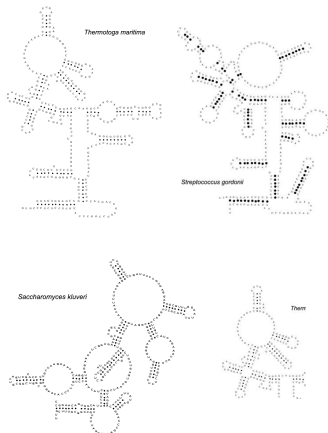


Pourquoi comparer des ARN ?



• A quel point ces structures sont-elles similaires (ou différentes ?)

→ classification

→ phylogénie

• Quelles parties des deux structures se ressemblent le plus ?

• La petite est-elle similaire à une partie de la grande ?

Comparaison → score + correspondance entre les structures

17/01/2010

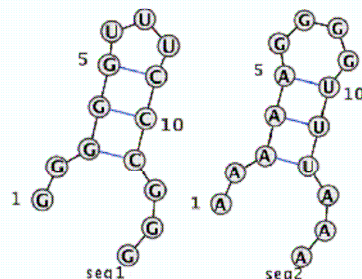
Projet Braserio - ANR Blanc 2006

11



Comparer les séquences ne suffit pas

- La fonction dépend de la structure.
- Des séquences différentes peuvent avoir la même structure.



Séquences :

[seq1] - ---GCGUUUCCCGGG
[seq2] - AAAAAAGGGUUU---AAA

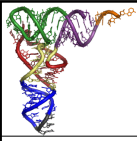
Séquences/structures :

[seq1] - ..(((.....)))...
[seq1] - GGGGG-UUUCCCGGG
[seq2] - AAAGGGGUUUAAA
[seq2] - ..(((.....)))...

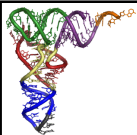
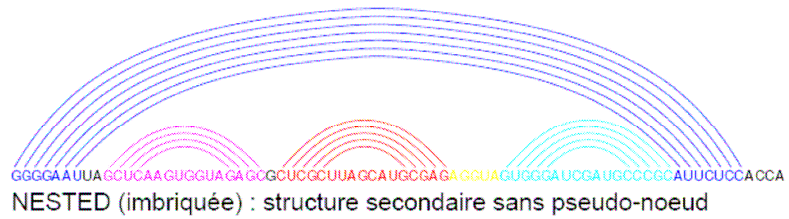
17/01/2010

Projet Braserio - ANR Blanc 2006

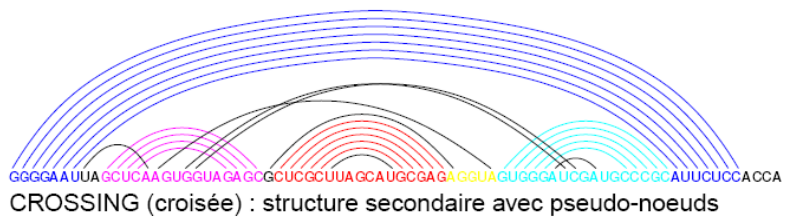
12

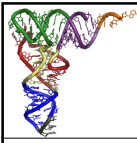


Séquences arc-annotées

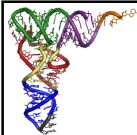
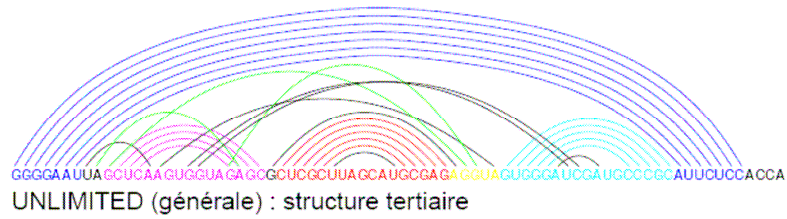


