

# Cluster algebras and snake modules

Bing Duan

(Lanzhou University and University of Connecticut)

joint with Jian-Rong Li & Yan-Feng Luo

A conference celebrating the 60th birthday of  
Prof. Vyjayanthi Chari

The Catholic University of America, Washington, D.C., June 7, 2018

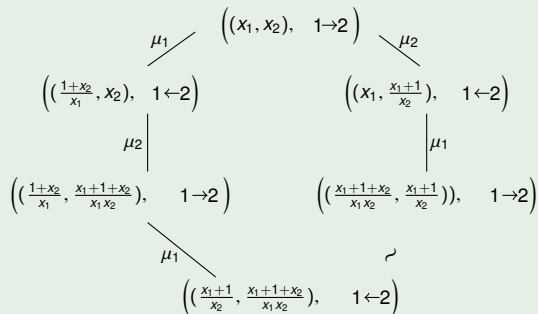
# Cluster algebras

Cluster algebras were introduced by Fomin and Zelevinsky around 2000.

## Example

Q: 1→2

$$\begin{pmatrix} 0 & 1 \\ -1 & 0 \end{pmatrix}$$



$$\mathcal{A}(Q) = \mathbb{Z}[x_1, x_2, \frac{1+x_2}{x_1}, \frac{x_1+1}{x_2}, \frac{x_1+x_2+1}{x_1 x_2}] \subseteq \mathbb{Q}(x_1, x_2)$$

# Representations of quantum affine algebras

- $\mathfrak{g}$  simple Lie algebra over  $\mathbb{C}$
- $I = \{1, 2, \dots, n\}$ , where  $n$  is the rank of  $\mathfrak{g}$
- $U_q(\hat{\mathfrak{g}})$  (untwisted) quantum affine algebra with quantum parameter  $q \in \mathbb{C}^*$  not a root of unity

Chari and Pressley classified the finite dimensional simple  $U_q(\hat{\mathfrak{g}})$ -modules in terms of Drinfeld polynomials, i.e.,  $I$ -tuples of polynomials  $(P_1, \dots, P_n)$ , where  $P_i, i = 1, \dots, n$ , are polynomials in one indeterminate  $u$  with coefficients in  $\mathbb{C}$  and constant term 1.

For example, in  $A_n$ ,

- $(1, \dots, 1, 1 - au, 1, \dots, 1)$  is a Drinfeld polynomial of a fundamental module
- $(1, \dots, 1, (1 - au)(1 - aq^2u), 1, \dots, 1)$  is a Drinfeld polynomial of a Kirillov-Reshetikhin module
- $(1, \dots, 1, (1 - au)(1 - aq^2u), (1 - aq^5u)(1 - aq^7u), 1, \dots, 1)$  is a Drinfeld polynomial of a minimal affinization.

Minimal affinizations were introduced by Chari in 1995.

Let  $\mathcal{P}$  be the free abelian multiplicative group of monomials in infinitely many formal variables  $(Y_{i,a})_{i \in I, a \in \mathbb{C}^*}$  and  $\mathcal{P}^+ \subseteq \mathcal{P}$  denote the set of all dominant monomials. Then

$$\begin{aligned} \mathcal{P}^+ &\longleftrightarrow \{\text{f. d. simple } U_q(\hat{\mathfrak{g}})\text{-modules}\} \\ m &\longmapsto L(m). \end{aligned}$$

Example, in  $A_n$ ,

Drinfeld polynomials	Dominant monomials	simple $U_q(\hat{\mathfrak{g}})$ -modules
$(1, \dots, 1, 1 - au, 1, \dots, 1)$	$Y_{i,a}$	$L(Y_{i,a})$
$(1, \dots, 1, (1 - au)_i(1 - aq^2u), 1, \dots, 1)$	$Y_{i,a} Y_{i,aq^2}$	$L(Y_{i,a} Y_{i,aq^2})$
$(1, \dots, 1, (1 - au)_i(1 - aq^2u), (1 - aq^5u)_{i+1}(1 - aq^7u), 1, \dots, 1)$	$Y_{i,a} Y_{i,aq^2} Y_{i+1,aq^5} Y_{i+1,aq^7}$	$L(Y_{i,a} Y_{i,aq^2} Y_{i+1,aq^5} Y_{i+1,aq^7})$
	$Y_{1,aq^{-13}} Y_{2,aq^{-10}} Y_{3,aq^{-7}} Y_{2,aq^{-4}} Y_{1,aq^{-1}}$	$L(Y_{1,aq^{-13}} Y_{2,aq^{-10}} Y_{3,aq^{-7}} Y_{2,aq^{-4}} Y_{1,aq^{-1}})$

