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### THE DIMENSION OF A QUASI-HEREDITARY ALGEBRA

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Quasi-hereditary algebras have been introduced by L. Scott [S] in order to study highest weight categories as they arise in the representation theory of complex Lie algebras and algebraic groups. They have been studied by Cline, Parshall and Scott [CPS], [PS], and in [DR1], [DR2]. Here, we are going to give lower and upper bounds for the dimension of a quasi-hereditary algebra in terms of its species, and we characterize those algebras where one of these bounds is attained: we call them the shallow and the deep quasi-hereditary algebras, respectively.

#### 1. Definitions and results

Let A be a basic semiprimary ring with radical N, let  $e_1, \ldots, e_n$  be a complete set of orthogonal primitive idempotents. The simple right A-module which is not annihilated by  $e_i$  will be denoted by E(i), its projective cover by  $P(i) = P_A(i)$ . The simple left A-module not annihilated by  $e_i$  is denoted by  $E^*(i)$ . The species of A is, by definition,  $\mathcal{S} = \mathcal{S}(A) = (F_i, iM_j)_{1 \le i,j \le n}$ , where  $F_i = e_i A e_i / e_i N e_i$ , and  $iM_j = e_i N e_j / e_i N^2 e_j$ . In our considerations, the total ordering of the index set  $\{1, \ldots, n\}$  of the species will usually be of importance, and in order to stress this, we will speak of a labelled species.

We recall that an ideal J of A is called a heredity ideal provided  $J^2 = J$ , JNJ = 0, and the right module  $J_A$  (or, equivalently, the left module  ${}_AJ$ ) is

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projective. And A is said to be quasi-hereditary provided there exists a chain  $\mathcal{J} = (J_i)_i$  of ideals

$$0 = J_0 \subset J_1 \subset \ldots \subset J_m = A$$

such that  $J_i/J_{i-1}$  is a heredity ideal of  $A/J_{i-1}$ ; such a chain will be called a heredity chain of A. Observe that any heredity ideal J is generated (as an ideal) by an idempotent, and if e is any idempotent in J, then the ideal  $\langle e \rangle$  generated by e is a heredity ideal of A, and  $J/\langle e \rangle$  is a heredity ideal of  $A/\langle e \rangle$ . It follows that we can refine any heredity chain of A to a heredity chain  $\mathcal{J}$  such that, in addition,  $J_i/J_{i-1}$  is generated by a primitive idempotent, and we call such a heredity chain a saturated one. So, let  $\mathcal{J}$  be a saturated heredity chain of A, and we always assume that the idempotents  $e_i$  are chosen in such a way that  $J_i = \langle e_{n-i+1} + \ldots + e_n \rangle$ , for  $0 \leq i \leq n$ . In this way, the quasi-hereditary algebra A together with the fixed saturated heredity chain determines uniquely  $\mathcal{S}(A)$  as a labelled species. Note that  $\mathcal{S}(A)$  is a species without loops.

Assume that A is quasi-hereditary, with heredity chain  $\mathcal{J} = \langle J_i \rangle_i$ , where  $J_i = \langle e_{n-i+1} + \ldots + e_n \rangle$ . Let  $A_i = A/J_{n-i}$ . Note that E(i) and  $E^*(i)$  are  $A_i$ -modules, and we denote their  $A_i$ -projective covers by  $\Delta(i) = \Delta_A(i)$  and  $\Delta^*(i) = \Delta_A^*(i)$ , respectively. Since we deal with a quasi-hereditary algebra, it follows that  $J_i/J_{i-1}$ , as a right A-module, is the direct sum of copies of  $\Delta(n-i+1)$  (so the modules  $\Delta(i)$  are just those modules which occur as building blocks in the standard filtrations of the projective right A-modules: the "Verma modules", or "induced modules"). Similarly,  $J_i/J_{i-1}$  is, as left A-module, the direct sum of copies  $\Delta^*(n-i+1)$ .