Séquences arc-annotées





Séquences arc-annotées



Séquences arc-annotées

GGGGAAUUA GCUCAA GUGGUA GAGCUCGCUUAGCAUGCAGAGGUAUGGGGAUCGAUGCCCGCAUUCUCCACCA

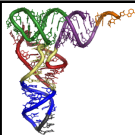
PLAIN (sans arcs) : séquence sans appariement



Opérations d'édition



- Opérations on bases:
 - Substitution: $A \rightarrow C$
 - Deletion / Insertion: $A \rightarrow \emptyset$
- Operations on arcs:
 - Arc-substitution: $\overset{\text{C}}{\text{C}} \overset{\text{G}}{\text{G}} \rightarrow \overset{\text{U}}{\text{U}} \overset{\text{A}}{\text{A}}$
 - Arc-deletion / Arc-insertion: $\overset{\text{C}}{\text{C}} \overset{\text{G}}{\text{G}} \rightarrow \emptyset$
 - Arc-breaking / : $\overset{\text{C}}{\text{C}} \overset{\text{G}}{\text{G}} \rightarrow \text{C} \text{ G}$
 - Arc-altering / : $\overset{\text{C}}{\text{C}} \overset{\text{G}}{\text{G}} \rightarrow \text{C} -$



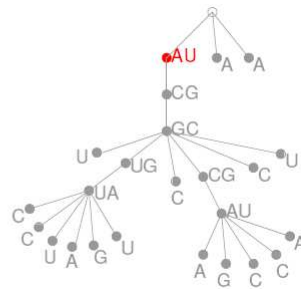
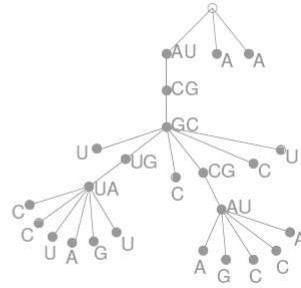
Une édition de deux structures

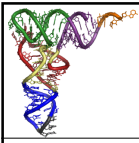




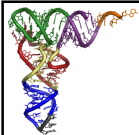
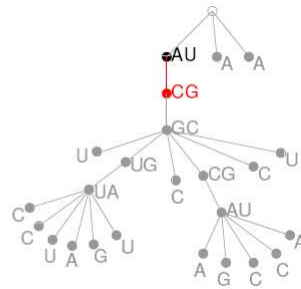
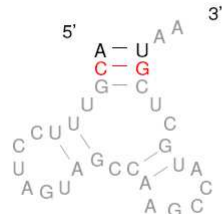
- Jiang, Lin, Ma, Zhang 2002
- Blin, Fertin, Rusu, Sinoquet 2003
- Crochemore, Landau, Ziv-Ukelson 2002



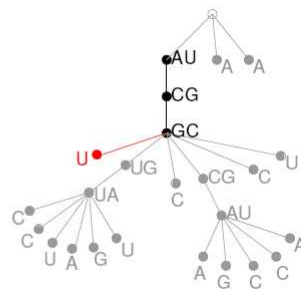
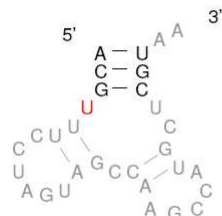


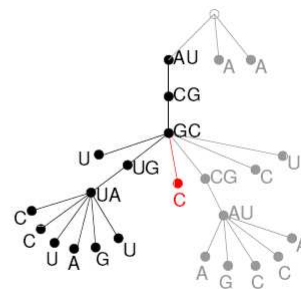
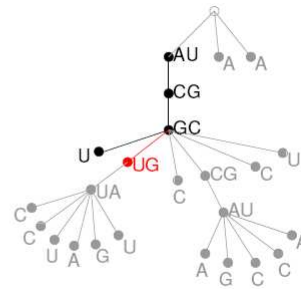


