# Selected topics in representation theory 2 The braid group action on the set of exceptional sequences $\frac{\text{WS }2005/06}{\text{WS }2005/06}$

#### 1 The braid group

**Definition.** Let  $n \geq 2$  be a natural number and define

$$B_n := \langle \sigma_i \mid 1 \leq i < n; \quad \sigma_i \sigma_{i+1} \sigma_i = \sigma_{i+1} \sigma_i \sigma_{i+1}, \quad \sigma_i \sigma_j = \sigma_j \sigma_i \text{ for } |i-j| \geq 2 \rangle$$

the group generated by n-1 elements, with relations as above.

**Remark.** The groups  $B_2$  is isomorphic to  $\mathbb{Z}$ . Morover, there exists a surjective group homomorphism  $B_n \longrightarrow S_n$  onto the symmetric group. There is a topological interpretation of this group in terms of equivalence classes of diagrams of braids.

### 2 The action of the braid group

We consider exceptional sequences of objects in a triangulated category  $\mathcal{D}^b$ , as introduced in the first lecture. In particular, we assume that all left and right mutations are defined in this category. We denote by  $\operatorname{Hom}^{\bullet}(E,F)$  the finite dimensional vector space  $\bigoplus_{l}\operatorname{Hom}(E,F[l])$ . Moreover, we write  $\varepsilon = (E_1,\ldots,E_r) \simeq \varepsilon' = (E_1,\ldots,E_r')$  if  $E_i \simeq E_i'$  for all  $i=1,\ldots,r$ .

**Theorem.** Let  $\varepsilon = (E_1, \dots, E_r)$  be an exceptional sequence. Then

- 1)  $L_i R_i \varepsilon \simeq R_i L_i \varepsilon \simeq \varepsilon$ ,
- 2)  $L_i L_{i+1} L_i \varepsilon \simeq L_{i+1} L_i L_{i+1} \varepsilon$ ,  $R_i R_{i+1} R_i \varepsilon \simeq R_{i+1} R_i R_{i+1} \varepsilon$ ,
- 3)  $L_i L_j \varepsilon \simeq L_j L_i \varepsilon$ , and  $R_i R_j \varepsilon \simeq R_j R_i \varepsilon$  for all  $|i j| \ge 2$ .

Corollary. The braid group  $B_r$  acts on the set of all exceptional sequences of length r via  $\sigma_i \varepsilon := L_i \varepsilon$  and  $\sigma_i^{-1} \varepsilon := R_i \varepsilon$ .

PROOF. 3) is obvious. We first prove 1). We consider the triangle defining the left mutation of a pair  $\varepsilon = (E, F)$ , the right mutation of the pair  $(L_E F, E)$ , and note that for  $L\varepsilon = (L_E F, E)$  we have already shown  $\operatorname{Hom}^{\bullet}(E, F) \simeq \operatorname{Hom}^{\bullet}(L_E F, E)^*$ :

The two triangles are obviously isomorphic in two positions, consequently we obtain  $F \simeq R_E L_E F$ . The other isomorphism can be shown analogously.

To show 2) we consider first the mutations of a triple (E, F, G) and obtain

$$L_1L_2L_1(E, F, G) = (L_{L_F}L_EG, L_EF, E)$$
 and  $L_2L_1L_2(E, F, G) = (L_EL_FG, L_EF, E)$ .

So we need to show  $L_{L_EF}L_EG \simeq L_EL_FG$ . For we need the following lemma.

**Lemma.** Let A be an exceptional object. Then  $L_A$  is a functor  $\mathcal{D}^b \longrightarrow \mathcal{D}^b$ .

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PROOF. Let  $E \longrightarrow F$  be a homomorphism. We can define the objects  $L_AE$  and  $L_AF$  as the mapping cone of the corresponding canonical map. Then there exists (using one of the axioms of triangulated categories) a map  $L_AE \longrightarrow L_AF$ . We have to show that this map is unique. For we consider the commutative diagram with triangles in the rows

$$E[-1] \longrightarrow L_A E \xrightarrow{can^*} \operatorname{Hom}(A, E) \otimes A \xrightarrow{can} E$$

$$\downarrow \qquad \qquad \downarrow \qquad \qquad \downarrow \qquad \qquad \downarrow$$

$$F[-1] \longrightarrow L_A F \xrightarrow{can^*} \operatorname{Hom}(A, F) \otimes A \xrightarrow{can} F.$$

A homomorphism  $E \longrightarrow F$  defines a homomorphism  $E[-1] \longrightarrow L_A F$  via composition. We show  $\operatorname{Hom}(L_A E, L_A F[l]) \simeq \operatorname{Hom}(E[-1], L_A F[l])$  for all l, so each homomorphism  $E \longrightarrow F[l]$  defines a unique homomorphism  $L_A E \longrightarrow L_A F[l]$ . For we consider the long exact sequence

$$(A \otimes (A, E), L_A F) \longrightarrow (L_A E, L_A F) \longrightarrow (E[-1], L_A F) \longrightarrow (A[-1] \otimes (A, E), L_A F).$$

Since  $(A[l], L_A F) = 0$  for all l we obtain the desired isomorphism  $(L_A E, L_A F) \longrightarrow (E[-1], L_A F)$ .

We proceed with the proof of the theorem: we have to show  $L_{L_EF}L_EG \simeq L_EL_FG$  for any exceptional sequence (E, F, G). We obtain two triangles, the first one obtained by applying the functor  $L_E$  to the triangle defining the left mutation  $L_FG$ , the second one defines the mutation  $L_{L_EF}L_EG$ 

Since  $\operatorname{Hom}^{\bullet}(F,G) \simeq \operatorname{Hom}^{\bullet}(L_{E}F,L_{E}G)$  both triangles are isomorphic, consequently we have proven the result.

## 3 Exceptional Sequences for hereditary algebras

For finite dimensional hereditary algebras we can use some modification for the mutations of exceptional sequences. Let  $\Lambda$  be a hereditary algebra and C be a complex in the derived category  $\mathcal{D}^b$  of bounded complexes of finitely generated  $\Lambda$ -modules. If C is indecomposable then C is isomorphic to an indecomposable module M up to some unique shift. If we define a stronger equivalence relation for exceptional sequences:  $\varepsilon = (E_1, \ldots, E_r) \sim \varepsilon' = (E'_1, \ldots, E'_r)$  iff  $E_i \simeq E'_i[l]$  for some l, then we can represent each exceptional sequence uniquely, up to isomorphism, by a sequence of modules. Since left and right mutations commute with shifts, we obtain an action of the braid group on the set of all exceptional sequences of  $\Lambda$ -modules. So we have already proven the first assertion of the following result.

**Theorem.** 1) The braid group  $B_r$  acts on the set of all exceptional sequences of  $\Lambda$ -modules of length r (via the modified action).

2) Let n be the number of isomorphism classes of pairwise non-isomorphic simple  $\Lambda$ -modules. Then the braid group  $B_n$  acts transitively on the set of all exceptional sequences of  $\Lambda$ -modules of length n.

The proof of the second part is due to Crawley-Boevey ([1]) for basic algebras and due to Ringel ([3]) in general. Instead of mutations the notion of perpendicular categories, developed by Schofield and Geigle, Lenzing ([2]) is used in the proof. The essential step in the proof shows that via a sequence of mutations we can reach the sequence consisting of simple  $\Lambda$ -modules. This sequences is unique, up to reordering. In fact, there is a natural partial order on the set of sequences, so that each sequence of simples is minimal with respect to this order.

#### References

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