# Major Index for 01-Fillings of Moon Polyominoes

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#### Abstract

We propose a major index statistic on 01-fillings of moon polyominoes which, when specialized to certain shapes, reduces to the major index for permutations and set partitions. We consider the set  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  of all 01-fillings of a moon polyomino  $\mathcal{M}$  with given column sum  $\mathbf{s}$  whose empty rows are A, and prove that this major index has the same distribution as the number of north-east chains, which are the natural extension of inversions (resp. crossings) for permutations (resp. set partitions). Hence our result generalizes the classical equidistribution results for the permutation statistics inv and maj. Two proofs are presented. The first is an algebraic one using generating functions, and the second is a bijection on 01-fillings of moon polyominoes in the spirit of Foata's second fundamental transformation on words and permutations.

### 1 Introduction

Given a multiset S of n positive integers, a word on S is a sequence  $w = w_1 w_2 \dots w_n$  that reorders the elements in S. When  $S = [n] := \{1, \dots, n\}$  the word is a permutation. A pair  $(w_i, w_j)$  is called an *inversion* of w if i < j and  $w_i > w_j$ . One well-known statistic on words and permutations is inv(w), defined as the number of inversions of w. The *descent set* and *descent statistic* of a word w are defined as

$$Des(w) = \{i : 1 \le i \le n - 1, w_i > w_{i+1}\}, \qquad des(w) = \#Des(w).$$

In [15] MacMahon defined the major index statistic for a word w as

$$\operatorname{maj}(w) = \sum_{i \in \operatorname{Des}(w)} i,$$

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and showed the remarkable result that its distribution over all words on S is equal to that of the inversion number over the same set. Precisely, for the set  $W_S$  of all words on S,

$$\sum_{w \in W_S} q^{\operatorname{maj}(w)} = \sum_{w \in W_S} q^{\operatorname{inv}(w)}.$$
(1.1)

The proof of (1.1) given by MacMahon relied on combinatorial analysis, while an elegant bijection  $\Phi: W_S \to W_S$  such that  $\operatorname{maj}(w) = \operatorname{inv}(\Phi(w))$  was constructed later by Foata [6].

A set partition of [n] is a family of nonempty sets  $B_1, \ldots, B_k$  which are pairwise disjoint and whose union is [n]. If all the blocks  $B_i$  have at most two elements, the set partition is called a matching. If the size of every block is exactly two, the matching is called a *perfect matching*. A set partition  $\pi$  can be represented as a simple graph with vertex set [n] drawn on a horizontal line, where two vertices are joined by an arc if and only if they represent consecutive elements in the same block of  $\pi$  under the numerical order. Two statistics, the numbers of crossings and nestings, arise naturally from such graphical representations, and have been the central topic in many recent research articles, e.g., [1], [12], [13], [17], [20], [16], [18], [5], [22], to list a few. A crossing (resp. nesting) of a set partition is a pair of arcs  $(i_1, j_1), (i_2, j_2)$  such that  $i_1 < i_2 < j_1 < j_2$ (resp.  $i_1 < i_2 < j_2 < j_1$ ). As explained in [2, Section 2], a crossing can be viewed as a natural generalization of an inversion in a permutation. Hence it is natural to ask what would be the major index for matchings (set partitions) that extends MacMahon's equidistribution results (1.1). A definition of such a statistic, denoted by pmaj, was proposed by Chen et al. in [2]. For completeness, we present the definition for perfect matchings here. The general case for set partitions is very similar. Label the arcs of a perfect matching M of [2n] by  $1, \ldots, n$  in decreasing order with respect to their left endpoints from left to right. Suppose  $r_1, \ldots, r_n$  are the right endpoints of arcs from right to left. Then to each right endpoint  $r_i$  of an arc associate a word  $w^{(i)}$  on [n] where the words  $w^{(i)}$  are defined backwards recursively: let  $w^{(1)} = a$ , where a is the label of the arc that ends with  $r_1$ . In general, after defining  $w^{(i)}$ , assume that the left endpoints of the arcs labeled  $a_1, \ldots, a_t$  lie between  $r_i$  and  $r_{i+1}$ . Then  $w^{(i+1)}$  is obtained from  $w^{(i)}$  by deleting entries  $a_1, \ldots, a_t$  and adding the entry b at the very beginning, where b is the label of the arc that ends with  $r_{i+1}$ . Then

$$pmaj(M) := \sum_{i=1}^{n} des(w^{(i)})$$

Chen et al. [2] computed the generating function for *pmaj* and showed that it is equally distributed with the number of crossings over all matchings with fixed left and right endpoints. A similar result holds for set partitions.

The aim of this paper is to extend the major index to certain 01-fillings of moon polyominoes which include words and set partitions. In Section 2 we introduce the necessary notations, and describe the definition of the major index for 01-fillings of moon polyominoes. We explain in Section 3 how the classical definition of major index on words and set partitions can be obtained by considering moon polyominoes of special shapes. In Section 4 we show that the maj statistic is equally distributed as ne, the number of north-east chains, by computing the corresponding generating functions. In the fillings of special shapes, ne corresponds to the number of inversions for words and crossings for set partitions. Therefore, our main result, Theorem 4.1, leads to a generalization of (1.1) and the analogous result for set partitions. In Section 5, we present a bijective proof for the equidistribution of maj and ne, which consists of three maps. The first is from the fillings of left-aligned stack polyominoes to itself that sends maj to ne, constructed in the spirit of Foata's second fundamental transformation. The other two are maps that transform a moon polyomino to a left-aligned stack polyomino with the same set of columns, while preserving the statistics maj and ne, respectively. Composing these three maps yields the desired bijection.

### 2 Definition of major index for fillings of moon polyominoes

A polyomino is a finite subset of  $\mathbb{Z}^2$ , where we represent every element of  $\mathbb{Z}^2$  by a square cell. The polyomino is *convex* if its intersection with any column or row is connected. It is *intersection-free* if every two columns are comparable, i.e., the row-coordinates of one column form a subset of those of the other column. Equivalently, it is intersection-free if every two rows are comparable. A moon polyomino is a convex intersection-free polyomino. If the rows (resp. columns) of the moon polyomino are left-aligned (resp. top-aligned), we will call it a *left-aligned stack polyomino* (resp. *top-aligned stack polyomino*). A *Ferrers diagram* is a left-aligned and top-aligned stack polyomino. See Figure 1 for an illustration. The term 'moon polyomino' was first used by Jonsson in [10] where he used fillings of such polyominoes to study generalized triangulations.



Figure 1: A moon polyomino, a left-aligned and a top-aligned stack polyomino, and a Ferrers diagram.

We are concerned with 01-fillings of moon polyominoes with restricted row sums. That is, given a moon polyomino  $\mathcal{M}$ , we assign a 0 or a 1 to each cell of  $\mathcal{M}$  so that there is at most one 1 in each row. Throughout this paper we will simply use the term *filling* to denote such 01-filling. We say a cell is empty if it is assigned a 0, and it is a 1-cell otherwise. Given a moon polyomino  $\mathcal{M}$ with m columns, we label them  $c_1, \ldots, c_m$  from left to right. Let  $\mathbf{s} = (s_1, \ldots, s_m)$  be an m-tuple of nonnegative integers and A be a subset of rows of  $\mathcal{M}$ . We denote by  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  the set of fillings M of  $\mathcal{M}$  such that the empty rows of M are exactly those in A and the column  $c_i$  has exactly  $s_i$ many 1's,  $1 \leq i \leq m$ . A north-east (NE) chain in a filling M of  $\mathcal{M}$  is a set of two 1-cells such that one of them is strictly above and to the right of the other and the smallest rectangle containing them is contained in  $\mathcal{M}$ . The number of NE chains of M will be denoted ne(M). See Figure 2.

The set  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  was first studied by Kasraoui [11], who showed that the numbers of northeast and south-east chains are equally distributed. The generating functions for these statistics are also presented in [11, Theorem 4.4].

To define the major index for fillings of moon polyominoes, we first state it for rectangular shapes, which is essentially the classical definition of major index for words, (c.f. Section 3.1).

Let R be a filling of a rectangle whose n nonempty rows  $r_1, \ldots, r_n$  are numbered from top to bottom. We define the *descent statistic* for R as

 $des(R) = |\{i \mid \text{ the 1-cells in rows } r_i \text{ and } r_{i+1} \text{ form an NE chain}\}|.$ 



Figure 2: Filling M of a moon polyomino  $\mathcal{M}$  for  $A = \{3\}$  and  $\mathbf{s} = (1, 1, 2, 1, 1, 0)$  with  $\operatorname{ne}(M) = 5$ .

That is, descents of a rectangular filling are NE chains in consecutive nonempty rows.

For each nonempty row  $r_i$  of R, let  $R(r_i)$  denote the rectangle that contains the row  $r_i$  and all rows in R above the row  $r_i$ .

**Definition 2.1.** The major index for the rectangular filling R is defined to be

$$\operatorname{maj}(R) = \sum_{i=1}^{n} \operatorname{des}(R(r_i)).$$
(2.1)

Clearly, the empty rows of R do not play any role in maj(R). That is, if we delete them, the major index of the resulting filling remains the same.

**Definition 2.2.** Let M be a filling of a moon polyomino  $\mathcal{M}$ . Let  $\mathcal{R}_1, \ldots, \mathcal{R}_r$  be the list of all the maximal rectangles contained in  $\mathcal{M}$  ordered increasingly by height, and denote by  $R_i$  the filling M restricted on the rectangle  $\mathcal{R}_i$ . Then the major index of M is defined to be

$$\operatorname{maj}(M) = \sum_{i=1}^{r} \operatorname{maj}(R_i) - \sum_{i=1}^{r-1} \operatorname{maj}(R_i \cap R_{i+1}),$$
(2.2)

where  $\operatorname{maj}(R_i)$  and  $\operatorname{maj}(R_i \cap R_{i+1})$  are defined by (2.1).

In particular,  $\operatorname{maj}(M) = \operatorname{maj}(R_1)$  when M is a rectangular shape. It is also clear that  $\operatorname{maj}(M)$  is always nonnegative since  $\operatorname{maj}(R_{i+1}) \ge \operatorname{maj}(R_i \cap R_{i+1}), i = 1, \ldots, r-1$ .

