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## ON THE RIGIDITY OF ONE-DIMENSIONAL SYSTEMS OF CONTRACTION SIMILITUDES

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Associated family of similitudes  $\mathfrak{F}(S)$  for a system  $S$  of contraction similitudes in  $\mathbb{R}^d$  was primarily introduced by C.Bandt and S.Graf in their work [1] as a tool for checking, whether the invariant set  $K(S)$  of the system  $S$  has positive Hausdorff measure. This approach was developed in [3],[5],[2], leading to formulation of weak separation property (WSP) and it's application to graph-directed systems.

Nevertheless, thorough consideration of the associated family of similitudes allows to throw light upon some other properties of self-similar sets. If the identity map is a limit point for the associated family  $\mathfrak{F}(S)$ , the system  $S$  demonstrates the rigidity phenomenon. We regard it in the case when the invariant set  $K(S)$  of the system  $S$  is the segment  $[0, 1]$ . Our main result is that in this situation any continuous map  $\varphi : K(S) \rightarrow K(T)$  of the attractor of the system  $S$  to the attractor of the other system of contraction similitudes  $T$  which agrees with the structure of self-similar set on the sets  $K(S)$  and  $K(T)$  is a linear map of  $[0, 1]$  to a straight line segment.

**1. Definitions and notation.** A compact set  $K = K(S)$  in  $\mathbb{R}^d$  is called an *attractor* or *invariant set* of the system  $S = \{S_1, \dots, S_m\}$  of contraction similitudes in  $\mathbb{R}^d$  if  $K = S_1(K) \cup \dots \cup S_m(K)$ . Existence and uniqueness of the set  $K$  follow from theorem of Hutchinson ([4, стр. 724]).

We denote by  $I = \{1, \dots, m\}$  a set of first  $m$  natural numbers, by  $I^k$  – the set of *multiindices* of length  $k$  and  $I^* = \bigcup_{k=1}^{\infty} I^k$ . Given a system  $S = \{S_1, \dots, S_m\}$  of maps of  $\mathbb{R}^d$  to itself, for each multiindex  $\mathbf{i} \in I^k$ ,  $\mathbf{i} = i_1 i_2 \dots i_k$ , we define  $S_{\mathbf{i}} = S_{i_1} \cdot S_{i_2} \cdot \dots \cdot S_{i_k} : \mathbb{R}^d \rightarrow \mathbb{R}^d$ .

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We denote by  $\text{Lip}(g)$  the expansion ratio of the similitude  $g$ , and by  $\text{fix}(g)$  we denote it's fixed point. If  $g$  is a translation we put  $\text{fix}(g) = \infty$ .

**1.1 The associated family.** A family of all transformations having the form  $S_i^{-1} \cdot S_j : \mathbb{R}^d \rightarrow \mathbb{R}^d$ , where  $\mathbf{i} \in I^*$ ,  $\mathbf{j} \in I^*$  and  $i_1 \neq j_1$  we call, following [1, p.996], the *associated family*  $\mathfrak{F}(S)$  of similitudes of the system  $S$ .

**1.2 Structure-preserving homeomorphisms.** Let  $S = \{S_1, \dots, S_m\}$  and  $T = \{T_1, \dots, T_m\}$  be two systems of contraction maps of complete metric spaces  $X_1$  and  $X_2$  to themselves. Let  $K(S) \subset X_1$  and  $K(T) \subset X_2$  be the attractors of these two systems. A continuous map  $\varphi : K(S) \rightarrow K(T)$  is said to be *structure-preserving*, if

$$(1) \quad \forall x \in K(S) \quad \forall i \in I \quad \varphi(S_i(x)) = T_i(\varphi(x))$$

It follows immediately from this definition, that for any multiindex  $\mathbf{j} \in I^*$ , and for any  $x \in [0, 1]$  the relation  $\varphi(S_{\mathbf{j}}(x)) = S_{\mathbf{j}}(\varphi(x))$  is also valid. The same is also true for the elements of associated families:

**Proposition 1.** *Let  $\varphi : K(S) \rightarrow K(T)$  - be a structure-preserving map of the attractors of systems  $S$  and  $T$ . Let  $g = S_i^{-1} \cdot S_j$  and  $h = T_i^{-1} \cdot T_j$  - be the elements of  $\mathfrak{F}(S)$  and  $\mathfrak{F}(T)$ . The relation  $\varphi \cdot g(x) = h \cdot \varphi(x)$  holds for any  $x \in K(S) \cap g^{-1}(K(S))$ .*

Two systems  $S = \{S_1, \dots, S_m\}$  and  $T = \{T_1, \dots, T_m\}$  are called *structurally equivalent*, if there is a structure-preserving homeomorphism  $\varphi : K(S) \rightarrow K(T)$ .

**2. The main results.**

The first statement belongs to the case when each of the invariant sets is  $[0,1]$ :

**Theorem 1.** *Let  $S = \{S_1, \dots, S_m\}$ ,  $T = \{T_1, \dots, T_m\}$  be systems of contraction similitudes in  $\mathbb{R}$ , invariant set of each being the segment  $[0, 1]$ , and let  $\varphi : K(S) \rightarrow K(T)$  be a structure-preserving homeomorphism for these two systems, such that  $\varphi(0) = 0$  and  $\varphi(1) = 1$ . If  $\text{Id}$  is a limit point of the associated family  $\mathfrak{F}(S)$  for the system  $S$ , then  $\varphi(x) \equiv x$ , and  $S = T$ .*

The scheme of the proof is the following:

Let  $g_n = S_i^{-1} S_j$  be such a sequence of elements of the family  $\mathfrak{F}(S)$ , that  $g_n \rightarrow \text{Id}$  (and  $g_n([0, 1]) \cap [0, 1] \neq \emptyset$ ), and let  $h_n = T_i^{-1} T_j$  be a sequence of elements of  $\mathfrak{F}(S)$ , which are conjugate to  $g_n$  via the homeomorphism  $\varphi$  in the sense of Proposition 1. We prove that this pair of conjugate sequences may be modified to satisfy the following conditions:

1. Each map  $g_n, h_n$  preserves orientation on  $\mathbb{R}$ .
2. Either (A): all similitudes  $g_n$  are translations  $g_n(x) = x + \Delta_n$ , where  $\Delta_n > 0$ , or (B): the expansion ratios  $p_n = \text{Lip}(g_n) \neq 1$ ,  $g_n(0) \geq 0$  and the sequence of points  $\text{fix}(g_n)$  converges to infinity;
3. Either (I): all similitudes  $h_n$  are translations  $h_n(x) = x + \delta_n$ ,  $\delta_n > 0$ , or (II): the expansion ratios  $q_n = \text{Lip}(h_n) \neq 1$ ,  $h_n(0) \geq 0$  and the sequence of points  $\text{fix}(h_n)$  converges to infinity;
4. In the case (,II) there is  $\lim_{n \rightarrow \infty} \log_{p_n} q_n = \alpha$ ;
5. For any  $n$ ,  $g_n(0) > 0, h_n(0) > 0$ .

Let  $\Gamma(\varphi) \subset [0, 1] \times [0, 1]$  be the graph of the homeomorphism  $\varphi$ . Take some  $\varepsilon \in (0, 1/4)$ .

For  $n$  sufficiently large, the points  $(g_n^k(0), h_n^k(0))$ , where  $k$  runs from 0 to  $M_n = \max\{k : g_n^k(0) \leq 1\}$ , form an  $\varepsilon$ -net in the set  $\Gamma(\varphi)$ . At the same time these points belong to an arc  $\gamma$ , specified by an equation  $y = \psi(x), x \in [0, 1]$ , where:

$\psi(x) = \frac{\Delta_n}{\delta_n}x$  in the case (AI);

$\psi(x) = L(e^{\lambda x} - 1)$ , where  $\lambda = \frac{\log q_n}{\Delta_n}$ ,  $L = -\text{fix}(h_n)$  in the case (AII);

$\psi(x) = L' \left( 1 - \left( 1 - \frac{x}{L} \right)^\lambda \right)$ ,  $\lambda = \frac{\log q_n}{\log p_n}$ ,  $L = -\text{fix}(g_n)$ ,  $L' = -\text{fix}(h_n)$  in the case (BII).