By definition, both  $\Delta(i)$  and  $\Delta^*(i)$  are local A-modules. In case all the modules  $\Delta(i)$  and  $\Delta^*(i)$ , with  $1 \le i \le n$ , have Loewy length at most 2, we call A shallow. Thus, A is shallow if and only if all the modules  $\operatorname{rad} \Delta(i)$  and  $\operatorname{rad} \Delta^*(i)$  are semisimple. Observe that these modules are actually  $A_{i-1}$ -modules, and we call A deep provided  $\operatorname{rad} \Delta(i)$  is a projective right  $A_{i-1}$ -module and  $\operatorname{rad} \Delta^*(i)$  is a projective left  $A_{i-1}$ -module, for all  $1 \le i \le n$ .

Now, conversely, let  $\mathscr S$  be a labelled species without loops, say  $\mathscr S=(F_i,{}_iM_j)_{1\leqslant i,j\leqslant n}$ , with  ${}_iM_i=0$  for all i. The tensor algebra  $\mathscr T(\mathscr S)$  can be decomposed as follows. Let T=T(n) be the set of all sequences  $(t_0,\,t_1,\,\ldots,t_m)$  where the  $t_i$  are integers with  $1\leqslant t_i\leqslant n$ , and  $m\geqslant 1$ , such that, moreover,  $t_{i-1}\neq t_i$  for  $1\leqslant i\leqslant m$ . For  $t=(t_0,\,t_1,\,\ldots,t_m)\in T$ , let

$$M(t) = {}_{t_0}M_{t_1} \underset{F_{t_1}}{\otimes} {}_{t_1}M_{t_2} \underset{F_{t_2}}{\otimes} \dots \underset{F_{t_{m-1}}}{\otimes} {}_{t_{m-1}}M_{t_m},$$

and for  $T' \subseteq T$ , let

$$M(T') = \bigoplus_{t \in T'} M(t).$$

Let  $\mathcal{T}_0(\mathcal{S}) = \prod_{i=1}^n F_i$  and  $\mathcal{T}_+(\mathcal{S}) = M(T)$ , thus  $\mathcal{T}(\mathcal{S}) = \mathcal{T}_0(\mathcal{S}) \oplus \mathcal{T}_+(\mathcal{S})$ .

We are going to define two factor algebras of  $\mathcal{F}(\mathcal{S})$  which will turn out to be quasi-hereditary. Both algebras will be of the form  $\mathcal{F}(\mathcal{S})/M(T')$  for suitable choices of T'. In order to define the first one, we define complementary subsets U,  $U^0$  of T as follows: Let

$$U = U(n) = \{(t_0, t_1) \in T\} \cup \{(t_0, t_1, t_2) \in T \mid t_0 < t_1 > t_2\},\$$

thus

$$U^{0} = \mathcal{F} \setminus U = \{(t_{0}, t_{1}, \dots, t_{m}) \in T \mid \text{there is } 0 < i < m \}$$
with  $t_{i} < \max(t_{i-1}, t_{i+1}) \}$ .

Obviously,  $M(U^0)$  is an ideal of  $\mathcal{F}(\mathcal{S})$ , and

$$(\mathscr{T}_{+}(\mathscr{S}))^{3} \subseteq M(U^{0}) \subseteq (\mathscr{T}_{+}(\mathscr{S}))^{2},$$

thus  $M(U^0)$  is an admissible ideal. We define  $S(\mathcal{S}) = T(\mathcal{S})/M(U^0)$ . Note that as abelian groups, we can identify  $S(\mathcal{S})$  and  $\mathcal{T}_0(\mathcal{S}) \oplus M(U)$ .

For the second algebra, we define complementary subsets V,  $V^0$  of T as follows: Let

$$V = V(n) = \{(t_0, \dots, t_m) \in T \mid \text{given } i < j \text{ with } t_i = t_j,$$
 there exists  $l$  with  $i < l < j$  and  $t_i < t_l\}$ , 
$$V^0 = T \setminus V = \{(t_0, \dots, t_m) \in T \mid \text{there are } i < j \text{ with } t_i = t_j \text{ and } t_l < t_i \text{ for all } i < l < j\}.$$