**Example 2.3.** Consider the filling from Figure 2. It has four maximal rectangles and maj(M) = (2+2+1+1) - (2+0+0) = 4.



Figure 3: Calculation of maj(M) for a moon polyomino using Definition 2.2.

The major index can be equivalently defined in a slightly more complicated way. However, this way is useful in proofs, especially when one uses induction on the number of columns of the filling, (c.f. Theorem 4.1). We state this equivalent definition next. First we need some notation. Let  $Left(\mathcal{M})$  be the set of columns of  $\mathcal{M}$  which are to left of the columns of maximal length and let  $Right(\mathcal{M})$  consist of the remaining columns of  $\mathcal{M}$ .

We order the columns of  $\mathcal{M}, c_1, \ldots, c_m$ , by a total order  $\prec$  as follows:  $c_i \prec c_j$  if and only if

- $|c_i| < |c_j|$  or
- $|c_i| = |c_j|, c_i \in Left(\mathcal{M}) \text{ and } c_j \in Right(\mathcal{M}), \text{ or}$
- $|c_i| = |c_j|, c_i, c_j \in Left(\mathcal{M})$  and  $c_i$  is to the left of  $c_j$ , or
- $|c_i| = |c_j|, c_i, c_j \in Right(\mathcal{M})$  and  $c_i$  is to the right of  $c_j$ ,

where |c| denotes the length of the column c. A similar ordering of rows was used in [11].

For every column  $c_i \in Left(\mathcal{M})$  define the rectangle  $\mathcal{M}(c_i)$  to be the largest rectangle that contains  $c_i$  as the leftmost column. For  $c_i \in Right(\mathcal{M})$ , the rectangle  $\mathcal{M}(c_i)$  is taken to be the largest rectangle that contains  $c_i$  as the rightmost column and does not contain any columns from  $Left(\mathcal{M})$  of same length as  $c_i$ .

**Definition 2.2'.** Let  $c_{i_1} \prec c_{i_2} \prec \cdots \prec c_{i_m}$  be the ordering of the columns of  $\mathcal{M}$  and let  $M_j$  be the restriction of M on the rectangle  $\mathcal{M}(c_{i_j}), 1 \leq j \leq m$ . Then

$$\operatorname{maj}(M) = \sum_{j=1}^{m} \operatorname{maj}(M_j) - \sum_{j=1}^{m-1} \operatorname{maj}(M_j \cap M_{j+1}),$$
(2.3)

where  $\operatorname{maj}(M_j)$  and  $\operatorname{maj}(M_j \cap M_{j+1})$  are defined by (2.1).

**Example 2.4.** Consider the filling M from Figure 2. The order  $\prec$  on the columns of  $\mathcal{M}$  is  $c_6 \prec c_1 \prec c_5 \prec c_4 \prec c_3 \prec c_2$ . So,  $\mathcal{M}_1 = \mathcal{M}(c_6)$ ,  $\mathcal{M}_2 = \mathcal{M}(c_1)$ ,  $\mathcal{M}_3 = \mathcal{M}(c_5)$ ,  $\mathcal{M}_4 = \mathcal{M}(c_4)$ ,  $\mathcal{M}_5 = \mathcal{M}(c_3)$ , and  $\mathcal{M}_6 = \mathcal{M}(c_2)$ , as illustrated in Figure 4. By Definition 2.2', maj $(\mathcal{M}) = (2+2+1+1+1+0) - (2+1+0+0+0) = 4$ .

**Proposition 2.5.** Definitions 2.2 and 2.2' are equivalent.

Proof. Some of the rectangular fillings  $M_j$  in Definition 2.2' may contain  $M_{j+1}$ , in which case  $\operatorname{maj}(M_{j+1}) - \operatorname{maj}(M_j \cap M_{j+1}) = 0$  and formula (2.3) can be simplified. More precisely, there are uniquely determined indices  $1 = j_1 < j_2 < \cdots < j_r \leq m$  such that  $M_1 = M_{j_1} \supseteq M_2 \supseteq \cdots \supseteq M_{j_2-1} \not\supseteq M_{j_2} \supseteq \cdots \supseteq M_{j_r-1} \not\supseteq M_{j_r} \supseteq \cdots \supseteq M_m$ . That is, if the rectangles  $M_i$  are ordered by containment, then  $M_{j_1}, M_{j_2}, \ldots, M_{j_r}$  are the maximal elements. Following the notation in Definition 2.2, the filling  $M_{j_k}$  is the filling  $R_k$  of the k-th maximal rectangle contained in  $\mathcal{M}$ . Then, after cancellation of some terms, the right-hand side of (2.3) becomes the right-hand side of (2.2). For instance, in the previous example, the rectangles  $M_3$  and  $M_6$  are not maximal and therefore  $\operatorname{maj}(M_3) - \operatorname{maj}(M_2 \cap M_3) = 0$  and  $\operatorname{maj}(M_6) - \operatorname{maj}(M_5 \cap M_6) = 0$ .



Figure 4: Calculation of maj(M) for a moon polyomino using Definition 2.2'.

## 3 maj(M) for special shapes $\mathcal{M}$

It is well-known that permutations and set-partitions can be represented as fillings of Ferrers diagrams (e.g. [14]). Here we will describe such presentations to show how to get the classical major index for words and set partitions from Definition 2.2.

### 3.1 When $\mathcal{M}$ is a rectangle: words and permutations

For any rectangle, label the rows from top to bottom, and columns from left to right. Fillings of rectangles are in bijection with words. More precisely, let  $w = w_1 \dots w_n$  be a word with letters in the set [m]. The word w can be represented as a filling M of an  $n \times m$  rectangle  $\mathcal{M}$  in which the cell in row n - i + 1 and column m - j + 1 is assigned the integer 1 if  $w_i = j$ , and is empty otherwise. Conversely, each filling of the rectangle  $\mathcal{M}$  corresponds to a word w. It is easy to check that  $\operatorname{ne}(M) = \operatorname{inv}(w)$ ,  $\operatorname{des}(M) = \operatorname{des}(w)$ , and  $\operatorname{maj}(M) = \operatorname{maj}(w)$ . The latter follows from the fact that  $\operatorname{maj}(w) = \sum_{i=1}^{n} \operatorname{des}(w_i w_{i+1} \dots w_n)$ .

#### 3.2 When $\mathcal{M}$ is a Ferrers diagram: matchings and set partitions

As explained in [4], general fillings of Ferrers diagrams correspond to multigraphs. Here we briefly describe the correspondence when restricted to 01-fillings with row sum at most 1. The Ferrers diagram is bounded by a vertical line from the left, a horizontal line from above, and a path consisting of east and north steps. Following this path starting from the bottom left end, we label the steps of the path by  $1, 2, \ldots, n$ . This gives a labeling of the columns and rows of the diagram. Draw n vertices on a horizontal line and draw an edge connecting vertices i and j if and only if there is a 1 in the cell of the column labeled i and the row labeled j (see Figure 5). The resulting graph has no loops or multiple edges and every vertex is a right endpoint of at most one edge. The NE chains correspond to crossings of edges in the graph. If the column sums are at most one,

the graph is a matching. If, moreover, in the labeling of the steps we allow to have a vertical step followed by a horizontal one with a same label, then the corresponding graph will have vertices which are both right and left endpoints of edges. Such graphs represent set partitions. In either case, the major index of the filling is equal to the major index of the corresponding matching or set partition as defined in [2]. Restricting to the triangular Ferrers diagram, the 01-fillings considered in this paper also include *linked partitions*, a combinatorial structure that was studied in [3, 11].



Figure 5: A filling of a Ferrers diagram and the corresponding graph.

#### 3.3 When $\mathcal{M}$ is a top-aligned stack polyomino

In the case when  $\mathcal{M}$  is a top-aligned stack polyomino, maj(M) has a simpler form in terms of the des statistic, just as in words and permutations. Explicitly, if the nonempty rows of M are  $r_1, \ldots, r_n$ , denote by  $M(r_i)$  the filling M restricted to the largest rectangle contained in  $\mathcal{M}$  whose bottom row is  $r_i$ .

**Proposition 3.1.** Let M be a filling of a top-aligned stack polyomino and  $M(r_i)$  be as defined above. Then

$$maj(M) = \sum_{i=1}^{n} des(M(r_i)).$$
 (3.1)

*Proof.* Let  $r_{k+1}, \ldots, r_n$  be all the nonempty rows of M that are of same length as the last row  $r_n$ . Denote by  $M' = M \setminus \{r_{k+1}, \ldots, r_n\}$  and by R the filling of the largest rectangle of  $\mathcal{M}$  containing  $r_n$ . We proceed by induction on the number n of rows of M. The claim is trivial when M has only one row or is a rectangle. Otherwise,

$$maj(M) = maj(M') + maj(R) - maj(M' \cap R)$$
(Definition 2.2)  
$$= \sum_{i=1}^{k} des(M(r_i)) + \sum_{i=k+1}^{n} des(M(r_i)),$$
(Induction hypothesis and Definition 2.1)

which completes the proof.

**Example 3.2.** Consider the filling M of a top-aligned stack polynomia in Figure 6 with  $A = \{3\}$  and  $\mathbf{s} = (0, 1, 2, 1, 1)$ . M has five nonempty rows and, by Proposition 3.1,  $\operatorname{maj}(M) = \sum_{i=1}^{5} \operatorname{des}(M(r_i)) = 0 + 1 + 0 + 1 + 1 = 3$ .

Proposition 3.1 will be used in the proof of Lemma 5.5.



Figure 6: Calculation of maj(M) using Proposition 3.1.

### 4 The generating function for the major index

In this section we state and prove the main theorem which gives the generating function for maj over  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$ , the set of all fillings of the moon polyomino  $\mathcal{M}$  with column sums given by the integer sequence  $\mathbf{s}$  and empty rows given by A. Let  $|c_1|, \ldots, |c_m|$  be the column lengths of M and let  $a_i$  the number of rows in A that intersect the column  $c_i$ . Suppose that, under the ordering defined in Section 2,  $c_{i_1} \prec c_{i_2} \prec \cdots \prec c_{i_m}$ . Then for  $j = 1, \ldots, m$ , define

$$h_{i_j} = |c_{i_j}| - a_{i_j} - (s_{i_1} + s_{i_2} + \dots + s_{i_{j-1}})$$

$$(4.1)$$

The numbers  $h_i$  have the following meaning: if one fills in the columns of M from smallest to largest according to the order  $\prec$ , then  $h_{i_j}$  is the number of available cells in the *j*-th column to be filled.