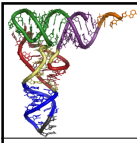
Structure 2^{aire} ↔ arbre



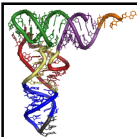
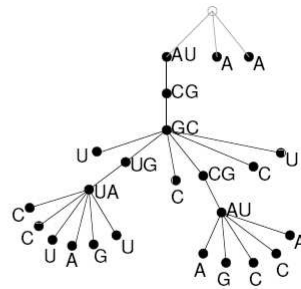
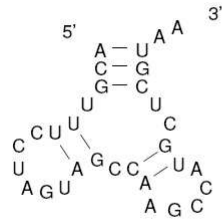
Structure 2^{aire} ↔ arbre



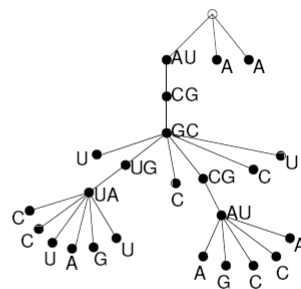
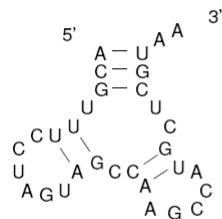




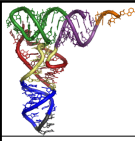
Structure 2^{aire} ↔ arbre



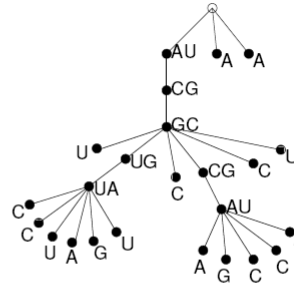
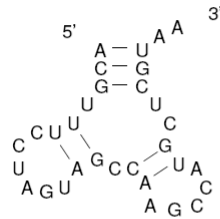
Structure 2^{aire} ↔ arbre



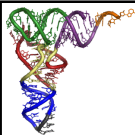
- Il est possible d'ajouter un racine pour obtenir un arbre.



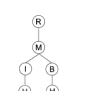
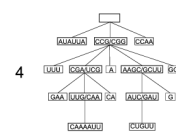
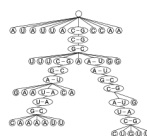
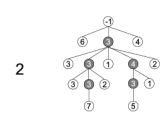
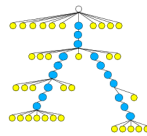
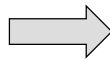
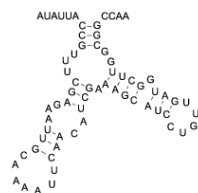
Structure 2^{aire} ↔ arbre

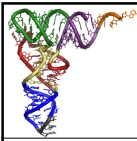


- ▶ Il est possible d'ajouter un racine pour obtenir un arbre.
- ▶ Les bases non appariées sont des feuilles.
- ▶ Les paires de bases sont des sommets internes.



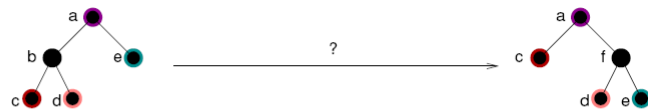
Différentes échelles de représentation



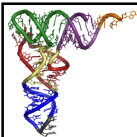
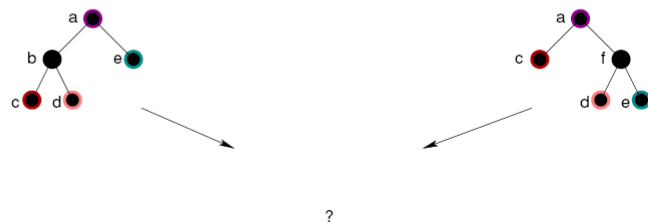


Edition et alignement d'arbres

Edition : Transformer un arbre en un autre.