As usual, we may consider a product on T by using the juxtaposition, thus  $(t_0, \ldots, t_m) \cdot (t'_0, \ldots, t'_{m'}) = (t_0, \ldots, t_m, t'_0, \ldots, t'_{m'})$ . Of course, for subsets T', T'' of T, we define  $T' \cdot T'' = \{t' \cdot t'' \mid t' \in T', t'' \in T'' \text{ and } t' \cdot t'' \in T\}$  and so on. Then, obviously, for  $n \ge 2$ 

$$V(n) = V(n-1) \cup V(n-1) \cdot n \cup n \cdot V(n-1) \cup V(n-1) \cdot n \cdot V(n-1).$$

By induction on n, we see that V(n) is finite. In particular, the sequences  $(t_0, \ldots, t_m) \in V(n)$  are of bounded length, say  $m \le v(n)$  for some v(n). Thus

$$(\mathscr{T}_{+}(\mathscr{S}))^{v(n)+1} \subseteq M(V^{0}) \subseteq (\mathscr{T}_{+}(\mathscr{S}))^{2},$$

so that  $M(V^0)$  is an admissible ideal. We define  $D(\mathcal{S}) = \mathcal{F}(\mathcal{S})/M(V^0)$ , and note that  $D(\mathcal{S})$  can be identified, as an abelian group, with  $\mathcal{F}_0(\mathcal{S}) \oplus M(V)$ .

THEOREM 1. Let  $\mathscr G$  be a labelled species without loops. The rings  $S(\mathscr G)$  and  $D(\mathscr G)$  are quasi-hereditary, with labelled species  $\mathscr G$ . The ring  $S(\mathscr G)$  is shallow, the ring  $D(\mathscr G)$  is deep.

In particular, we see that the nonexistence of loops is the only condition on a species for being realizable as the species of a quasi-hereditary ring.

Let k be a (commutative) field. In case  $\mathscr S$  is a finite-dimensional k-species, labelled and without loops, we denote by  $s_k(\mathscr S)$  and  $d_k(\mathscr S)$  the k-dimension of  $S(\mathscr S)$  and  $D(\mathscr S)$ , respectively. We are going to formulate an estimate for the Cartan invariants of a quasi-hereditary algebra A in terms of the Cartan invariants of the corresponding algebras  $S(\mathscr S)$  and  $D(\mathscr S)$ . In this way, we deduce that the dimension of A is bounded from below by  $s_k(\mathscr S)$  and from above by  $d_k(\mathscr S)$ .

THEOREM 2. Let A be a basic, finite-dimensional k-algebra which is quasi-hereditary with labelled species  $\mathcal S$ . Then, for any i,j

$$\dim_{k}(e_{i}S(\mathcal{S})e_{j}) \leq \dim_{k}(e_{i}Ae_{j}) \leq \dim_{k}(e_{i}D(\mathcal{S})e_{j}).$$

In particular,

$$s_k(\mathcal{S}) \leqslant \dim_k A \leqslant d_k(\mathcal{S}).$$

We have  $s_k(\mathcal{S}) = \dim_k A$  if and only if A is shallow, and  $d_k(\mathcal{S}) = \dim_k A$  if and only if A is deep.

The proof of Theorem 1 is given in Section 2, the proof of Theorem 2 in Section 3. We add examples showing that besides the algebras  $S(\mathcal{S})$  and  $D(\mathcal{S})$ , there are other shallow or deep algebras. A detailed study of the ring-theoretical and homological properties of quasi-hereditary rings which are shallow or deep will be given in a subsequent publication.

# 2. The rings $S(\mathcal{S})$ and $D(\mathcal{S})$

The aim of this section is a proof of Theorem 1. Thus, let  $\mathscr S$  be a labelled species without loops, with index set  $\{1, \ldots, n\}$ . The proof is by induction on n. If n = 1, then  $S(\mathscr S) = D(\mathscr S) = F_1$ , thus quasi-hereditary (and trivially both shallow and deep). Thus, let  $n \ge 2$ , and let  $\mathscr S'$  be the restriction of  $\mathscr S$  to  $\{1, \ldots, n-1\}$ .

