# Unfolding Simple Chains Inside Circles

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#### Abstract

It is an open problem to determined whether a polygonal chain can be straightened inside a confining region if its links are not allowed to cross. In this paper we propose a special case: whether a polygonal chain can be straightened inside a circle without allowing its links to cross. We prove that this is possible if the straightened configuration can fit within circle. Then we show that these simple chains have just one equivalence class of configurations.

Key words: folding, polygonal chain, reconfiguration

#### 1. Introduction

A chain is a sequence of rigid rods or links consecutively connected at their endpoints, about which they may rotate freely. The link between  $A_{i-1}$  and  $A_i$   $(1 \le i \le n)$  is denoted by  $L_i$  and the length of  $L_i$  is denoted by  $l_i$ . The angle at intermediate joint  $A_i, \theta_i \in [0, 2\pi)$ , is that determined by rotating  $L_i$  about  $A_i$  counterclockwise to bring  $L_i$  to  $L_{i+1}$ . The chain  $\Gamma$  is simple if it is non-selfcrossing and non-self-touching. The subchain of  $\Gamma$ with joints  $A_i, ..., A_j$  is denoted by  $\Gamma[i, j]$ .

We say a *bend operation* is performed at joint  $A_i$ , when the joint angle  $\theta_i$  is changed between  $\theta_i$  and  $\pi$ . Throughout this paper, we assume that the only bend operations allowed are *single-joint* bend operations, in which only one joint angle is altered at a time. A bend operation is *complete* if, at the end of the operation the joint angle is  $\pi$ . We then say that the joint has been *straightened*. A bend Operation that is not complete is called a *partial bend*. A sequence of bend operations is said to be *monotonic* if no operation increases the absolute deviation from straightenes,  $|\theta_i - \pi|$ , for a joint  $A_i$ .

Let  $\sigma = (i_1, i_2, ..., i_{n-1})$  be a permutation of the indices  $\{1, 2, ..., n-1\}$ . For a simple chain  $\Gamma$ , we say that a sequence  $(A_{i_1}, A_{i_2}, ..., A_{i_{n-1}})$  of joints is *unfoldable*, if  $\Gamma$  can be straightened into a straight line segment L using the joints in the sequence in

turn, such that  $\Gamma$  remains simple and all of the bend operations are complete. A simple chain  $\Gamma$  is called *unfoldable chain*, if it has a unfoldable sequence of joints. An intermediate joint  $A_i$  is called *unfoldable joint*, if a complete bend operation can be performed at  $A_i$  such that during the performing bend operation,  $\Gamma$  remains simple.

The union chain,  $\Gamma_U$ , of a chain  $\Gamma$  is a chain which is obtained from  $\Gamma$  in the following way: if none of the joints of  $\Gamma$  is straight joint,  $\Gamma_U = \Gamma$ ; if  $\Gamma$  has at least one straight joint, for any straight joints  $A_i$ , we delete joint  $A_i$  and put  $A_{i-1}A_{i+1}$  as a single link.

Reconfiguration problem and in particular, folding problem, been raised independently by several researchers. [3] has considered reconfiguration of robot arms inside a circle, with allowing its links to cross. In [4], Pei has proved that for a chain  $\Gamma$ inside a circle whose radius is sufficiently big, there is just one equivalence class when its links are allowed to cross. In [5] and [2], straightening a simple chain in the plane is studied and is proved that any simple chain can be straightened in the plane. And in [1] Arkin, Fekete and Mitchell have given an efficient algorithm to determined if a simple chain can be straightened by performing complete bend operations. In this paper, we study straightening a simple chain within a circle. we give a quadratictime algorithm to straighten a simple unfoldable chain within a circle whose radius is sufficiently big. Then we prove that all of simple chains can

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be straightened within a circle, if and only if their straightened configuration can fit in the circle. Finally we show that any two configuration of these simple chains are equivalent.