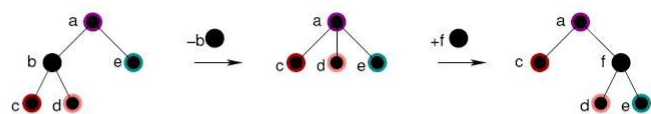


Alignement : Construire un super-arbre commun à deux arbres.

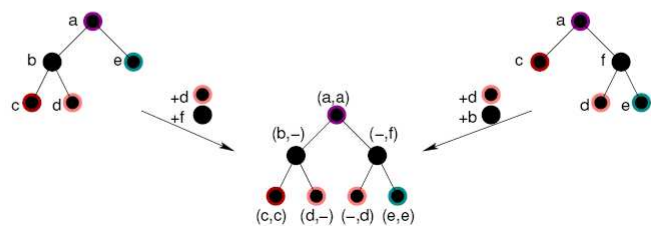


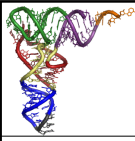
Edition et alignement d'arbres

Edition : Transformer un arbre en un autre.



Alignement : Construire un super-arbre commun à deux arbres.



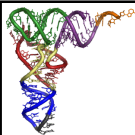
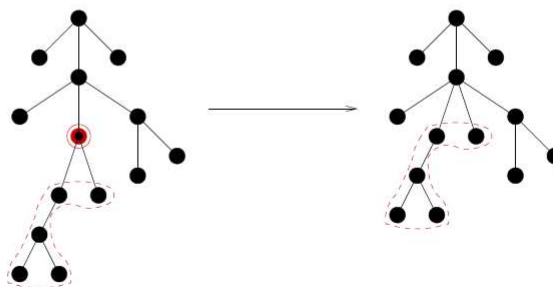


Opérations classiques pour les arbres

Suppression du sommet v :
les fils de v deviennent

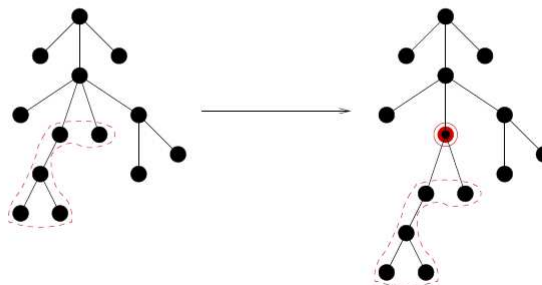
1. fils du père de v ,
2. frères droits des frères gauches de v ,
3. frères gauches des frères droits de v ,

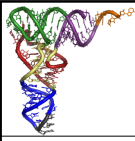
et leur ordre est conservé.



Opérations classiques pour les arbres

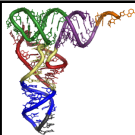
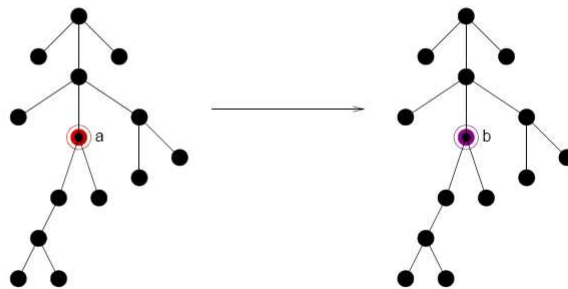
Insertion du sommet v :
un sommet v est ajouté comme père d'un certain nombre de frères consécutifs (opération symétrique à la suppression).





Opérations classiques pour les arbres

Substitution du sommet v :
l'étiquette de v est modifiée.

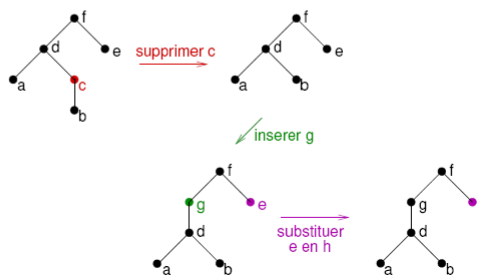


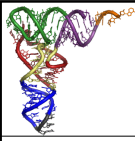
Opérations classiques pour les arbres

Chaque opération d'édition a un coût.