Consider first  $S(\mathcal{S})$ . Given  $m \in \mathbb{N}$ , let  $[1, m] = \{i \in \mathbb{N} \mid 1 \le i \le m\}$ . Then

$$S(\mathcal{S})e_{n} = F_{n} \oplus M([1, n-1] \cdot n),$$

$$e_{n}S(\mathcal{S}) = F_{n} \oplus M(n \cdot [1, n-1]),$$

$$\langle e_{n} \rangle = F_{n} \oplus M(\{t \in U \mid t_{i} = n \text{ for some } i\})$$

$$= F_{n} \oplus M([1, n-1] \cdot n \cup n \cdot [1, n-1] \cup [1, n-1] \cdot n \cdot [1, n-1])$$

$$= (F_{n} \oplus M([1, n-1] \cdot n)) \otimes_{F_{n}} (F_{n} \oplus M(n \cdot [1, n-1]))$$

$$= S(\mathcal{S})e_{n} \otimes_{F_{n}} e_{n}S(\mathcal{S}).$$

In particular,  $e_n S(\mathcal{S})e_n = F_n$ , and the equalities above show that  $\langle e_n \rangle$  is a heredity ideal. Of course, rad  $\Delta(n) = M(n \cdot [1, n-1])$  is a semisimple right

module,  $\operatorname{rad} \Delta^*(n) = M([1, n-1] \cdot n)$  is a semisimple left module. Since  $S(\mathcal{S})/\langle e_n \rangle = S(\mathcal{S}')$ , we use induction and conclude that  $S(\mathcal{S})$  is a shallow quasi-hereditary ring.

Next, we consider  $D(\mathcal{S})$ . We have

$$\begin{split} D(\mathcal{S})e_n &= F_n \oplus M\big(V(n-1)\cdot n\big), \\ e_n D(\mathcal{S}) &= F_n \oplus M\big(n\cdot V(n-1)\big), \\ \langle e_n \rangle &= F_n \oplus M\big(V(n-1)\cdot n \cup n\cdot V(n-1) \cup V(n-1)\cdot n\cdot V(n-1)\big) \\ &= \big(F_n \oplus M\big(V(n-1)\cdot n\big)\big) \otimes_{F_n} \big(F_n \oplus M\big(n\cdot V(n-1)\big)\big) \\ &= D(\mathcal{S})e_n \otimes_{F_n} e_n D(\mathcal{S}), \end{split}$$

so that  $e_n D(\mathcal{S}) e_n = F_n$ , and  $\langle e_n \rangle$  is a heredity ideal. Since  $D(\mathcal{S})/\langle e_n \rangle = D(\mathcal{S}')$ , it follows by induction that  $D(\mathcal{S})$  is quasi-hereditary. Now

$$\operatorname{rad} \Delta(n) = M(n \cdot V(n-1)) = \bigoplus_{i=1}^{n-1} {}_{n}M_{i} \otimes_{F_{i}} P_{D(\mathscr{S}')}(i),$$

thus  $\Delta(n)$  is a projective right  $D(\mathcal{S}')$ -module. Similarly, rad  $\Delta^*(n)$  is a projective left  $D(\mathcal{S}')$ -module. By induction, it follows that  $D(\mathcal{S})$  is deep.

## 3. Quasi-hereditary k-algebras

Let k be a field, and A a basic finite-dimensional quasi-hereditary k-algebra with labelled species  $\mathscr{G}$ . Let  $\{1, \ldots, n\}$  be the index set of  $\mathscr{G}$ . Note that  $e_n A e_n = F_n$ , and, in the same way,  $e_n S(\mathscr{G}) e_n = e_n D(\mathscr{G}) e_n = F_n$ . In particular, for the proof of the dimension inequalities, we may assume  $n \ge 2$ . Let  $\mathscr{G}'$  be the restriction of  $\mathscr{G}$  to  $\{1, \ldots, n-1\}$ ; clearly, this is the labelled species for  $B = A/\langle e_n \rangle$ . By induction, we know that

$$\dim_{k}(e_{i}S(\mathcal{S}')e_{j}) \leq \dim_{k}(e_{i}Be_{j}) \leq \dim_{k}(e_{i}D(\mathcal{S}')e_{j}),$$

for all  $i, j \leq n-1$ .