**Theorem 4.1.** For a moon polyomino  $\mathcal{M}$  with m columns,

$$\sum_{M \in \boldsymbol{F}(\mathcal{M}, \boldsymbol{s}; A)} q^{\operatorname{maj}(M)} = \prod_{i=1}^{m} \begin{bmatrix} h_i \\ s_i \end{bmatrix}_q$$
(4.2)

where  $h_i$  is defined by (4.1).

We postpone the proof of Theorem 4.1 until the end of this section.

**Example 4.2.** Suppose  $\mathcal{M}$  is the first moon polyomino in Figure 1,  $A = \{5\}$ , and  $\mathbf{s} = (1, 0, 2, 1, 1)$ . The  $\prec$  order on the columns of  $\mathcal{M}$  is:  $c_1 \prec c_5 \prec c_2 \prec c_4 \prec c_3$ . The fifth row intersects all columns except  $c_1$ , so,  $a_1 = 0$  and  $a_2 = a_3 = a_4 = a_5 = 1$ . Therefore,

$$\begin{split} h_1 &= h_{i_1} = |c_1| - a_1 = 2 - 0 = 2 \\ h_5 &= h_{i_2} = |c_5| - a_5 - s_1 = 3 - 1 - 1 = 1 \\ h_2 &= h_{i_3} = |c_2| - a_2 - (s_1 + s_5) = 4 - 1 - (1 + 1) = 1 \\ h_4 &= h_{i_4} = |c_4| - a_4 - (s_1 + s_5 + s_2) = 6 - 1 - (1 + 1 + 0) = 3 \\ h_3 &= h_{i_5} = |c_3| - a_3 - (s_1 + s_5 + s_2 + s_4) = 6 - 1 - (1 + 1 + 0 + 1) = 2 \end{split}$$

By Theorem 4.1,  $\sum_{M \in F(\mathcal{M}, \mathbf{s}; A)} q^{\max(M)} = \prod_{i=1}^{m} {h_i \brack s_i}_q = {2 \brack 1}_q {1 \brack 2}_q {2 \brack 2}_q {2 \brack 1}_q {1 \brack 2}_q {1 \brack 2}_q {1 \brack 2}_q {1 \atop 2}_q {1 \atop$ 



Figure 7: Illustration of Theorem 4.1.

**Corollary 4.3.** Let  $\sigma$  be a permutation of [m] and let  $\mathcal{M}$  be a moon polyomino with columns  $c_1, \ldots, c_m$ . Suppose the shape  $\mathcal{N}$  with columns  $c'_i = c_{\sigma(i)}$  is also a moon polyomino and  $s' = (s'_1, \ldots, s'_m)$  with  $s'_i = s_{\sigma(i)}$ . Then

$$\sum_{M \in \boldsymbol{F}(\mathcal{M}, \boldsymbol{s}; A)} q^{\operatorname{maj}(M)} = \sum_{N \in \boldsymbol{F}(\mathcal{N}, \boldsymbol{s}'; A)} q^{\operatorname{maj}(N)}.$$
(4.3)

That is, the generating function  $\sum_{M \in F(\mathcal{M}, s; A)} q^{\operatorname{maj}(M)}$  does not depend on the order of the columns of  $\mathcal{M}$ .

*Proof.* The formula (4.2) seems to depend on the order  $\prec$  of the columns of the moon polyomino. However, note that  $c_i \prec c_j$  if  $|c_i| < |c_j|$ ; and columns of same length are consecutive in the order  $\prec$ . Hence it suffices to compare the terms in the right-hand side of (4.2) that come from columns of fixed length. If  $c_{i_{j+1}} \prec \cdots \prec c_{i_{j+k}}$  are all the columns of  $\mathcal{M}$  of fixed length, then

$$h_{i_{j+r}} = h_{i_{j+1}} - (s_{i_{j+1}} + \dots + s_{i_{j+r-1}}), \ 2 \le r \le k.$$

$$(4.4)$$

Thus the contribution of these columns to the right-hand side of (4.2) is

$$\prod_{r=1}^{k} \begin{bmatrix} h_{i_{j+r}} \\ s_{i_{j+r}} \end{bmatrix}_{q} = \begin{bmatrix} h_{i_{j+1}} \\ s_{i_{j+1}}, \dots, s_{i_{j+k}}, h_{i_{j+1}} - (s_{i_{j+1}} + \dots + s_{i_{j+k}}) \end{bmatrix}_{q}.$$
(4.5)

The last q-multinomial coefficient is invariant under permutations of  $s_{i_{j+1}}, \ldots, s_{i_{j+k}}$ , and the claim follows.

The right-hand side of formula (4.2) also appeared in the generating function of NE chains, as shown in [11]:

Theorem 4.4 (Kasraoui).

. . .

$$\sum_{M \in \boldsymbol{F}(\mathcal{M}, \boldsymbol{s}; A)} q^{\operatorname{ne}(M)} = \prod_{i=1}^{m} \begin{bmatrix} h_i \\ s_i \end{bmatrix}_q.$$

Combining Theorems 4.1 and 4.4, we have

Theorem 4.5.

$$\sum_{M \in \boldsymbol{F}(\mathcal{M}, \boldsymbol{s}; A)} q^{\operatorname{maj}(M)} = \sum_{M \in \boldsymbol{F}(\mathcal{M}, \boldsymbol{s}; A)} q^{\operatorname{ne}(M)}$$

That is, the maj statistic has the same distribution as the ne statistic over the set  $\mathbf{F}(\mathcal{M}, \mathbf{s}, A)$ . Since  $\operatorname{ne}(\mathcal{M})$  is the natural analogue of inv for words and permutations, Theorem 4.5 generalizes MacMahon's equidistribution result. Unfortunately, as noted in [2], the refinement of MacMahon's theorem which asserts that the joint distribution of maj and inv over permutations is symmetric does not hold even for matchings.

We shall prove a lemma about the major index for words and then use it to prove Theorem 4.1.

**Lemma 4.6.** Let  $w = w_1 \dots w_k$  be a word such that  $w_i < n$  for all i. Consider the set S(w) of all the words w' that can be obtained by inserting m many n's between the letters of w. Then the difference  $\operatorname{maj}(w') - \operatorname{maj}(w)$  ranges over the multiset  $\{i_1 + i_2 + \dots + i_m \mid 0 \leq i_1 \leq i_2 \leq \dots \leq i_m \leq k\}$ . It follows that

$$\sum_{w' \in S(w)} q^{\max(w')} = q^{\max(w)} \sum_{0 \le i_1 \le i_2 \le \dots \le i_m \le k} q^{i_1 + i_2 + \dots + i_m} = q^{\max(w)} {k+m \brack m}_q.$$
 (4.6)

The same statement holds if  $w = w_1 \dots w_k$  is a word such that  $w_i > 1$  for all i and S(w) is the set of all the words w' obtained by inserting m many 1's between the letters of w.

*Proof.* We give an elementary proof for the case of inserting n's. The case of inserting 1's is dealt with similarly. For a word  $w = w_1 \dots w_k$  we define the *descent sequence*  $desseq(w) = a_1 \dots a_{k+1}$  by letting  $a_i = maj(w^{(i)}) - maj(w)$ , where  $w^{(i)}$  is obtained by inserting one n in the *i*-th gap of w, i.e.,  $w^{(i)} = w_1 \dots w_{i-1} n w_i \dots w_k$ . Note that

$$\operatorname{maj}(w^{(i)}) = \operatorname{maj}(w) + \begin{cases} 1 + \operatorname{des}(w_i \dots w_k), & \text{if } w_{i-1} > w_i, \ 2 \le i \le k \text{ or } i = 1\\ i + \operatorname{des}(w_i \dots w_k), & \text{if } w_{i-1} \le w_i, \ 2 \le i \le k\\ 0, & \text{if } i = k+1. \end{cases}$$

Hence, if t = des(w) then desseq(w) is a shuffle of the sequences t + 1, t, ..., 0 and t + 2, t + 3, ..., kwhich begins with t + 1 and ends with 0. Moreover, every such sequence is desseq(w) for some word w. By definition, the multiset  $\{maj(w') - maj(w) \mid w' \in S(w)\}$  depends only on desseq(w) and not on the letters of w.

First we consider the case when desseq(w) = k(k-1)...0, i.e., Des(w) = [k-1]. Inserting b n's between  $w_{i-1}$  and  $w_i$  increases the major index by b(k-i+1). Hence, if one inserts  $b_i$  many n's between  $w_{i-1}$  and  $w_i$  for  $1 \le i \le k+1$ , the major index is increased by

$$\sum_{i=1}^{k+1} b_i(k-i+1) = \underbrace{k+\dots+k}_{b_1} + \underbrace{(k-1)+\dots+(k-1)}_{b_2} + \dots + \underbrace{0+\dots+0}_{b_{k+1}}.$$

Letting  $(b_1, \ldots, b_{k+1})$  range over all (k+1)-tuples with  $\sum_{i=1}^{k+1} b_i = m$ , the increased amount ranges over the multiset  $\{i_1 + i_2 + \cdots + i_m \mid 0 \le i_1 \le i_2 \le \cdots \le i_m \le k\}$ .