## 2. Preliminaries

Let  $\Gamma$  be a simple chain inside circle C(O, r) with joints  $A_0...A_n$ . For fitting straightened configuration of  $\Gamma$  in C, we must have  $\sum_{i=1}^{n} l_i \leq 2r$ . From now on, we suppose  $\Gamma$  is a simple chain inside Csuch that  $\sum_{i=1}^{n} l_i < 2r$ .

For a circle C(O, r) and two points  $x, y \in \partial C$ , we use  $\widehat{xy}$  to denote the clockwise arc from x to y. For a point  $A_i \in \partial C$  we denote the other endpoint of the diameter of C which is containing  $A_i$ , by  $M_i$ . **Definition 1** A joint  $A_{r_i}$  is called rim joint if it lies on boundary of circle C. We denote the set of all rim joints of chain  $\Gamma$  by  $A_{Rim} = \{A_{r_0}, A_{r_1}, ..., A_{r_s}\}$ .

**Definition 2** For any rim joint  $A_i$  of chain  $\Gamma$ , the vector  $\overrightarrow{OM}_i$  is called radius vector of  $A_i$  and is denoted by  $\mathbf{r}_i$ .

**Lemma 3** There is a diameter s = ab of circle C such that all of rim joints of chain  $\Gamma$  belong to one of the arcs  $\overrightarrow{ab}$  or  $\overrightarrow{ba}$ .

**PROOF.** If  $A_{Rim} = \emptyset$ , there is nothing to prove. Let  $A_i$  be a rim joint of  $\Gamma$  and X be a moving object which is walking along arc  $A_iM_i$  clockwise, starting at the point  $A_i$ . Suppose  $A_r$  is the last rim joint of  $\Gamma$  that is visited by X. Diameter  $s = A_rM_r$  is a solution. Because arc  $A_rM_r$  contains no rim joint of  $\Gamma$ , except  $A_r$ .  $\Box$ 

**Definition 4** Suppose  $A_{Rim}$  has at least two point and rim joints of  $\Gamma$  belong to ab. The nearest rim joints to points a and b are denoted by  $A_f$  and  $A_e$ , respectively. These joints are called limit-joints.

It is clear that all of the other rim joints of  $\Gamma$  are on arc  $A_f A_e$ .

**Definition 5** Let  $A_e$  and  $A_f$  be limit-joints of  $\Gamma$ . Vectors  $\mathbf{r_e}$  and  $\mathbf{r_f}$  are called direction vectors.

**Definition 6** The sum of direction vectors,  $\mathbf{r}_{e}$  and  $\mathbf{r}_{f}$ , is called central direction and denoted by  $\mathbf{d}$ , *i.e.*,  $\mathbf{d} = \mathbf{r}_{e} + \mathbf{r}_{f}$ .

**Central Translation:** We draw *n* vectors parallel to **d** from any joint  $A_i$  until hit circle at points  $N_i$ , then put  $\varepsilon_i = ||\overrightarrow{A_iN_i}||$  and  $\varepsilon = \min\{\varepsilon_i \mid 0 \le i \le n\}$ . Translation of  $\Gamma$  inside *C* along the vector  $\mathbf{d}_{\varepsilon} = \frac{\varepsilon}{||\mathbf{d}||} \mathbf{d}$ , is called *central translation* of  $\Gamma$ . New positions of  $\Gamma$  and any joint  $A_i$ , after the central translation, are denoted by  $\Gamma'$  and  $A'_i$ . It is clear that  $\varepsilon_e = \varepsilon_f$ .

### 3. Unfoldable Simple Chains

Suppose  $\sigma = (A_{i_1}, A_{i_2}, ..., A_{i_{n-1}})$  is an unfoldable sequence of joints of  $\Gamma$ . For straightening  $\Gamma$ inside circle C(O, r), we propose the following algorithm which contains three steps:

**Algorithm 1** Unfolding Simple Chain  $\Gamma$ :

step 1.  $\Gamma' = \Gamma$ ; j = 0. step 2.  $\Gamma' = \Gamma'_U$ ; j = j + 1. If j = n, stop. else  $k = i_j$ ;

**step 3.** Straighten  $A_k$ . Go to step 2.

Now for step 3, straightening joint  $A_k$  within circle C, we propose the following algorithm which contains four steps:

**Algorithm 2** Straightening joint  $A_k$ :

step 1.  $\Gamma_0 = \Gamma[0, k]; \Gamma_n = \Gamma[k, n];$ 

**step 2.** Rotate  $\Gamma_0$  about  $A_k$  until  $A_k$  straightens or one of joints of  $\Gamma_0$  hits C and  $\Gamma_0$  can not rotate more about  $A_k$ . If  $A_k$  straightens, stop; else go to step 3. **step 3.** Rotate  $\Gamma_n$  about  $A_k$  until  $A_k$  straightens or one of joints of  $\Gamma_n$  hits C and  $\Gamma_n$  can not rotate more about  $A_k$ . If  $A_k$  straightens, stop; else go to step 4.

**step 4.** Calculate  $\mathbf{d}_{\varepsilon}$  and transmit  $\Gamma$  by  $\mathbf{d}_{\varepsilon}$ . Go to step 2.

## 4. Correctness of Algorithm 2

Any repeat of algorithm 2 is called a *phase* and the joint angle at  $A_k$ , at the end of phase *i*, is denoted by  $\alpha_i$ . For showing correctness of algorithm 2, we show that during the algorithm,  $\Gamma$  remains simple and it remains inside *C*. And we prove that by using algorithm 2, after a finite number of repeats,  $A_k$  straightens. Furthermore, this finite number is independent of *n*.

Chain  $\Gamma$  remains simple, because  $A_k$  is an unfold-

able joint in the plane. Now for showing that  $\Gamma$  remains inside C, we first prove that central translation always can be done. Lemma 7  $\mathbf{d}_{\varepsilon} \neq \mathbf{0}$ .

**PROOF.** If  $\mathbf{d} = \mathbf{0}$ , we have  $\mathbf{r_f} = -\mathbf{r_e}$ . That is implies  $A_f = M_e$  and  $A_e = M_f$ , i.e.,  $A_f A_e = 2r$ . Therefore  $\sum_{i=1}^n l_i \geq \sum_{l_i \in \Gamma[e,f]} l_i \geq A_f A_e = 2r$ . That is a contradiction. Thus  $\mathbf{d} \neq \mathbf{0}$ .

Because the angles between **d** and its components are less than  $\pi/2$  and all of radius vectors lie between vectors  $\mathbf{r_e}$  and  $\mathbf{r_f}$ , the angle between **d** and radius vectors are less than  $\pi/2$ , too. Thus any rim joint can transmit in direction **d** inside *C*. Any interior points of *C* also can transmit in all directions inside *C*. Therefore  $\varepsilon \neq 0$ . Consequently  $\mathbf{d}_{\varepsilon} \neq \mathbf{0}$ .  $\Box$ 

It is clear that during the step 1 and step 2, all of the joints remain inside circle. At step 3, because  $\varepsilon = \min{\{\varepsilon_i \mid 0 \le i \le n\}}$  and the angle between radius vectors and **d** are less than  $\pi/2$ ,  $\Gamma$  remains inside C.

Now to show that after a finite number of repeats, algorithm 2 is terminated, we first show that at the end of any phase of algorithm 2,  $\alpha_i$  becomes strongly close to  $\pi$ , i.e.,  $|\pi - \alpha_{i+1}| < |\pi - \alpha_i|$ . So we have to prove at the end of central translation of  $\Gamma$ , at least one of the subchains  $\Gamma_0$  or  $\Gamma_n$ , can rotate about  $A_k$  such that joint angle at  $A_k$  has became close to  $\pi$ . Note that at the end of step 1 and step 2, if  $A_k$  does not straighten,  $A_{Rim}$  has at least two points, one point from  $\Gamma_0$  and the other point from  $\Gamma_n$ .

**Lemma 8** Let  $A_e$  and  $A_f$  be the limit-joints of  $\Gamma$  at phase *i*. If both of  $A_e$  and  $A_f$  belong to one of the subchains  $\Gamma_0$  or  $\Gamma_n$ , then at the end of translation, none of the joints of the other subchain lie on  $\partial C$ . Furthermore, this subchain can rotate about  $A_k$  at phase i + 1.