Edition : Transformer un arbre en un autre

- ▶ par une suite d'opérations d'édition,
- ▶ en minimisant le coût.

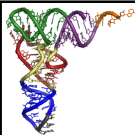
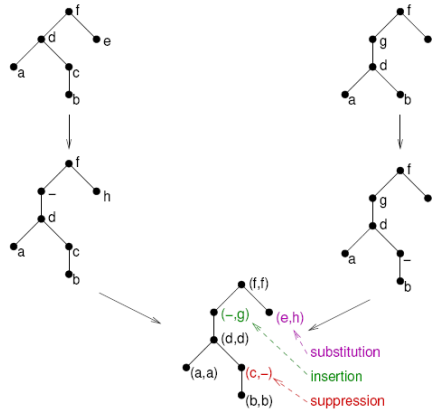




Opérations classiques pour les arbres

Alignement : Construire un super-arbre commun à deux arbres

- ▶ par une suite d'opérations d'édition,
- ▶ en minimisant le coût.



Tree edition algorithm

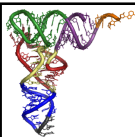
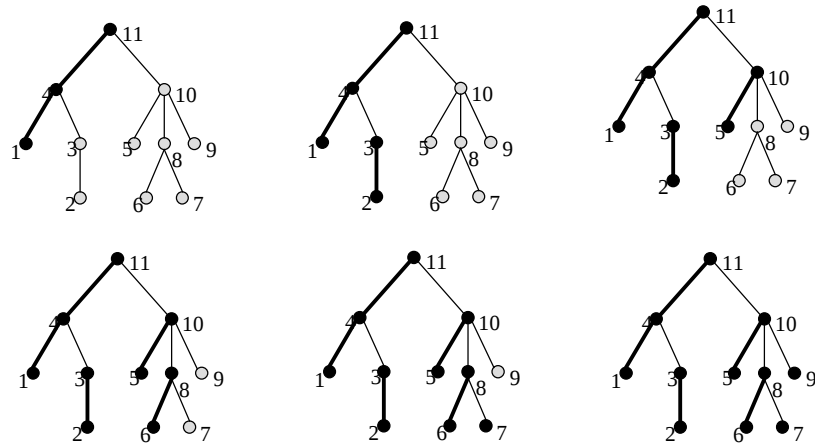
Zhang, Shasha 1989

$$\begin{aligned}
 & Tdist (\text{Tree 1}, \text{Tree 2}) \\
 &= \text{Min} \left\{ \begin{aligned} & Fdist (\text{Tree 1}, \text{Tree 2}) + \text{Change} (\text{Node 1}, \text{Node 2}), \\ & Fdist (\text{Tree 1}, \text{Tree 2}) + \text{Delete} (\text{Node 1}), \\ & Fdist (\text{Tree 1}, \text{Tree 2}) + \text{Insert} (\text{Node 2}) \end{aligned} \right\} \\
 & Fdist (\text{Tree 1}, \text{Tree 2}) \\
 &= \text{Min} \left\{ \begin{aligned} & Fdist (\text{Tree 1}, \text{Tree 2}) + Tdist (\text{Tree 1}, \text{Tree 2}), \\ & Fdist (\text{Tree 1}, \text{Tree 2}) + \text{Delete} (\text{Node 1}), \\ & Fdist (\text{Tree 1}, \text{Tree 2}) + \text{Insert} (\text{Node 2}) \end{aligned} \right\}
 \end{aligned}$$