Snake modules were introduced by Mukhin and Young in 2012. Snake modules corresponds to ladder representations in the setting of representations of p-adic groups. Ladder representations were introduced by Lapid and Minguez in 2014.

Kirillov–Reshetikhin modules  $\subset$  minimal affinizations  $\subset$  prime snake modules  $\subset$  snake modules

Frenkel and Reshetikhin introduced the theory of  $q$ -characters. The  $q$ -character is an injective ring homomorphism from Grothendick ring of the category of f. d. simple  $U_q(\hat{\mathfrak{g}})$ -modules to Laurent polynomial ring  $\mathbb{Z}[Y_{i,a}^{\pm}]_{i \in I, a \in \mathbb{C}^*}$ . For example,  $\mathfrak{g} = \mathfrak{sl}_2$ ,

$$\chi_q(L(Y_{1,a})) = Y_{1,a} + Y_{1,aq^2}^{-1}.$$

Frenkel-Mukhin algorithm can be used to compute the  $q$ -characters of a large family of modules. For example, the modules whose  $q$ -characters have only one dominant monomial.



The simple module

$$L(Y_{i_T, aq^{k_T}} \cdots Y_{i_2, aq^{k_2}} Y_{i_1, aq^{k_1}})$$

is called a **snake module/prime snake module/minimal snake module** if for all  $2 \leq t \leq T$ , the point  $(i_t, k_t)$  is in **snake position/prime snake position/minimal snake position** with respect to  $(i_{t-1}, k_{t-1})$ .

A point  $(i', k')$  is said to be in **snake position** (respectively, **prime snake position**) with respect to  $(i, k)$  if

Type  $A_n$  :  $k' - k \geq |i' - i| + 2$  and  $k' - k \equiv |i' - i| \pmod{2}$   
(respectively,  $\min\{2n + 2 - i - i', i + i'\} \geq k' - k \geq |i' - i| + 2$  and  $k' - k \equiv |i' - i| \pmod{2}$ ).

The point  $(i', k')$  is in **minimal snake position** to  $(i, k)$  if  $k' - k$  is equal to the given lower bound.

# Motivations

- Hernandez-Leclerc Conjecture: a certain subcategory  $C_\ell$  of the category of all finite dimensional  $U_q(\hat{\mathfrak{g}})$ -modules has a cluster algebra structure. In summer school, Leclerc introduced the conjecture and listed known results.
- The fact that prime snake modules are prime was proved by Mukhin and Young.

## Our results

We proved the following result.

### Theorem

*Prime snake modules of type  $A_n, B_n$  are real.*

Idea of the proof: we use "path description of q-characters" introduced by Mukhin and Young in 2012.

### Remark

*Prime snake modules of type  $A_n$  are real also follows from a recent result of Lapid and Minguez in 2017. They proved (in the language of representations of  $p$ -adic groups) that a large family of  $U_q(\hat{\mathfrak{g}})$ -modules (containing prime snake modules) of type  $A_n$  are real.*

Our main results are

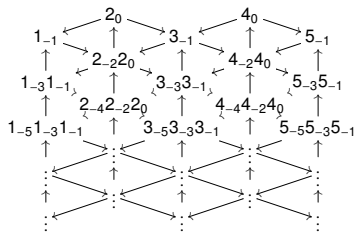
### Theorem

- (1) *Prime snake modules can be obtained using a sequence of mutations from the initial cluster of the cluster algebras constructed by Hernandez and Leclerc.*
- (2) *We obtained a system of equations satisfied by  $q$ -characters of prime snake modules. The system of equations contains all prime snake modules and only contains prime snake modules. It is a natural generalization of  $T$ -systems.*

I will show how to find mutation sequences.