First, consider  $e_n A e_j$ , with  $1 \le j \le n-1$ . Let  $X = \bigoplus_{j=1}^{n-1} e_n A e_j$ , thus X is the radical of the right A-module  $e_n A$ ; this is a B-module with top  $\overline{X} = \bigoplus_{i=1}^{n-1} {}_n M_i$ . Let  $d_i = \dim({}_n M_i)_{F_i}$ . We denote by P the B-projective cover of X, thus P is the direct sum of  $d_i$  copies of  $e_i B$ , for  $1 \le i \le n-1$ . The epimorphisms  $P \to X \to \overline{X}$  yield epimorphisms  $P e_j \to X e_j \to \overline{X} e_j$ . Now,  $\overline{X} e_j = {}_n M_j$ ,  $X e_j = e_n A e_j$ , and  $P e_j = \bigoplus_{i=1}^{n-1} (e_i B e_j)^{d_i}$ , thus

$$\dim_k({}_nM_j)\leqslant\dim_k(e_nAe_j)\leqslant\sum_{i=1}^{n-1}d_i\cdot\dim_k(e_iBe_j).$$

However,  $e_n S(\mathcal{S}) e_j = {}_n M_j$ , so the left-hand term is the desired one. Now,

 $rad(e_n D(\mathcal{S})_{D(\mathcal{S})})$  is the  $D(\mathcal{S}')$ -projective module with top  $\bigoplus_{i=1}^{n-1} {}_n M_i$ , thus

$$\mathrm{rad}(e_n D(\mathscr{S})_{D(\mathscr{S})}) = \bigoplus_{i=1}^{n-1} (e_i D(\mathscr{S}'))^{d_i}.$$

It follows that  $e_n D(\mathcal{S}) e_j = \bigoplus_{i=1}^{n-1} (e_i D(\mathcal{S}') e_j)^{d_i}$ , and therefore

$$\sum_{i=1}^{n-1} d_i \cdot \dim_k(e_i B e_j) \leqslant \sum_{i=1}^{n-1} d_i \cdot \dim_k(e_i D(\mathcal{S}') e_j) = \dim_k(e_n D(\mathcal{S}) e_j).$$

This finishes the proof for  $e_n A e_j$ . The dual proof yields the similar inequality for  $e_j A e_n$ , where  $1 \le j \le n-1$ .

It remains to consider  $e_i A e_j$ , where  $1 \le i, j \le n-1$ . Since  $\langle e_n \rangle$  $=Ae_n\otimes_{F_n}e_nA$ , there is the exact sequence

$$0 \rightarrow e_i A e_n \otimes_{F_n} e_n A e_j \rightarrow e_i A e_j \rightarrow e_i B e_j \rightarrow 0,$$

and similar ones for  $S(\mathcal{S})$  and  $D(\mathcal{S})$ , namely

$$\begin{split} 0 &\to e_i S(\mathcal{S}) e_{\mathbf{n}} \otimes_{F_{\mathbf{n}}} e_{\mathbf{n}} S(\mathcal{S}) e_j \to e_i S(\mathcal{S}) e_j \to e_i S(\mathcal{S}') e_j \to 0, \\ 0 &\to e_i D(\mathcal{S}) e_{\mathbf{n}} \otimes_{F_{\mathbf{n}}} e_{\mathbf{n}} D(\mathcal{S}) e_j \to e_i D(\mathcal{S}) e_j \to e_i D(\mathcal{S}') e_j \to 0. \end{split}$$

The desired inequalities follow from the inequalities for  $e_i A e_n$ ,  $e_n A e_j$ , and  $e_i B e_j$ , by taking into account that for a right  $F_n$ -space X and a left  $F_n$ -space Y, we have

$$\dim_k X \otimes_{F_n} Y = \frac{1}{\dim_k F_n} \dim_k X \cdot \dim_k Y.$$

This finishes the proof of the first part of Theorem 2.

Now assume that A is shallow. By induction, we know that  $\dim_k(e_iS(\mathcal{S}')e_j) = \dim_k(e_iBe_j)$ , for  $i,j \leq n-1$ . Since  $X = \overline{X}$ , we have  $e_n S(\mathcal{S}) e_j = {}_n M_j = e_n A e_j$ , for  $j \le n-1$ , and similarly  $e_j S(\mathcal{S}) e_n = e_j A e_n$  for  $j \le n-1$ . It follows that  $\dim_k(e_i S(\mathcal{S})e_j) = \dim_k(e_i Ae_j)$ , for all i, j.