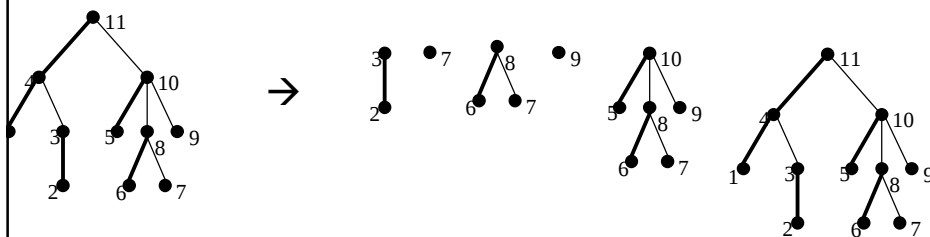
Algorithme d'édition

Décomposition en branches gauches



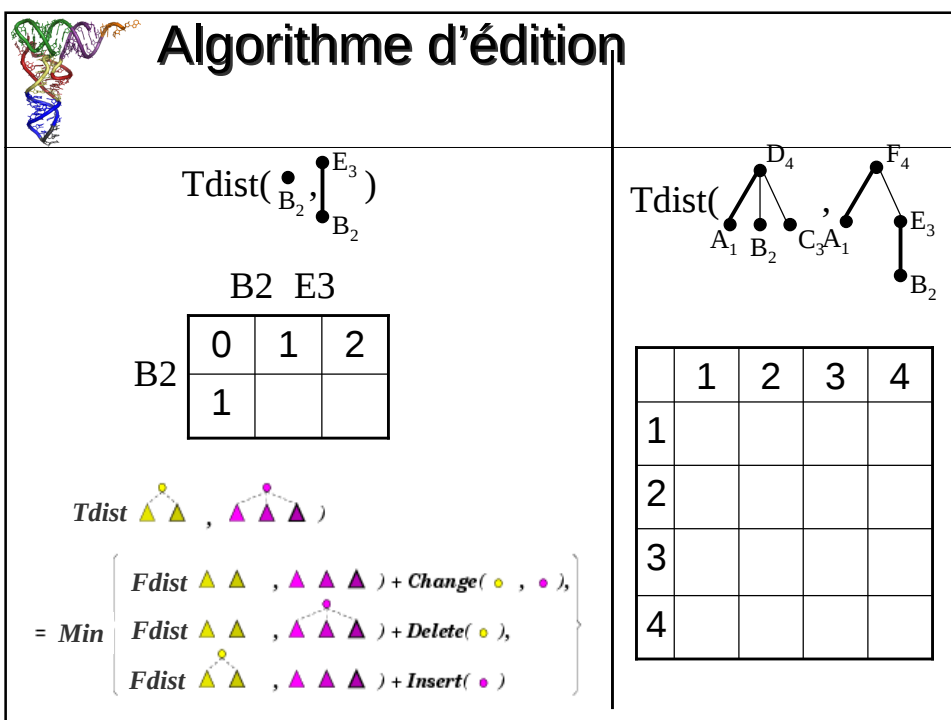
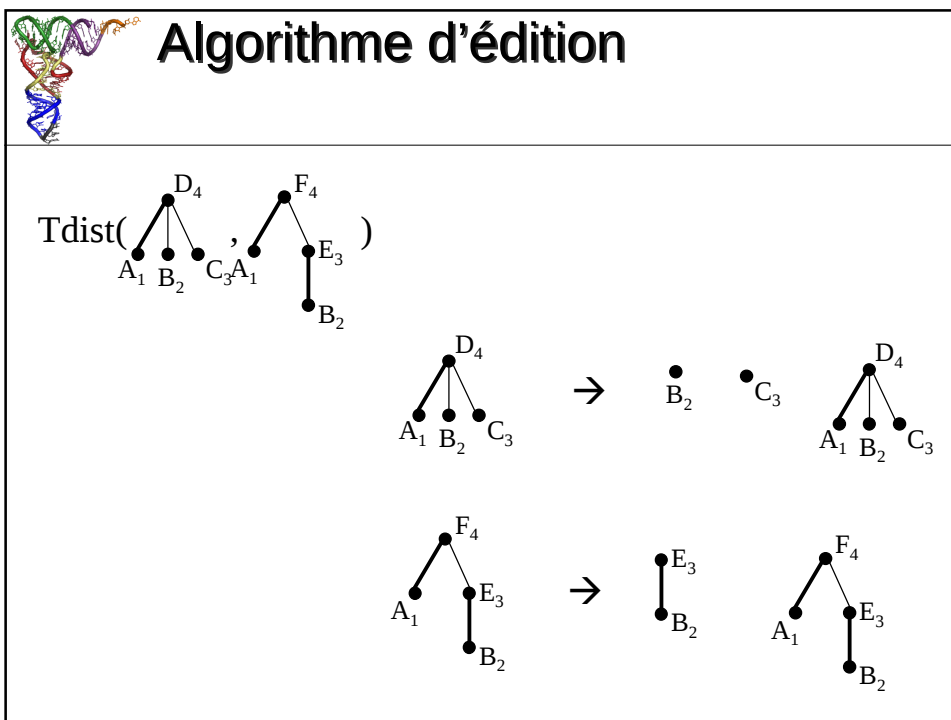
Algorithme d'édition