## Mutation sequences and some examples

For every finite dimensional Lie algebra  $\mathfrak{g}$ , Hernandez and Leclerc constructed a cluster algebra with an infinite quiver  $Q$ , for example, in type  $A_5$ , see Figure 1.

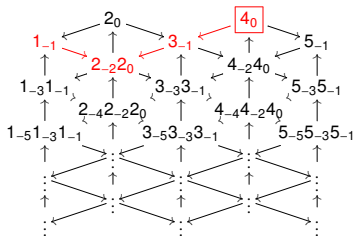


We denote  $4_{-2}4_0 = L(Y_{4,aq^{-2}} Y_{4,a})$ .

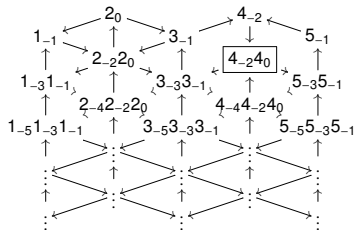
Figure: Quiver  $A_5$

How to get

- $L(2_{-4}2_0)$ , it is a non-minimal prime snake module, and
- $L(3_{-7}2_{-4}2_0)$ .

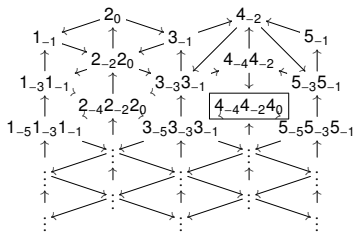


$$[4_0][4_{-2}] = [4_{-2}4_0] + [3_{-1}][5_{-1}]$$

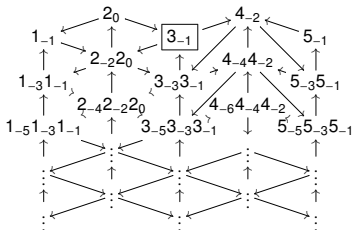


$$[4_{-2}4_0][4_{-4}4_{-2}] = [4_{-2}][4_{-4}4_{-2}4_0] + [3_{-3}3_{-1}][5_{-3}5_{-1}]$$

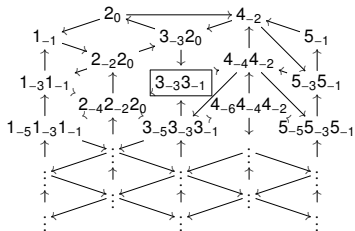




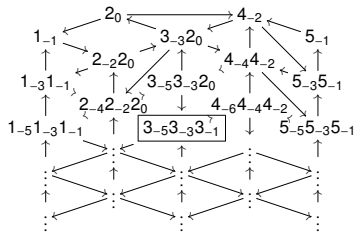
$$[4_{-4}4_{-2}4_0][4_{-6}4_{-4}4_{-2}] = [4_{-4}4_{-2}][4_{-6}4_{-4}4_{-2}4_0] \\ + [3_{-5}3_{-3}3_{-1}][5_{-5}5_{-3}5_{-1}]$$



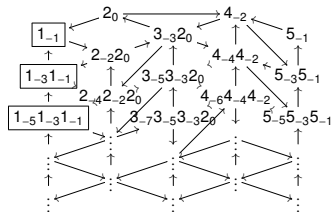
$$[3_{-1}][3_{-3}2_0] = [2_0][3_{-3}3_{-1}] + [2_{-2}2_0][4_{-2}]$$



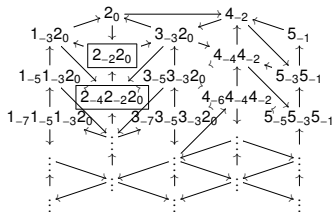
$$[3_{-3}3_{-1}][3_{-5}3_{-3}2_0] = [3_{-3}2_0][3_{-5}3_{-3}3_{-1}] + [2_{-4}2_{-2}2_0][4_{-4}4_{-2}]$$