Similarly, if we assume that A is deep, then, by induction,  $\dim_k(e_iBe_j) = \dim_k(e_iD(\mathcal{S}')e_j)$ , for  $i, j \le n-1$ . On the other hand, we have in this case X = P, thus  $e_n A e_j' = \bigoplus_{i=1}^{n-1} (e_i B e_j)^{d_i}$ , and therefore

$$\dim_{\mathbf{k}}(e_nAe_j) = \sum_{i=1}^{n-1} d_i \cdot \dim_{\mathbf{k}}(e_iBe_j) = \sum_{i=1}^{n-1} d_i \cdot \dim_{\mathbf{k}}(e_iD(\mathscr{S}')e_j) = \dim_{\mathbf{k}}(e_nD(\mathscr{S})e_j).$$

It follows that  $\dim_k(e_iAe_j) = \dim_k(e_iD(\mathcal{S})e_j)$ . Note that  $\dim_k A = \sum_{i,j} \dim_k(e_iAe_j)$ , thus always  $s_k(\mathcal{S}) \leq \dim_k A \leq d_k(\mathcal{S})$ . Let us first assme  $s_k(\mathscr{S}) = \dim_k A$ , thus  $\dim_k (e_i A e_j) = \dim_k (e_i S(\mathscr{S}) e_j)$ , for all i,j. If  $i,j \leq n-1$ , a proper inequality  $\dim_k(e_iS(\mathcal{S}')e_j) < \dim_k(e_iBe_j)$  would yield that  $\dim_k(e_iS(\mathcal{S})e_j) < \dim_k(e_iAe_j)$  for the same pair i, j of indices, since

$$\dim_{k}(e_{i}Ae_{j})-\dim_{k}(e_{i}S(\mathcal{S})e_{i})=\dim_{k}(e_{i}Be_{j})-\dim_{k}(e_{i}S(\mathcal{S}')e_{j})+a,$$

with

$$a = \dim_{\mathbf{k}}(e_i A e_n \otimes_{F_n} e_n A e_j) - \dim_{\mathbf{k}}(e_i S(\mathcal{S}) e_n \otimes_{F_n} e_n S(\mathcal{S}) e_j) \geqslant 0.$$

Thus  $s_k(\mathcal{S}') = \dim_k B$ , and B is shallow by induction. On the other hand,  $\dim_k(e_n S(\mathcal{S})e_j) = \dim_k(e_n Ae_j)$  implies that  $Xe_j = \overline{X}e_j$ , for all  $1 \le j < n$ , and therefore  $X = \overline{X}$  is semisimple. This shows that the right A-module  $e_n A$  has Loewy length at most 2. Similarly, the left A-module  $Ae_n$  has Loewy length at most 2. As a consequence, A is shallow.

In the same way, we proceed in case  $\dim_k A = d_k(\mathcal{S})$ . We see immediately that  $\dim_k(e_iAe_j) = \dim_k(e_iD(\mathcal{S})e_j)$ , for all i,j, and conclude that  $\dim_k B = d_k(\mathcal{S}')$ . Thus B is deep by induction. On the other hand,  $\dim_k(e_nAe_j) = \dim_k(e_nD(\mathcal{S})e_j)$  implies that  $Pe_j = Xe_j$ , for all  $1 \le j \le n-1$ , and therefore X = P is a projective right B-module. Similarly, the radical of the left A-module  $Ae_n$  is projective as a left B-module. Thus A is deep.