To deal with the other possible descent sequences, we note that any descent sequence desseq(w) can be obtained by applying a series of adjacent transpositions to  $k(k-1)\ldots 0$ , where we only transpose adjacent elements i, j which correspond to ascent and descent positions, respectively. Suppose desseq $(w) = a_1 \ldots a_{i-1}a_{i+1}a_ia_{i+2} \ldots a_{k+1}$  is obtained by transposing the *i*-th and (i+1)-st element in desseq $(v) = a_1 \ldots a_{i-1}a_ia_{i+1}a_ia_{i+2} \ldots a_{k+1}$ , where  $a_i$  and  $a_{i+1}$  correspond to an ascent and a descent in v, respectively. Consequently,  $w_{i-1} > w_i \leq w_{i+1}$  and  $v_{i-1} \leq v_i > v_{i+1}$ , while  $w_j \leq w_{j+1}$  if and only if  $v_j \leq v_{j+1}$  for  $j \neq i-1, i$ . It suffices to show that the multisets  $\{\text{maj}(w') - \text{maj}(w) \mid w' \in S(w)\}$  and  $\{\text{maj}(v') - \text{maj}(v) \mid v' \in S(v)\}$  are equal. Suppose  $w' \in S(w)$  is obtained by inserting  $b_j$  many n's in the j-th gap of w,  $1 \leq j \leq k+1$ . Let  $v' \in S(v)$  be the word obtained by inserting  $c_j n$ 's in the j-th gap of v where  $c_j$  are defined as follows.

$$c_{j} = \begin{cases} 0, & \text{if } j = i \\ b_{i}, & \text{if } j = i+1 \\ b_{j}, & \text{if } j \neq i, i+1. \end{cases}$$
(4.7)

If  $b_{i+1} > 0$  then

$$c_{j} = \begin{cases} b_{i} + 1, & \text{if } j = i \\ b_{i+1} - 1, & \text{if } j = i + 1 \\ b_{j}, & \text{if } j \neq i, i + 1. \end{cases}$$
(4.8)

This defines a map between sequences  $(b_j)$  with  $b_{i+1} = 0$  (resp.  $b_{i+1} > 0$ ), and  $(c_j)$  with  $c_i = 0$  (resp.  $c_i > 0$ ), which is a bijection. The *n*'s inserted in the *j*-th gap of *w* for  $j \neq i, i+1$  contribute to maj(w') - maj(w) the same amount as do the *n*'s inserted in the same gaps in *v* to maj(v') - maj(v). The major index contributed to maj(w') by the segment

$$w_{i-1} \underbrace{n \dots n}_{b_i} w_i \underbrace{n \dots n}_{b_{i+1}}$$

is the sum of i-1 (which equals the contribution of  $w_{i-1}w_i$  to maj(w)) and

$$\begin{cases} (b_1 + \dots + b_i) + (b_1 + \dots + b_{i+1} + i), & \text{if } b_{i+1} > 0\\ (b_1 + \dots + b_i), & \text{if } b_{i+1} = 0. \end{cases}$$

Similarly, the major index contributed to maj(v') by the segment

$$v_{i-1} \underbrace{n \dots n}_{c_i} v_i \underbrace{n \dots n}_{c_{i+1}}$$

is the sum of i (which equals the contribution of  $v_{i-1}v_i$  to maj(v)) and

$$\begin{cases} (c_1 + \dots + c_i + i - 1) + (c_1 + \dots + c_{i+1}), & \text{if } c_i > 0\\ (c_1 + \dots + c_{i+1}), & \text{if } c_i = 0. \end{cases}$$

Now, one readily checks that  $\operatorname{maj}(w') - \operatorname{maj}(w) = \operatorname{maj}(v') - \operatorname{maj}(v)$ . This completes the proof.

**Remark 4.7.** Lemma 4.6 can be proved in many ways, for example, in [8, Theorem 1.2] Foata and Han constructed a transformation on the shuffle class  $Sh(0^m v)$ , where v is a nonempty word whose letters are positive integers. Alternatively, one can use the theory of P-partitions [9], which leads to a stronger result, namely, Equation (4.6) holds if  $w = w_1 \dots w_k$  with  $w_i \neq n_0$ ,  $i = 1, \dots, k$ and S(w) is the set of all words w' obtained by inserting m copies of  $n_0$  between the letters of w. To show this, it is enough to consider the case when w is a permutation of length k on k distinct numbers between 1 and k + m such that the integers  $N_1 < N_2 < \cdots < N_m$  are missing from it and w' varies over the set S(w) of all permutations obtained by inserting  $N_1, N_2, \dots, N_m$  in w from left to right. We outline the proof in this general case assuming that the reader is familiar with the theory of P-partitions. We adopt the notation and definitions from Section 4.5 in [21], which is also a good reference for basic results on P-partitions.

Let  $P_1$  be the antichain with k elements labeled with the letters of w. Let  $P_2$  be a disjoint union of  $P_1$  and a chain of m elements labeled  $N_1 < N_2 < \cdots < N_m$ . We denote by  $F_w(q_1, \ldots, q_k)$  the generating function of all w-compatible functions  $f : [k] \to \mathbb{N}$  (see [21] for the definitions). Lemma 4.5.2 in [21] asserts that

$$F_w(q, \dots, q) = \frac{q^{\max(w)}}{(1-q)\cdots(1-q^k)}.$$
(4.9)

All  $P_2$ -partitions compatible with some  $w' \in S(w)$  are obtained by combining a  $P_1$ -partition compatible with w and a integer partition with at most m parts. Therefore,

$$\sum_{w' \in S(w)} F_{w'}(q, \dots, q) = \frac{F_w(q, \dots, q)}{(1-q)\cdots(1-q^m)}.$$
(4.10)

Applying Lemma 4.5.2 in [21] to all  $w' \in S(w)$ , we have

$$\sum_{w' \in S(w)} F_{w'}(q, \dots, q) = \sum_{w' \in S(w)} \frac{q^{\operatorname{maj}(w')}}{(1-q)\cdots(1-q^{k+m})}.$$
(4.11)

Combining equations (4.9), (4.10), and (4.11) gives (4.6).

**Proof of Theorem 4.1.** Let c be the smallest column of  $\mathcal{M}$  in the order  $\prec$  and  $\mathcal{M}' = \mathcal{M} \setminus c$ . Note that c is the leftmost or the rightmost column of  $\mathcal{M}$ . Let M be a filling of  $\mathcal{M}$  with s nonempty cells in c and let M' be its restriction on  $\mathcal{M}'$ . From Definition 2.2', we derive that

$$\operatorname{maj}(M) = \operatorname{maj}(M') + \operatorname{maj}(R_c) - \operatorname{maj}(R_c \cap M'),$$

where  $R_c$  is the filling of the rectangle  $\mathcal{M}(c)$  determined by the column c.  $R_c$  is obtained by adding a column with s many 1-cells to the rectangular filling  $R_c \cap M'$ . Using the bijection between fillings of rectangles and words described in Section 3.1, one sees that if c is the leftmost (resp. rightmost) column in M, this corresponds to inserting s maximal (resp. minimal) elements in the word determined by the filling  $R_c \cap M'$ . When the column c varies over all possible  $\binom{h}{s}$  fillings, by Lemma 4.6, the value maj $(R_c) - \text{maj}(R_c \cap M')$  varies over the multiset  $\{i_1 + i_2 + \cdots + i_s \mid 0 \le i_1 \le i_2 \le \cdots \le i_s \le (h-s)\}$  with generating function  $\sum_{0 \le i_1 \le \cdots \le i_s \le (h-s)} q^{i_1 + \cdots + i_s} = \begin{bmatrix} h \\ s \end{bmatrix}_q$ , where h is the value  $h_1$  (resp.  $h_m$ ) if c is the leftmost (resp. the rightmost) column of M, as defined in (4.1). Therefore,

$$\sum_{M \in \mathbf{F}(\mathcal{M}, \mathbf{s}; A)} q^{\operatorname{maj}(M)} = \begin{bmatrix} h \\ s \end{bmatrix}_q \sum_{M' \in \mathbf{F}(\mathcal{M}', \mathbf{s}'; A)} q^{\operatorname{maj}(M')}$$

where  $\mathbf{s}'$  is obtained from  $\mathbf{s}$  by removing the first (resp. last) component if c is the leftmost (resp. rightmost) column of M. Equation (4.2) now follows by induction on the number of columns of  $\mathcal{M}$ .

### 5 The Foata-type bijection for moon polyominoes

The objective of this section is to give a bijective proof for Theorem 4.5. For the set  $W_S$  of all words of a multiset S, the equidistribution of maj and inv was first proved by MacMahon by combinatorial analysis. This raised the question of constructing a canonical bijection  $\Phi: W_S \to W_S$  such that  $\operatorname{maj}(w) = \operatorname{inv}(\Phi(w))$ . Foata answered the question by constructing an elegant map  $\Phi$  [6], which is referred to as the second fundamental transformation [7]. We begin this section by reviewing Foata's map  $\Phi: W_S \to W_S$ .

Let  $w = w_1 w_2 \cdots w_n$  be a word on N and let *a* be an integer. If  $w_n \leq a$ , the *a*-factorization of *w* is  $w = v_1 b_1 \cdots v_p b_p$ , where each  $b_i$  is a letter less than or equal to *a*, and each  $v_i$  is a word (possibly empty), all of whose letters are greater than *a*. Similarly, if  $w_n > a$ , the *a*-factorization of *w* is  $w = v_1 b_1 \cdots v_p b_p$ , where each  $b_i$  is a letter greater than *a*, and each  $v_i$  is a word (possibly empty), all of whose letters are less than or equal to *a*. In both cases one defines

$$\gamma_a(w) = b_1 v_1 \cdots b_p v_p$$

With the above notation, let  $a = w_n$  and let  $w' = w_1 \cdots w_{n-1}$ . The second fundamental transformation  $\Phi$  is defined recursively by  $\Phi(w) = w$  if w has length 1, and

$$\Phi(w) = \gamma_a(\Phi(w'))a,$$

if w has length n > 1. The map  $\Phi$  has the property that it preserves the last letter of the word, and  $inv(\Phi(w)) = maj(w)$ .

Foata's map  $\Phi$  is constructed recursively with certain "local operations" to eliminate the difference caused by adding the last letter in the words. Inspired by this idea, we construct a Foata-type bijection  $\phi : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  with the property maj $(\mathcal{M}) = \operatorname{ne}(\phi(\mathcal{M}))$ . The map  $\phi$  can be defined directly for left-aligned stack polyominoes. But we describe it first for Ferrers diagrams in Section 5.1, because this case contains all the essential steps and is easy to understand. This map is a revision of the Foata-type bijection presented in the preprint of [2] for set partitions, which correspond to fillings with row and column sums at most 1. Then in Section 5.2 we extend the construction to left-aligned stack polyominoes and prove that  $\operatorname{maj}(\mathcal{M}) = \operatorname{ne}(\phi(\mathcal{M}))$ . In Section 5.3, we construct two bijections, f and g, that transform a filling of a moon polyomino to a filling of a left-aligned stack polyomino and preserve the statistics maj and ne, respectively. Composing these maps with  $\phi$  defined on left-aligned stack polyominoes yields a bijection on  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  that sends maj to ne.