**PROOF.** Assume without loss of generality that  $A_f, A_e \in \Gamma_0$ . Suppose for a contradiction,  $A_t$  is a joint of  $\Gamma_n$  such that  $A'_t \in \partial C$ . At the beginning of translation,  $\Gamma_n$  has at least one rim joint,  $A_m$ , which lies on arc  $A_fA_e$ . If  $A_k$  is in the exterior of closed curve  $\delta = A_fA_e \cup \Gamma_0[e, f], \Gamma_n[m, k]$  and  $\Gamma_0[e, f]$  will be intersecting. Thus  $A_k$  is in the interior of  $\delta$ . See figure 1. Therefore  $A'_k$  is in the interior of the closed curve  $\delta' = \beta \cup \Gamma'_0[f, e]$  where

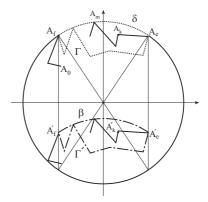


Fig. 1. If  $\Gamma_n$  contains no limit-joints,  $\Gamma'_n$  has no rim joint.

 $\beta$  is the translation of arc  $A_f A_e$  by the vector  $\mathbf{d}_{\varepsilon}$ . But  $A'_t$  is in the exterior of  $\delta'$ . So  $\Gamma'_n[k, t]$  intersects boundary of  $\delta'$ . That is a contradiction. Because  $\Gamma_n[k, t]$  does not intersect boundary of  $\delta$ .  $\Box$ 

By lemma 8, we suppose  $A_e$  and  $A_f$  don't belong to the same subchain. From now on, the subchain which contains  $A_e$  is denoted by  $\Gamma_e$  and the other subchain which contains  $A_f$  is denoted by  $\Gamma_f$ . We have the following theorem.

**Theorem 9** At the end of translation, at least one of the subchains  $\Gamma_e$  and  $\Gamma_f$  can rotate about  $A_k$ .

**PROOF.** Refer to full paper.  $\Box$ 

**Corollary 10**  $|\alpha_i - \pi| > |\alpha_{i+1} - \pi|.$ 

By corollary 10, the configuration of  $\Gamma$  in two consecutive phase is different. Thus during the algorithm 2, straightening  $A_k$  is strongly progressed and cycling is not possible. Now for showing that after a finite number of repeats, algorithm 2 is terminated, we use simplicity of  $\Gamma$ . Assume without loss of generality that  $\theta_k < \pi$ . Thus according to definition of joint angle,  $\Gamma_0$  must rotate about  $A_k$ clockwise. First suppose there is no confining region. So  $A_k$  can be straightened and then  $\Gamma_0$  can rotate about  $A_k$  clockwise more, until first selftouching is occurred. This operation is called  $\pi$ passage motion and the joint angle at  $A_k$  is denoted by  $\pi + \tau_k$ , that  $\tau_k > 0$ . Now suppose  $\Gamma$  is inside C(O, r). We change the stopping criteria of algorithm 2, from achieving  $\pi$  to achieving  $\pi + \tau_k$  and use this new algorithm on  $A_k$ . All of above proofs also hold for this new algorithm. So by corollary 10 we also have:

$$|\alpha_{i+1} - \pi - \tau_k| < |\alpha_i - \pi - \tau_k|$$
 (\*)

