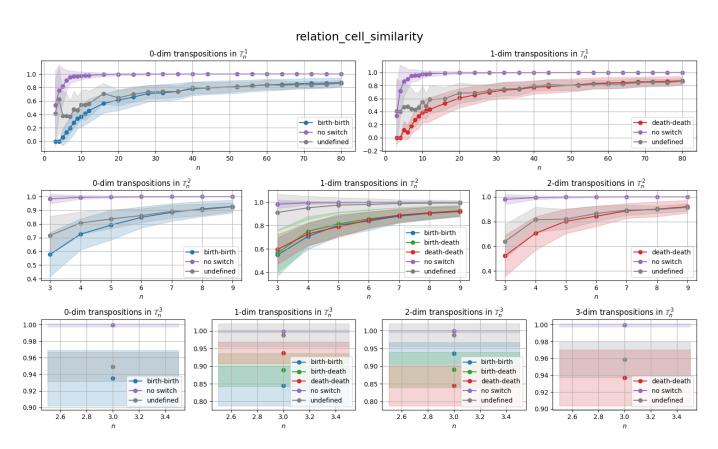


Figure 12: relation_cell_similarity values

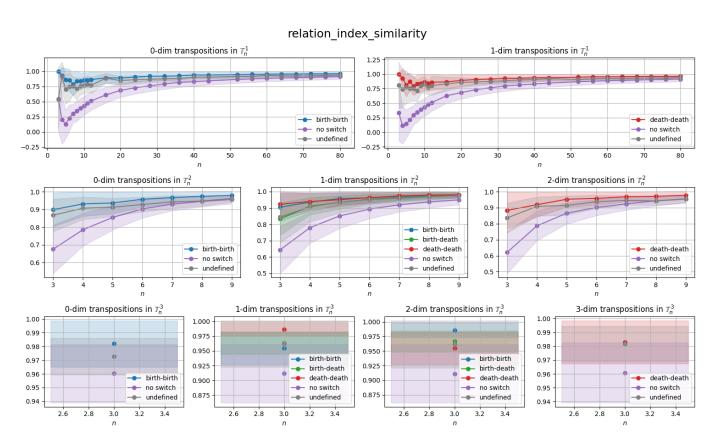


Figure 13: relation_index_similarity values