#### 5.1 The bijection $\phi$ for Ferrers diagrams

The empty rows in the filling  $M \in \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  do not play any role in the definitions of maj(M) or  $\operatorname{ne}(M)$ . Therefore, in what follows we assume that  $A = \emptyset$  and describe  $\phi$  for fillings without empty rows. For  $A \neq \emptyset$ , one can first delete the empty rows of the filling, apply the map  $\phi$ , and reinsert the empty rows back.

Let  $\mathcal{F}$  be a Ferrers diagram and F a filling of  $\mathcal{F}$ . The bijection  $\phi$  is defined inductively on the number of rows of F. If F has only one row, then  $\phi(F) = F$ . Otherwise, we denote by  $F_1$  the filling obtained by deleting the top row r of F. Let  $F'_1 = \phi(F_1)$  and let  $F_2$  be the filling obtained by performing the algorithm  $\gamma_r$  described below. Then  $F' = \phi(F)$  is obtained from  $F_2$  by adding the top row r. So, by definition,  $\phi$  preserves the first row of F.

#### The algorithm $\gamma_r$

If C is the 1-cell of r, then denote by  $\mathcal{R}$  the set of all rows of  $F'_1$  that intersect the column of C. The 1-cells in  $\mathcal{R}$  that are strictly to the left of C are called left and the 1-cells in  $\mathcal{R}$  that are weakly to the right of C are called right. The cell C is neither left nor right.

Let CLC (critical left cell) be the topmost left 1-cell, and CRC (critical right cell) be the leftmost right 1-cell that is above CLC. If there is more than one such cell the CRC is defined to be the lowest one. Note that CLC and CRC need not exist. Denote by  $\mathcal{R}_1$  the set of all rows weakly below the row of CRC that intersect the column of CRC and  $\mathcal{R}_2$  is the set of all rows in  $\mathcal{R}$ that do not intersect the column of CRC. If CLC does not exist then both  $\mathcal{R}_1$  and  $\mathcal{R}_2$  are empty. If CLC exists but CRC does not, then  $\mathcal{R}_1$  is empty and  $\mathcal{R}_2$  contains all the rows in  $\mathcal{R}$  weakly below CLC. See Figure 8 for an illustration.



Figure 8: The *CRC* and *CLC* are in rows 5 and 6, respectively.  $\mathcal{R}_1$  contains rows 5, 6, and 7, while  $\mathcal{R}_2$  contains rows 8 and 9.

**Definition 5.1.** Let  $C_1$  and  $C_2$  be two 1-cells with coordinates  $(i_1, j_1)$  and  $(i_2, j_2)$ , respectively. We swap the cells  $C_1$  and  $C_2$  by deleting the 1's from these two cells and write 1's in the cells with coordinates  $(i_1, j_2)$  and  $(i_2, j_1)$ .

For a cell C, denote by col(C) the column of C, and |col(C)| the length of col(C).

## Algorithm $\gamma_r^1$ on $\mathcal{R}_1$

Let  $ptr_1$  and  $ptr_2$  be two pointers.

- (A) Set  $ptr_1$  on the highest row of  $\mathcal{R}_1$  and  $ptr_2$  on the next row in  $\mathcal{R}_1$  below  $ptr_1$ .
- (B) If  $ptr_2$  is null, then go to (D). Otherwise, the pointers  $ptr_1$  and  $ptr_2$  point at 1-cells  $C_1$  and  $C_2$ , respectively.

- (B1) If  $C_2$  is a left cell then swap the cells  $C_1$  and  $C_2$  and move  $ptr_1$  to the row of  $ptr_2$ .
- (B2) If  $C_2$  is a right cell, then
  - (B2.1) If  $|col(C_1)| = |col(C_2)|$  then move  $ptr_1$  to the row of  $ptr_2$ .
  - (B2.2) If  $C_1$  is to the left of  $C_2$  and  $|col(C_1)| > |col(C_2)|$  then do nothing.
  - (B2.3) If  $C_1$  is to the right of  $C_2$  and  $|col(C_1)| < |col(C_2)|$ , then find the lowest left 1-cell L that is above  $C_1$ . Suppose that the row-column coordinates of the cells L,  $C_1$ , and  $C_2$  are  $(i_1, j_1)$ ,  $(i_2, j_2)$ , and  $(i_3, j_3)$ , respectively. Delete the 1's from these three cells and write them in the cells with coordinates  $(i_1, j_3)$ ,  $(i_2, j_1)$ , and  $(i_3, j_2)$ . Move  $ptr_1$  to the row of  $ptr_2$ .<sup>†</sup>
- (C) Move  $ptr_2$  to the next row in  $\mathcal{R}_1$ . Go to (B).

(D) Stop.

See Figure 9 for illustration of the steps.

<sup>†</sup>**Note:** This step is well-defined since it cannot occur before  $ptr_2$  reaches CLC (that would contradict the definition of CRC) and, after that,  $ptr_1$  is always below CLC. Therefore, the 1-cell L always exists. The fact that the square  $(i_3, j_2)$  belongs to  $\mathcal{F}$  follows from the definition of  $\mathcal{R}_1$  and Lemma 5.2 (c).

When the algorithm  $\gamma_r^1$  stops, we continue processing the rows of  $\mathcal{R}_2$  by algorithm  $\gamma_r^2$ .

### The algorithm $\gamma_r^2$ on $\mathcal{R}_2$

- (A') Set  $ptr_1$  on the highest row of  $\mathcal{R}_2$  and  $ptr_2$  on the next row in  $\mathcal{R}_2$  below  $ptr_1$ .
- (B') (*Borrowing*) If  $ptr_1$  points to a right 1-cell then find the lowest left 1-cell above it and swap them. Now  $ptr_1$  points to a left 1-cell.
- (C') If  $ptr_2$  is null, then go to (E'). Otherwise, the pointers  $ptr_1$  and  $ptr_2$  point at 1-cells  $C_1$  and  $C_2$  respectively.
  - (C'1) If  $C_2$  is a right cell then swap  $C_1$  and  $C_2$ .
  - (C'2) If  $C_2$  is a left cell then do nothing.
- (D') Move  $ptr_1$  to the row of  $ptr_2$  and  $ptr_2$  to the next row in  $\mathcal{R}_2$  below. Go to (C').
- (E') Stop.

See Figure 10 for illustration of the steps in  $\gamma_r^2$ .

Let us note the following easy but useful properties of the algorithm  $\gamma_r$ .

- 1. The pointers  $ptr_1$  and  $ptr_2$  process the rows of  $\mathcal{R}_1$  and  $\mathcal{R}_2$  from top to bottom. However, while  $ptr_2$  always moves from one row to the next one below it,  $ptr_1$  sometimes stays on the same row (cf. step (B2.2)) and sometimes "jumps" several rows below (cf. step (B1)).
- 2. Pointer  $ptr_1$  always points to a right 1-cell in  $\gamma_r^1$  and to a left 1-cell in  $\gamma_r^2$ .

Further we show some nontrivial properties which will be used to show that the map  $\phi$  has the desired property maj $(F) = \operatorname{ne}(\phi(F))$ .



Step (B1).



Figure 9: The algorithm  $\gamma_r^1$  on  $\mathcal{R}_1$ .

**Lemma 5.2.** Suppose in  $\gamma_r^1$  ptr<sub>1</sub> is pointing to  $C_1$  and ptr<sub>2</sub> is pointing to  $C_2$ . Then

- (a) In the rows between  $C_1$  and  $C_2$  there are no 1's weakly to the left of  $C_1$ .
- (b) If L is the lowest left 1-cell above  $C_1$ , then in the rows between L and  $C_1$  there are no 1's weakly to the left of  $C_1$ .
- (c) The 1-cell  $C_1$  is in a column with same length as the column of the critical right cell CRC. Moreover, when  $\gamma_r^1$  stops  $ptr_1$  points to the last row in  $\mathcal{R}_1$ .

*Proof.* The first two parts are proved by induction on the number of steps performed in the algorithm.

- (a) The only step which leaves the gap between  $ptr_1$  and  $ptr_2$  nonempty is (B2.2). But after this step, the number of 1's in the rows between  $ptr_1$  and  $ptr_2$  weakly to the left of  $C_1$  does not change.
- (b) The case when the last step is (B2.2) follows from the induction hypothesis. If the last step is (B1) or (B2.3) the claim follows from part (a). The case when the last step was (B2.1)



Figure 10: Algorithm  $\gamma_r^2$  on  $\mathcal{R}_2$ .

and the cells  $C_1$  and  $C_2$  formed an NE chain the claim also follows from part (a) and the induction hypothesis. If in (B2.1), the cells  $C_1$  and  $C_2$  do not form an NE chain, in addition to the induction hypothesis, one uses the fact that there are no 1's in the rows between these two cells which are weakly to the left of  $C_2$ .

(c) The first part follows immediately from the definition of the steps. If the second part is not true, then the last step must be (B2.2). and the column of  $C_1$  is longer than the column of  $C_2$ . Since this is the last step of  $\gamma_r^1$ , it follows that the column of  $C_1$  intersects the top row of  $\mathcal{R}_2$ . However, the column of CRC does not intersect the top row of  $\mathcal{R}_2$  by definition of  $\mathcal{R}_2$ . This contradicts the fact that  $C_1$  and CRC are in columns of same length.

**Proposition 5.3.** During algorithm  $\gamma_r^1$  the number of NE chains decreases by the total number of left 1-cells in  $\mathcal{R}_1$ , whereas during algorithm  $\gamma_r^2$  the number of NE chains increases by the total number of right 1-cells in  $\mathcal{R}_2$ .

*Proof.* To prove the first part note that pointer  $ptr_2$  points to each left 1-cell in  $\mathcal{R}_1$  exactly once in the algorithm  $\gamma_r^1$ . When that happens, step (B1) is performed during which, by Lemma 5.2(a), the number of NE chains decreases by one. In the other steps, the number of NE chains remains unchanged. This is trivial for steps (B2.1) and (B2.2), whereas for step (B2.3) it follows from Lemma 5.2 (a) and (b).