Assumption  $\theta_k < \pi$  yields: for every  $i \ge 0$ ,  $\alpha_i \le \pi + \tau_k \cdot So(*)$  yields  $\pi + \tau_k - \alpha_{i+1} < \pi + \tau_k - \alpha_i$ . In the other words,  $\{\alpha_i\}_{i\ge 0}$  is a bounded and monotone sequence. Therefore it converges to its suprimum,  $\pi + \tau_k$ . Thus for every  $\varepsilon > 0$  exists a finite natural number N > 0 such that for every  $i \ge N$  we have  $|\alpha_i - \pi - \tau_k| < \varepsilon$ , i.e., for every  $i \ge N$ ,  $\pi + \tau_k - \alpha_i < \varepsilon$ . Thus for  $\varepsilon = \tau_k$ , there is a finite number  $N_\tau$  such that for all  $i \ge N_\tau$ ,  $\pi + \tau_k - \alpha_i < \tau_k$ . So for  $i = N_\tau$ , we have  $\pi + \tau_k - \alpha_{N_\tau} < \tau_k$ , i.e.,  $\alpha_{N_\tau} > \pi$ . Because  $N_\tau$  is the smallest natural number that  $\pi + \tau_k - \alpha_i < \tau_k$ , we have  $\pi + \tau_k - \alpha_{N_\tau - 1} \ge \tau_k$ , i.e.,  $\alpha_{N_\tau - 1} \le \pi$ . Therefore  $A_k$  can straighten in phase  $N_\tau$  or  $N_\tau - 1$ . Because  $\alpha_{N_\tau - 1} \le \pi$  and  $\alpha_{N_\tau} > \pi$ . It is clear that  $N_\tau$  is independent of n.

Therefore proof of correctness of algorithm 2 is terminated. Complexity of algorithm 1 is  $O(n^2)$ , because complexity of each step is O(n) and the number of repeats is n-1.

## 5. Arbitrary Simple Chains

Now we prove that an arbitrary simple chain can be straightened inside a circle. First we have the following theorem.

**Theorem 11** Any simple chain  $\Gamma$  can be straightened using a finite number of monotonic singlejoint bend operations.

**PROOF.** See [1] and [2].  $\Box$ 

Theorem 11 is true, if chain is inside a circle as a confining region.

**Theorem 12** A simple chain  $\Gamma$  can be straightened inside a circle using a finite number of mono-

tonic single-joint bend operations, if  $\sum_{i=0}^{n} l_i < 2r$ .

**PROOF.** By theorem 11,  $\Gamma$  can be straightened in the plane using a finite number of monotonic single-joint bend operations. If all of these bend operations are complete,  $\Gamma$  can be straightened by using algorithm 1. But if at least one of the bend operations is not complete, these bend operations will be in accordance with a sequence of motions,  $M = \{M_j\}_{j=1}^k$ , such that  $M_j$  is a partial bend operation and  $\Gamma$  can be straightened by using M. Any operation  $M_j$  is single-joint, so it is in accordance with a joint  $A_{i_j}$  and this accordance is not one to one, because  $M_j$  s are not complete. Suppose any bend operation  $M_j$  is changed joint angle at  $A_{i_j}$  by  $\Delta(A_{i_j}; j)$ . Now note that any operation  $M_j$  is monotone; so if we change the stopping criteria of algorithm 2, from achieving  $\pi$  to achieving  $\theta_{i_j} + \Delta(A_{i_j}; j)$ , this new algorithm can be used to perform any bend operation  $M_j$  inside C. Therefore  $\Gamma$  can be straightened inside C by performing  $M_j$  s in turn, because k is finite.  $\Box$ 

## 6. Conclusion

Assume that  $\Gamma$  is a simple chain such that  $\sum_{i=1}^{n} l_i < 2r$  and  $\Gamma_1$  and  $\Gamma_2$  are two configuration of  $\Gamma$  inside circle C(O, r). We denote their straight configurations by  $L_1$  and  $L_2$ , respectively. Let M be a sequence of bend operations for straightening  $\Gamma_2$  inside C and  $M^R$  be the reverse of Motion M. It is clear that by performing  $M^R$ ,  $L_2$  can be reconfigured to  $\Gamma_2$ . Now by theorem 11, we can reconfigure  $\Gamma_1$  to  $L_1$ , then we can reconfigure  $L_1$  to  $L_2$  by translation and rotation operations and finally we can reconfigure  $L_2$  to  $\Gamma_2$  by  $M^R$ . So  $\Gamma_1$  can be reconfigured to  $\Gamma_2$  inside C, i.e., for a simple chain  $\Gamma$ , if  $\sum_{i=1}^{n} l_i < 2r$ , then any two configurations

of  $\Gamma$ , inside circle C(O, r) are equivalent.

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