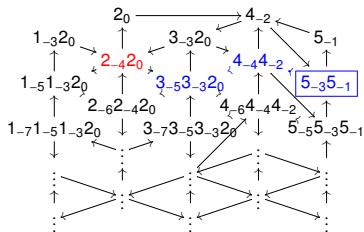
$$[3_{-5}3_{-3}3_{-1}][3_{-7}3_{-5}3_{-3}2_0] = [3_{-5}3_{-3}2_0][3_{-7}3_{-5}3_{-3}3_{-1}] + [2_{-6}2_{-4}2_{-2}2_0][4_{-6}4_{-4}4_{-2}]$$



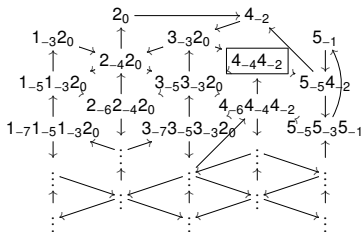
$$\begin{aligned}
 [1_{-1}][1_{-3}2_0] &= [2_0][1_{-3}1_{-1}] + [2_{-2}2_0], \\
 [1_{-3}1_{-1}][1_{-5}1_{-3}2_0] &= [1_{-3}2_0][1_{-5}1_{-3}1_{-1}] + [2_{-4}2_{-2}2_0], \\
 [1_{-5}1_{-3}1_{-1}][1_{-7}1_{-5}1_{-3}2_0] &= [1_{-5}1_{-3}2_0][1_{-7}1_{-5}1_{-3}1_{-1}] + [2_{-6}2_{-4}2_{-2}2_0]
 \end{aligned}$$



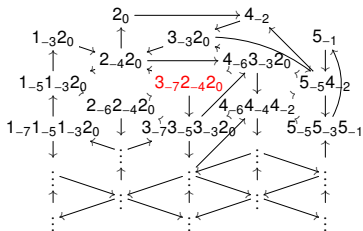
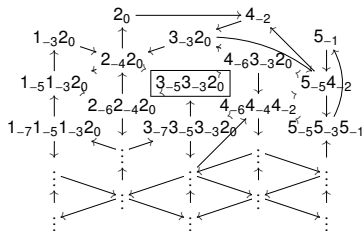
$$\begin{aligned}
 [2_{-2}2_0][2_{-4}2_0] &= [2_{-4}2_{-2}2_0][2_0] + [1_{-3}2_0][3_{-3}2_0], \\
 [2_{-4}2_{-2}2_0][2_{-6}2_{-4}2_0] &= [2_{-4}2_0][2_{-6}2_{-4}2_{-2}2_0] + [1_{-5}1_{-3}2_0][3_{-5}3_{-3}2_0]
 \end{aligned}$$



$$[5_{-3}5_{-1}][5_{-5}4_{-2}] = [5_{-5}5_{-3}5_{-1}][4_{-2}] + [4_{-4}4_{-2}].$$



$$[4_{-4}4_{-2}][4_{-6}3_{-3}2_0] = [3_{-3}2_0][4_{-6}4_{-4}4_{-2}] + [5_{-5}4_{-2}][3_{-5}3_{-3}2_0]$$



$$[3_{-5}3_{-3}2_0][3_{-7}2_{-4}2_0] = [3_{-7}3_{-5}3_{-3}2_0][2_{-4}2_0] + [4_{-6}3_{-3}2_0][2_{-6}2_{-4}2_0].$$

# S-systems

We obtained a series of exchange relations in the process of mutations. They form a system of equations which we called **S-systems**, which is a natural generalization of *T*-systems. These equations in *S*-systems are of the form

$$[S_1][S_2] = [S_3][S_4] + [S_5][S_6],$$

where  $S_i$ ,  $i = 1, \dots, 6$ , are certain prime snake modules. For example:

- $$[L(3_{-5}3_{-3}2_0)][L(3_{-7}2_{-4}2_0)] = [L(3_{-7}3_{-5}3_{-3}2_0)][L(2_{-4}2_0)] + [L(4_{-6}3_{-3}2_0)][L(2_{-6}2_{-4}2_0)]$$

# Thank you !