For the second part, note that pointer  $ptr_2$  points to each right 1-cell in  $\mathcal{R}_2$  exactly once in the algorithm  $\gamma_r^2$  with possible exception being the top 1-cell in  $\mathcal{R}_2$  to which it never points. In those steps the number of NE chains is increased by one, while otherwise it remains the same. When the top 1-cell is right, then Borrowing occurs. It follows from parts (b) and (c) of Lemma 5.2 that the number of NE chains is also increased by one.

**Theorem 5.4.** The map  $\phi : \mathbf{F}(\mathcal{F}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{F}, \mathbf{s}; A)$  is a bijection.

The proof is quite technical and is given in the appendix. Here we only present an example of the bijection  $\phi$  in Figure 11 and explain how it includes Foata's second fundamental transformation. If the moon polyomino is rectangular, then the algorithm  $\gamma_r$  has a simpler description because some of the cases can never arise. Namely, if the cell in the second row forms an NE chain with the top cell (a descent) then  $\mathcal{R}_1 = \emptyset$ , and only the steps (C'1) and (C'2) are performed. Otherwise,  $\mathcal{R}_2 = \emptyset$ and, since all columns are of equal height, only the steps (B1) and (B2.1) are performed. In both



Figure 11: Example of the map  $\phi$  applied inductively on  $F = F_7$ . The fillings  $F_i$  are restrictions of F on the last i rows.

these cases, via the correspondence between words and rectangular fillings (c.f. Section 3.1) the algorithms  $\gamma_r^2$  and  $\gamma_r^1$  for rectangles are equivalent to Foata's transformation for words [6].

### 5.2 The bijection $\phi$ for left-aligned stack polyominoes

Next we extend the bijection  $\phi$  to fillings of left-aligned stack polyominoes which sends maj to ne. Let  $\mathcal{M}$  be a left-aligned stack polyomino. As in the case of Ferrers diagrams, we can assume that  $A = \emptyset$  and only consider fillings without empty rows. Suppose M is a filling of  $\mathcal{M}$  with top row r. The map, which is again denoted by  $\phi$ , is defined inductively on the number of rows of M. If M has only one row, then  $\phi(M) = M$ . Otherwise, let  $\mathcal{F}$  be the maximal Ferrers diagram in  $\mathcal{M}$  that contains the top row and  $\mathcal{F}_1$  be  $\mathcal{F}$  without the top row. Let  $F = M \cap \mathcal{F}$  and  $F_1 = M \cap \mathcal{F}_1$ . To obtain  $\phi(M)$  we perform the following steps. (See Figure 12).

- 1. Delete the top row of M and get  $M_1 = M \setminus r$ .
- 2. Apply  $\phi$  to  $M_1$  and get  $M'_1 = \phi(M_1)$ .
- 3. Apply the algorithm  $\gamma_r$  to the filling  $F'_1 = M'_1 \cap \mathcal{F}_1$  and leave the cells in  $M'_1$  outside of  $F'_1$  unchanged. Denote the resulting filling by  $M_2$ .
- 4.  $\phi(M)$  is obtained by adding row r back to  $M_2$ .

It is a bijection from  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  to itself since every step is invertible. That the Step 3 is invertible follows from the proof of Theorem 5.4.



Figure 12: Illustration of  $\phi$  for left-aligned stack polyominoes.

Lemma 5.5. Using the notation introduced above,

$$\operatorname{maj}(M) - \operatorname{maj}(M_1) = \operatorname{maj}(F) - \operatorname{maj}(F_1) = \#\mathcal{R}_2(F_1)$$
 (5.1)

where  $\mathcal{R}_2(F_1)$  is defined with respect to the top row r of F as in Section 5.1.

*Proof.* Let S be the filling M restricted to the rows which are not completely contained in  $\mathcal{F}$ . From the definition of maj, we have

$$\operatorname{maj}(M) = \operatorname{maj}(F) + \operatorname{maj}(S) - \operatorname{maj}(F \cap S)$$
(5.2)

$$\operatorname{maj}(M_1) = \operatorname{maj}(F_1) + \operatorname{maj}(S) - \operatorname{maj}(F_1 \cap S)$$
(5.3)

Note that  $F \cap S = F_1 \cap S$ , hence (5.2) and (5.3) give the first equality in (5.1). The second one is readily checked using Proposition 3.1 for major index of stack polyminoes.

The rest of this subsection is devoted to proving that the map  $\phi$  has the desired property  $\operatorname{maj}(M) = \operatorname{ne}(\phi(M))$ . We shall need one more lemma.

Let C be a 1-cell in a filling M of a left-aligned stack polyomino. We say that C is maximal if there is no 1 in M which is strictly above and weakly to the left of C. Clearly, a maximal cell must be the highest 1-cell in its column. For a maximal cell C in M, we use t(C, M) to denote the index of the leftmost column (the columns are numbered from left to right) to the right of C that contains a 1-cell which together with C forms an NE chain. If such a column does not exist set  $t(C, M) = \infty$ .

**Lemma 5.6.** A 1-cell C in M is maximal if and only if the highest 1-cell C' in the same column in  $M' = \phi(M)$  exists and is maximal. Moreover, t(C, M) = t(C', M') when they are both maximal.

*Proof.* We proceed by induction on the number of rows of M and keep the notation as in Figure 11. The case when M has one row is trivial.

Suppose that the top row r of M has a 1-cell  $C^*$  in the column  $i^*$ . The cell  $C^*$  is maximal in both M and M' and  $t(C^*, M) = \infty = t(C^*, M')$ . Note that a 1-cell  $C \neq C^*$  in M (resp. M') is maximal if and only if C is in column  $i < i^*$  and is maximal in  $M_1$  (resp.  $M_2$ ).

Denote by  $C_1$ ,  $C'_1$ , and  $C_2$  the highest 1-cells in column  $i < i^*$  in the fillings  $M_1$ ,  $M'_1 = \phi(M_1)$ , and  $M_2$ , respectively. By the induction hypothesis,  $C_1$  is maximal in  $M_1$  if and only if  $C'_1$  is maximal in  $M'_1$ .  $M_2$  is obtained from  $M'_1$  by applying the algorithm  $\gamma_r$  with respect to the row r to  $F'_1$ .  $C'_1$  is a left cell in this algorithm, and since  $\gamma_r$  does not change the relative position of the left 1-cells,  $C_2$  is maximal in  $M_2$ .

It only remains to prove that t(C, M) = t(C', M'). Suppose  $t(C_1, M_1) = t(C'_1, M'_1) = a$  (they are equal by the inductive hypothesis).

If  $a < i^*$  then t(C, M) = a. In  $M'_1$  it corresponds to a 1-cell D' in a column left of  $C^*$  such that  $C'_1, D'$  form an NE chain. Since the algorithm  $\gamma_r$  preserves the column sums and the relative positions of left 1-cells, we have  $t(C_2, M_2) = a$ , and hence t(C', M') = a.

If  $a \ge i^*$  then  $t(C, M) = i^*$ . The fact that  $t(C'_1, M'_1) = a \ge i^*$  means that there is no left 1-cell in  $M'_1$  above and to the right of  $C'_1$  and, since  $\gamma_r$  does not change the relative positions of the left 1-cells, there will be no left 1-cell above and to the right of  $C_2$  in  $M_2$ . So, in this case, we conclude  $t(C', M') = i^* = t(C, M)$ .

**Theorem 5.7.** If M is a filling of a left-aligned stack polynomial  $\mathcal{M}$ , then  $\operatorname{maj}(M) = \operatorname{ne}(\phi(M))$ .

*Proof.* Again we proceed by induction on the number of rows of M. The case when M has one row is trivial. We use the notation as in Figure 11, and let  $F_2 = M_2 \cap \mathcal{F}_1$ .

By Proposition 5.3, the algorithm  $\gamma_r$  on  $M'_1$  decreases ne by one for each left 1-cell in  $\mathcal{R}_1(M'_1)$ and increases it by one for each right 1-cell in  $\mathcal{R}_2(M'_1)$ . Therefore,

$$ne(\phi(M)) = ne(M_2) + \#\{\text{left 1-cells in } \mathcal{R}(F_2)\} \\ = ne(M'_1) - \#\{\text{left 1-cells in } \mathcal{R}_1(F'_1)\} + \#\{\text{right 1-cells in } \mathcal{R}_2(F'_1)\} \\ + \#\{\text{left 1-cells in } \mathcal{R}(F_2)\}$$

Since  $\gamma_r$  preserves the column sums,

$$\#\{\text{left 1-cells in } \mathcal{R}(F_2)\} = \#\{\text{left 1-cells in } \mathcal{R}(F_1)\}$$

hence

$$\operatorname{ne}(\phi(M)) = \operatorname{ne}(M_1') + \#\mathcal{R}_2(F_1')$$

One the other hand, from Lemma 5.5 and the induction hypothesis, one gets

$$\operatorname{maj}(M) = \operatorname{maj}(M_1) + \#\mathcal{R}_2(F_1) = \operatorname{ne}(M'_1) + \#\mathcal{R}_2(F_1).$$

So, it suffices to show that

$$#\mathcal{R}_2(F_1) = #\mathcal{R}_2(F_1'). \tag{5.4}$$

To show equation (5.4), note that for a filling of a Ferrers diagram F the number of rows in  $\mathcal{R}_2(F)$  is determined by the column indices of CLC (critical left cell) and CRC (critical right cell). The CLC is the topmost left 1-cell which, if exists, is the topmost maximal cell besides the 1-cell in the first row. By Lemma 5.6, CLC in  $F_1 \subseteq M_1$  is in the same column as CLC' in  $F'_1 \subseteq M'_1$ , or neither of them exists. The latter case is trivial, as  $\mathcal{R}_2(F)$  is empty. In the first case, let |r| denote the length of row r. Then CRC, the critical right cell of  $F_1$ , is in the column  $t(CLC, M_1)$  if  $t(CLC, M_1) \leq |r|$  and does not exist otherwise. Similarly, the critical right cell of  $F'_1$ , CRC', is in the column  $t(CLC', M'_1)$  if  $t(CLC, M'_1) \leq |r|$ , and does not exist otherwise. By Lemma 5.6,  $t(CLC, M_1) = t(CLC', M'_1)$ , and this implies (5.4).