#### 4. Examples

The bounds  $s_k(\mathcal{S}) \leq \dim_k A \leq d_k(\mathcal{S})$  are optimal, but we should remark that usually  $d_k(\mathcal{S}) - s_k(\mathcal{S})$  may be rather large. As an example, consider the k-species  $\mathcal{S}_n = (F_i, {}_iM_j)_{1 \leq i,j \leq n}$  with  $F_i = k$  and  ${}_iM_i = 0$  for all i, whereas  ${}_iM_j = k$  for all  $i \neq j$ ; thus  $T(\mathcal{S}_n)$  is the path algebra for the quiver with n vertices, a unique arrow  $i \to j$  for  $i \neq j$ , and no loops. We are going to exhibit  $s(n) := s_k(\mathcal{S}_n)$  and  $d(n) := d_k(\mathcal{S}_n)$ . It suffices to calculate the cardinalities of the index sets U(n) and V(n), since

$$s(n) = n + U(n), \quad d(n) = n + V(n).$$

Clearly, |U(1)| = 0 = |V(1)|. For  $n \ge 2$ , we have

$$U(n) = U(n-1) \cup [1, n-1] \cdot n \cup n \cdot [1, n-1] \cup [1, n-1] \cdot n \cdot [1, n-1],$$

thus

$$|U(n)| = |U(n-1)| + 2(n-1) + (n-1)^2 = |U(n-1)| + n^2 - 1,$$

and consequently,

$$|U(n)| = -n + \sum_{t=1}^{n} t^2 = -n + \frac{1}{6}n(n+1)(2n+1).$$

Similarly, from

$$V(n) = V(n-1) \cup V(n-1) \cdot n \cup n \cdot V(n-1) \cup V(n-1) \cdot n \cdot V(n-1)$$

for  $n \ge 2$ , we obtain

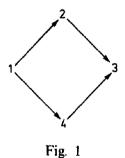
$$|V(n)| = 3|V(n-1)| + |V(n-1)|^2.$$

It follows that  $s(n) = \frac{1}{6}(n+1)(2n+1)$ , and that d(n) is given recursively by

d(1) = 1, and  $d(n) = d(n-1) + (d(n-1)+1)^2$  for  $n \ge 2$ . The first values for s(n) and d(n) are the following:

$$s(1) = 1,$$
  $d(1) = 1,$   
 $s(2) = 5,$   $d(2) = 5,$   
 $s(3) = 14,$   $d(3) = 41,$   
 $s(4) = 30,$   $d(4) = 1805,$   
 $s(5) = 55,$   $d(5) = 3263441.$ 

Let  $\mathscr G$  be a labelled species without loops. Let us assume that there are even no oriented cycles. Then  $D(\mathscr G)$  is the tensor algebra of  $\mathscr G$ . In particular, if  $\mathscr G$  is, in addition, a finite-dimensional k-algebra where k is a perfect field, then  $D(\mathscr G)$  is the only deep quasi-hereditary algebra with species  $\mathscr G$ . If the labelling is chosen in such a way that  ${}_iM_j=0$  for i>j, then  $S(\mathscr G)=T(\mathscr G)/T_+(\mathscr G)^2$ , so again  $S(\mathscr G)$  is the only shallow quasi-hereditary algebra with labelled species  $\mathscr G$ . Of course, in general there may be shallow rings which are not of the form  $S(\mathscr G)$ , the first example is the path algebra of the quiver of Fig. 1 with the commutativity relation.



For a labelled species  $\mathscr S$  without loops but with oriented cycles there usually also will exist deep rings which are not of the form  $D(\mathscr S)$ . For example, consider the algebra A given by the quiver of Fig. 2 with relations  $\beta \alpha - \gamma \delta = 0$ 

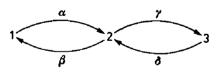


Fig. 2

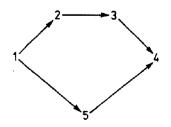


Fig. 3

and  $\delta \gamma = 0$ . The labelled species corresponding to this quiver will be denoted by  $\mathscr{S}$ . Then A is deep with labelled species  $\mathscr{S}$ , but not isomorphic to  $D(\mathscr{S})$ .

Also, we should remark that there are quasi-hereditary algebras A with radical N such that no ideal  $I \subseteq N^2$  yields a shallow algebra A/I. A typical example is the algebra A given by the quiver of Fig. 3 with the commutativity relation. Note that A has a unique minimal nonzero ideal I. An ideal I with A/I shallow must contain I, but there is no ideal I with  $I \subseteq I \subseteq I$  such that I is quasi-hereditary with respect to the given ordering of the vertices.

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