In each of these three cases we prove that for  $n$  sufficiently large, Hausdorff distance between  $\gamma$  and a segment  $C = \{y = x, x \in [0, 1]\}$ , is not greater than  $\varepsilon$ . Since  $\varepsilon$  may be arbitrarily small,  $\gamma$  and  $C$  (and therefore  $\Gamma(\varphi)$  and  $C$ ) coincide, hence  $\varphi(x) \equiv x$ .

The second statement treats structure-preserving homeomorphisms  $\varphi : K(S) \rightarrow K(T)$  in the case, when  $K(S) = [0, 1]$ , and  $K(T)$  is a Jordan arc.

**Theorem 2.** *Let  $S = \{S_1, \dots, S_m\}$  be a system of contraction similitudes in  $\mathbb{R}^d$ , whose attractor is segment  $[0, 1]$ . Let Jordan arc  $\gamma \in \mathbb{R}^d$  with endpoints  $a_0, a_1$  is the attractor of a system  $T = \{T_1, \dots, T_m\}$  of contraction similitudes in  $\mathbb{R}^d$ . Let  $\varphi : [0, 1] \rightarrow \gamma$  be a structure-preserving homeomorphism for the systems  $S$  and  $T$ .*

*If  $Id$  is a limit point for the family  $\mathfrak{F}(S)$ , then the arc  $\gamma$  is a straight line segment with endpoints  $a_0, a_1$ .*

The method of the proof of this theorem is similar to the previous one. Assuming, that the affine hull of  $\gamma$  is  $\mathbb{R}^d$ , we build such sequence  $g_n \in \mathfrak{F}(S)$ , that for it and for the sequence  $h_n \in \mathfrak{F}(T)$ , conjugate to  $g_n$  via homeomorphism  $\varphi$ , the following conditions are satisfied:

1.  $g_n$  and  $h_n$  preserve orientation in  $[0, 1]$  and  $\gamma$  correspondingly;
2.  $g_n(0) > 0$ ;
3. a sequence of distances  $\Delta_n$  from the point  $a_0$  to the closest proper strictly invariant affine codimension 2 subspace  $V_n$  of similitude  $h_n$  converges to infinity.

Take  $\varepsilon > 0$ . For  $n$  sufficiently large, the points  $h_n^k(a_0)$ , where  $k$  runs from 0 to  $M_n = \max\{k : g_n^k(0) \leq 1\}$  form an  $\varepsilon$ -net in  $\gamma$ . At the same time they all are contained in an arc of a loxodrome  $L_n = \{h_n^t(a_0), t \in [0, M_n]\}$ . By virtue of condition 3,  $n$  may be chosen so that Hausdorff distance between the arc  $L_n$  and a straight line segment  $[a_0, a_1]$  is no greater than  $\varepsilon$ . Since  $\varepsilon$  may be arbitrarily small, this means that  $\gamma$  is equal to the segment  $[a_0, a_1]$ .

Using the result of Theorem 2 and slightly modifying the argument in the proof of Theorem 3, we obtain:

**Theorem 3.** *Let  $S = \{S_1, \dots, S_m\}$  be a system of contraction similitudes in  $\mathbb{R}^d$ , whose attractor is segment  $[0, 1]$ . Let a continuum  $\gamma \in \mathbb{R}^d$  be the attractor of a system  $T = \{T_1, \dots, T_m\}$  of contraction similitudes in  $\mathbb{R}^d$ . Let  $\varphi : [0, 1] \rightarrow \gamma$  be a structure-preserving map of the attractors of the systems  $S$  and  $T$ .*

*If the identity map  $Id$  is a limit point for the family  $\mathfrak{F}(S)$ , then  $\gamma$  is a straight line segment, and  $\varphi$  is a linear map.*

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