#### 5.3 The case of a general moon polyomino

Now we consider the case when  $\mathcal{M}$  is a general moon polyomino. We label the rows of  $\mathcal{M}$  by  $r_1, \ldots, r_n$  from top to bottom. Let  $\mathcal{N}$  be the unique left-aligned stack polyomino whose sequence of row lengths is equal to  $|r_1|, \ldots, |r_n|$  from top to bottom. In other words,  $\mathcal{N}$  is the left-aligned polyomino obtained by rearranging the columns of  $\mathcal{M}$  by length in weakly decreasing order from left to right. For the definitions that follow, we use an idea of Rubey [19] to describe an algorithm that rearranges the columns of  $\mathcal{M}$  to obtain  $\mathcal{N}$  (Figure 13).

Algorithm  $\alpha$  for rearranging  $\mathcal{M}$ :

1. Set  $\mathcal{M}' = \mathcal{M}$ .

2. If  $\mathcal{M}'$  is left aligned go to (4).

3. If  $\mathcal{M}'$  is not left-aligned consider the largest rectangle  $\mathcal{R}$  completely contained in  $\mathcal{M}'$  that contains  $c_1$ , the leftmost column of  $\mathcal{M}'$ . Update  $\mathcal{M}'$  by letting  $\mathcal{M}'$  be the polyomino obtained by moving the leftmost column of  $\mathcal{R}$  to the right end. Go to (2). 4. Set  $\mathcal{N} = \mathcal{M}'$ .



Figure 13: The algorithm  $\alpha$ .

In this section we give two bijections  $f, g : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{N}, \mathbf{s}'; A)$  which preserve the major index and the number of NE chains, respectively. The sequence  $\mathbf{s}'$  is obtained by rearranging the sequence  $\mathbf{s}$  in the same way  $\mathcal{N}$  is obtained by rearranging the columns of  $\mathcal{M}$ .

**5.3.1** Bijection  $f : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{N}, \mathbf{s}'; A)$  such that  $\operatorname{maj}(M) = \operatorname{maj}(f(M))$ 

Let R be a filling of a rectangle  $\mathcal{R}$  with column sums  $s_1, s_2, \ldots, s_m$ . First, we describe a transformation  $\tau$  which gives a filling of  $\mathcal{R}$  with column sums  $s_2, s_3, \ldots, s_m, s_1$  and preserves the major index. Recall that a descent in R is a pair of 1's in consecutive nonempty rows that form an NE chain. Define an *ascent* to be a pair of 1's in two consecutive nonempty rows that does not form an NE chain. Suppose the first column of R has k many 1-cells:  $C_1, \ldots, C_k$  from top to bottom.

Transformation  $\tau$  on rectangles:

- 1. Let  $R^*$  be the rectangle obtained by adding one empty column to R from right. Process the 1-cells  $C_1, \ldots, C_k$  from bottom to top (see Figure 14):
- 2. For  $r = k, k 1, \ldots, 1$  do the following:

Let  $D_a$  be the lowest 1-cell above  $C_r$  in  $R^*$  and  $D_b$  be the highest 1-cell below  $C_r$  in  $R^*$ .



Figure 14: Illustration of the transformation  $\tau$  for rectangles.

- (a) If  $D_a$  does not exist or  $D_a = C_{r-1}$ , and  $D_b$  does not exist, just move  $C_r$  horizontally to the last column of  $R^*$ ;
- (b) If (i) D<sub>a</sub> does not exist or D<sub>a</sub> = C<sub>r-1</sub>, but D<sub>b</sub> exists, or (ii) both D<sub>a</sub> and D<sub>b</sub> exist, and D<sub>a</sub> is strictly to the right of D<sub>b</sub>:
  Let D<sub>1</sub> = C<sub>r</sub>, D<sub>2</sub> = D<sub>b</sub>, D<sub>3</sub>,..., D<sub>p</sub> be the maximal chain of consecutive ascents in R<sup>\*</sup>. Move D<sub>i</sub> horizontally to the column of D<sub>i+1</sub>, for i = 1,..., p − 1, and move D<sub>p</sub> horizontally to the last column of R<sup>\*</sup>. (If D<sub>p</sub> is in the last column of R<sup>\*</sup>, it remains
  - there.)
- (c) If  $D_a \neq C_{r-1}$ , and  $(D_a \text{ is weakly to the left of } D_b \text{ or } D_b \text{ does not exist})$ : Let  $D_1 = C_r, D_2 = D_a, D_3, \ldots, D_p$  be the maximal chain of consecutive descents in  $R^*$ . Move  $D_i$  horizontally to the column of  $D_{i+1}$ , for  $i = 1, \ldots, p-1$ , and move  $D_p$  horizontally to the last column of  $R^*$ .
- 3. Delete the first column of  $R^*$ .

Note that each iteration in Step 2 decreases the number of 1's in the first column of  $R^*$  by one and increases the number of 1's in the last column by one, while the other column sums remain unchanged.

**Lemma 5.8.** The transformation  $\tau$  is invertible. Moreover, it preserves the descents and hence  $\operatorname{maj}(R) = \operatorname{maj}(\tau(R))$ .

*Proof.* The map  $\tau$  is invertible since  $\tau^{-1}$  can be obtained by taking  $\rho \circ \tau \circ \rho$ , where  $\rho$  is the rotation of rectangles by 180°. Moreover, each iteration of Step 2 of  $\tau$  preserves the positions of descents, hence the second part of the claim holds.

Now we are ready to define the map f. Suppose  $M \in \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$ . We perform the algorithm  $\alpha$  on M to transform the shape  $\mathcal{M}$  to  $\mathcal{N}$ . While we are in Step 3, instead of just moving the first column of  $\mathcal{R}$  to the right end, we perform the algorithm  $\tau$  on the filling R of  $\mathcal{R}$ . f(M) is defined to be the resulting filling of  $\mathcal{N}$ .

**Proposition 5.9.** The map  $f : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{N}, \mathbf{s}'; A)$  is a bijection and  $\operatorname{maj}(M) = \operatorname{maj}(f(M))$ .

Proof. The first part follows from Lemma 5.8 and the fact that  $\alpha$  is invertible. The filling f(M) is obtained after several iterations of Step 3 in the algorithm  $\alpha$  combined with application of  $\tau$ . For the second part of the claim, it suffices to show that maj is preserved after one such iteration. Let M' be the filling of the shape  $\mathcal{M}'$  obtained from M after one step of f in which the rectangle R is transformed into  $\tau(R)$ . We shall use Definition 2.2 to compare maj(M) and maj(M'). Let  $R_1, \ldots, R_k$  (resp.  $R'_1, \ldots, R'_k$ ) be the maximal rectangles of M (resp. M'), ordered by height. In particular,  $R_i$  and  $R'_i$  are of same size. Suppose  $R = R_s$ , then  $R'_s = \tau(R)$ . We partition the set of rectangles into three subsets according to i < s, i = s, and i > s, and compare the contribution of  $R_i$ 's in each subset.

1. For  $1 \leq i \leq s-1$ , the rectangles  $R_i$  and  $R_i \cap R_{i+1}$  are of smaller height than R and contain several consecutive rows of R. Using Lemma 5.8 we know that the descents in  $R'_i \cap \tau(R)$ (resp.  $R'_i \cap R'_{i+1} \cap \tau(R)$ )  $1 \leq i \leq s-1$ , are in the same rows as in  $R_i \cap R$  (resp.  $R_i \cap R_{i+1} \cap R$ ). And descents formed by one cell in R and one cell on the right of R are preserved since  $\tau$ preserves the set of rows that contain a non-zero entry. Consequently,

$$\operatorname{maj}(R_i) = \operatorname{maj}(R'_i), \ \operatorname{maj}(R_i \cap R_{i+1}) = \operatorname{maj}(R'_i \cap R'_{i+1}), \ 1 \le i \le s - 1.$$
(5.5)

2. For i = s, by Lemma 5.8, we have

$$\operatorname{maj}(R_s) = \operatorname{maj}(R'_s). \tag{5.6}$$

3. For i > s, the height of the rectangle  $R_i$  or  $R_{i-1} \cap R_i$  is greater than or equal to that of R. Each of these rectangles contains several consecutive columns of R excluding the first one. The filling of each subrectangle of R consisting of the columns  $c_j, \ldots, c_{j+m}$  (j > 1) is equal, up to a rearrangement of the empty rows, to the filling of the subrectangle of  $\tau(R)$  consisting of the columns  $c_{j-1}, \ldots, c_{j+m-1}$ . To see this, note that the shifting of the 1-cells  $D_1, \cdots, D_p$ horizontally in the algorithm  $\tau$  can be viewed as moving the 1-cell  $D_i$  vertically to the row of  $D_{i-1}$  and moving  $D_1$  to the last column of  $R^*$  in the row of  $D_p$ . Thus, the descent positions in the fillings  $R'_i \cap \tau(R)$  (resp.  $R'_{i-1} \cap R'_i \cap \tau(R)$ ), i > s are the same as in  $R_i \cap R$  (resp.  $R_{i-1} \cap R_i \cap R$ ). Finally, one checks that the descents formed by one cell in R and one cell outside R are also preserved. Therefore,

$$maj(R_i) = maj(R'_i), \ maj(R_{i-1} \cap R_i) = maj(R'_{i-1} \cap R'_i), \ i > s.$$
(5.7)

Combining (5.5), (5.6), and (5.7) gives maj(M) = maj(M').

### **5.3.2** Bijection $g : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{N}, \mathbf{s}'; A)$ such that ne(M) = ne(g(M))

Suppose  $M \in \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$ . To obtain g(M), we perform the algorithm  $\alpha$  to transform the shape  $\mathcal{M}$  to  $\mathcal{N}$  and change the filling when we move columns in Step 3 so that the number of 1's in each row and column is preserved.

Let R be the rectangular filling in Step 3 of  $\alpha$  that contains the column  $c_1$  of the current filling. Suppose  $c_1$  contains k many 1-cells  $C_1, \ldots, C_k$  from top to bottom. Shade the empty rows of R and the cells in R to the right of  $C_1, \ldots, C_k$ . Let  $l_i$  denote the number of empty white cells in R above  $C_i$ . If  $\mathcal{R}'$  is the rectangle obtained by moving the column  $c_1$  from first to last place, fill it to obtain a filling R' as follows.

- 1. The rows that were empty remain empty. Shade these rows.
- 2. Write k many 1's in the last column from bottom to top so that there are  $l_i$  white empty cells below the *i*-th 1,  $1 \le i \le k$ .
- 3. Shade the cells to the left of the nonempty cells in the last column.
- 4. Fill in the rest of the rectangle by writing 1's in the unshaded rows of  $\mathcal{R}'$  so that the unshaded part of R to the right of the first column is the same as the unshaded part of R' to the left of the last column.

See Figure 15 for an example.



Figure 15: One step of the map g.

**Proposition 5.10.** The map  $g: \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{N}, \mathbf{s}'; A)$  is a bijection and  $\operatorname{ne}(M) = \operatorname{ne}(g(M))$ .

*Proof.* Clearly, every step of g is invertible. To see that ne is preserved, it suffices to show that it is preserved in each step of g, when one column is moved from left to right. First note that ne(R) = ne(R'). This follows from the fact that the 1's in the first column of R form  $l_1, \ldots, l_k$  NE chains from top to bottom, while the 1's in the last column of R' form  $l_1, \ldots, l_k$  NE chains from bottom to top and the remaining parts of R and R' are essentially equal. The numbers of NE chains in M and M' respectively made of at least one 1 which is outside of R and R' respectively are equal because each step of g preserves the row and column sums.

**Theorem 5.11.** Let  $\phi$  denote the Foata-type bijection described in Section 5.2. For a general moon polyomino M the map  $\psi = g^{-1} \circ \phi \circ f : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  is a bijection with the property

$$\operatorname{maj}(M) = \operatorname{ne}(\psi(M)) \tag{5.8}$$

*Proof.* The maps f and g permute the columns of M together with the corresponding number of 1's, while  $\phi$  preserves the column sums. Moreover, all three of them preserve the empty rows. Therefore,  $\psi$  is indeed a map from  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  to itself. Finally, (5.8) follows from Proposition 5.9, Proposition 5.10, and Theorem 5.7.

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The authors would like to thank Ira Gessel [9] for suggesting a connection between the enumeration of permutations by major index and the theory of P-partitions.

## Appendix

To show that the map  $\phi : \mathbf{F}(\mathcal{F}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{F}, \mathbf{s}; A)$  is a bijection, it suffices to describe the inverse of the algorithm  $\gamma_r$ . First, we need to determine what  $\mathcal{R}_1$  and  $\mathcal{R}_2$  were. For that we use the following properties of  $\gamma_r$ .

- 1. When  $\gamma_r^1$  stops, the pointer  $ptr_1$  is at the lowest row of  $\mathcal{R}_1$  pointing at a right 1-cell  $C_1$ . The algorithm  $\gamma_r^2$  does not change the cell  $C_1$  and when the whole algorithm  $\gamma_r$  terminates, there is no 1-cell below  $C_1$  that together with it forms an NE chain.
- 2. After performing  $\gamma_r^2$ , all the right 1-cells in  $\mathcal{R}_2$  have at least one 1-cell below them with which they form an NE chain.
- 3. If  $\mathcal{R}_2$  was nonempty then, after performing  $\gamma_r$ , the lowest row of  $\mathcal{R}_2$  contains a left 1-cell.
- 4. If any borrowing occurred and the right cell  $C_1 = (i_1, j_1)$  and the left cell  $C_2 = (i_2, j_2)$  were swapped, then right after this step the cell  $(i_1, j_2)$  is a right 1-cell in  $\mathcal{R}_1$  which forms an NE chain with a left 1-cell from  $\mathcal{R}_2$  and this is the lowest 1-cell in  $\mathcal{R}_1$  with this property.

Therefore, if  $\mathcal{R}$  has no left 1-cells then  $\mathcal{R}_2 = \mathcal{R}_1 = \emptyset$ . Otherwise, find the lowest right 1-cell  $C^*$ in  $\mathcal{R}$  such that there is no 1-cell in  $\mathcal{R}$  below it that together with  $C^*$  forms an NE chain. Then, by Properties 1 and 2 all the nonempty rows in  $\mathcal{R}$  below  $C^*$  are in  $\mathcal{R}_2$ . If there is no 1-cell  $C^*$  with that property, then  $\mathcal{R}_2 = \mathcal{R}$ .

### Algorithm $\delta_r^2$ : the inverse of $\gamma_r^2$

- (IA') Initially, set  $ptr_1$  to the lowest row of  $\mathcal{R}_2$  and  $ptr_2$  to the next row in  $\mathcal{R}_2$  above  $ptr_1$ .
- (IB') If  $ptr_2$  is null then go to step (ID'). Otherwise, suppose  $ptr_1$  and  $ptr_2$  are pointing at 1-cells  $C_1$  and  $C_2$ .
  - (IB'1) If  $C_2$  is a left cell do nothing.
  - (IB'2) If  $C_2$  is a right cell then swap the cells  $C_1$  and  $C_2$ .
- (IC') Move  $ptr_1$  to the row of  $ptr_2$  and  $ptr_2$  to the next row in  $\mathcal{R}_2$  above it. Go to (IB').

- (ID') (*Inverse borrowing*) Note that  $ptr_1$  always points to a left 1-cell. When it reaches the highest row of  $\mathcal{R}_2$  and points to a 1-cell  $C_1$  first we need to determine whether there was any borrowing. For that purpose find the lowest right 1-cell  $C_2$  above  $C_1$  such that the two cells form an NE chain. Using Property 4 we conclude:
  - (ID'1) If there is no such a cell  $C_2$ , then there was no borrowing so do nothing.
  - (ID'2) If there is such a cell  $C_2$ , and there is a left 1-cell in the rows between  $C_1$  and  $C_2$ , then there was no borrowing so do nothing.
  - (ID'3) If there is such a cell  $C_2$ , and there is no left 1-cell between  $C_1$  and  $C_2$ , then swap  $C_1$  and  $C_2$ .

**(IE')** Stop.

When the algorithm  $\delta_r^2$  stops, continue by applying the algorithm  $\delta_r^1$  on  $\mathcal{R} \setminus \mathcal{R}_2$ .

Algorithm  $\delta_r^1$ : the inverse of  $\gamma_r^1$ 

- (IA) Position  $ptr_1$  on the lowest row in  $\mathcal{R} \setminus \mathcal{R}_2$  and  $ptr_2$  on the lowest nonempty row in  $\mathcal{R} \setminus \mathcal{R}_2$  above it.
- (IB) If  $ptr_2$  is null then go to step (ID). Otherwise, suppose that  $ptr_1$  and  $ptr_2$  point at the 1-cells  $C_1$  and  $C_2$  respectively.
  - (IB1) If  $C_2$  is a left cell then check whether there exists a right 1-cell R above  $C_2$  in a column longer than the column of  $C_1$  such that there are no left 1-cells between R and  $C_2$ .
    - (IB1.1) If such a cell R exists, then suppose that the row-column coordinates of  $C_1$ ,  $C_2$ , and R are  $(i_1, j_1)$ ,  $(i_2, j_2)$ , and  $(i_3, j_3)$ , respectively. Delete the 1's from these three cells and write them in the cells with coordinates  $(i_1, j_3)$ ,  $(i_2, j_1)$ , and  $(i_3, j_2)$ . Move  $ptr_1$  to the row of  $ptr_2$ .
    - (IB1.1) If there is no such a cell R, then swap  $C_1$  and  $C_2$  and move  $ptr_1$  to the row of  $ptr_2$ .
  - (IB2) If  $C_2$  is a right cell, then

(IB2.1) If  $|col(C_1)| = |col(C_2)|$  then move  $ptr_1$  to the row of  $ptr_2$ .

- (IB2.2) If  $|col(C_1)| \neq |col(C_2)|$  then do nothing.
- (IC) Move  $ptr_2$  to the next row in  $\mathcal{R} \setminus \mathcal{R}_2$  above it. Go to (IB).
- (ID) Stop.

Suppose M' is a filling obtained by applying the algorithm  $\gamma_r$  to the filling  $M \in \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$ . Next we show that by applying  $\delta_r^2$  and  $\delta_r^1$  to M' one obtains the filling M.

Steps (IB'1) and (IB'2) clearly invert the steps (C'2) and (C'1), respectively. Immediately after the borrowing step in  $\gamma_r^2$ , by Lemma 5.2, (b) and (c), and Property 4, we can conclude that borrowing has been performed if and only if there is a right 1-cell in  $\mathcal{R}\setminus\mathcal{R}_2$  which forms an NE

chain with a left 1-cell from  $\mathcal{R}_2$  and does not form an NE chain with any left cells in  $\mathcal{R}_1$ . It is clear that step (ID') detects whether there was any borrowing and inverts it if it occurred. So, the first algorithm  $\delta_r^2$  is indeed the inverse of  $\gamma_r^2$ .

In  $\delta_r^1$ , when  $ptr_2$  points to a left 1-cell we need to invert either (B1) or (B2.3). Using Lemma 5.2 (c) one sees that step (IB1) detects exactly which of (B1) and (B2.3) occurred and inverts it. If  $ptr_2$  points to a right cell, it is clear that (IB2.1) and (IB2.2) invert the steps (B2.1) and (B2.2) of  $\gamma_r^1$ . Moreover, if both  $ptr_1$  and  $ptr_2$  point to right 1-cells, and there is no left 1-cell above them, then clearly the algorithm  $\delta_r^1$  leaves them unchanged. That is, the algorithm  $\delta_r^1$  preserves all rows that are above the rows in  $\mathcal{R}_1$ . Therefore, we have

$$M \xrightarrow{\gamma_r} M' \xrightarrow{\delta_r^2 + \delta_r^1} M.$$

This implies that  $\gamma_r : \mathbf{F}(\mathcal{M}, \mathbf{s}; A) \to \mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  is injective. Since  $\mathbf{F}(\mathcal{M}, \mathbf{s}; A)$  is finite, it follows that  $\gamma_r$  is bijective.

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