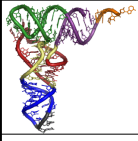
Sous-arbres spéciaux



LINUX → L LI LIN LINU LINUX

Ce n'est pas la seule décomposition possible [Klein 1998 ; Dulucq, Touzet 2003]





Algorithme d'édition

$$\text{Tdist}(\text{B}_2, \text{E}_3)$$

B2 E3

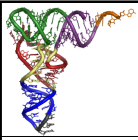
	B2	E3
B2	0	1
	1	0

$\text{Tdist}(\text{A}_1, \text{B}_2)$

$$= \text{Min} \left\{ \begin{array}{l} \text{Fdist}(\text{A}_1, \text{B}_2) + \text{Change}(\text{A}_1, \text{B}_2), \\ \text{Fdist}(\text{A}_1, \text{B}_2) + \text{Delete}(\text{A}_1), \\ \text{Fdist}(\text{A}_1, \text{B}_2) + \text{Insert}(\text{B}_2) \end{array} \right\}$$

$$\text{Tdist}(\text{A}_1, \text{B}_2, \text{C}_3, \text{A}_1, \text{E}_3, \text{B}_2)$$

	1	2	3	4
1				
2		0		
3				
4				



Algorithme d'édition

$$\text{Tdist}(\text{B}_2, \text{E}_3)$$

B2 E3

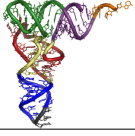
	B2	E3
B2	0	1
	1	0 ← 1

$\text{Tdist}(\text{A}_1, \text{B}_2)$

$$= \text{Min} \left\{ \begin{array}{l} \text{Fdist}(\text{A}_1, \text{B}_2) + \text{Change}(\text{A}_1, \text{B}_2), \\ \text{Fdist}(\text{A}_1, \text{B}_2) + \text{Delete}(\text{A}_1), \\ \text{Fdist}(\text{A}_1, \text{B}_2) + \text{Insert}(\text{B}_2) \end{array} \right\}$$

$$\text{Tdist}(\text{A}_1, \text{B}_2, \text{C}_3, \text{A}_1, \text{E}_3, \text{B}_2)$$

	1	2	3	4
1				
2		0	1	
3				
4				



Algorithme d'édition

$Tdist(\text{B}_2, \text{A}_1)$

$\text{A1} \quad \text{B2} \quad \text{E3} \quad \text{F4}$

	0	1	2	3	4
B2	1				

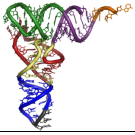
$Tdist(\text{A}_1, \text{B}_2)$

$= \min \left\{ \begin{array}{l} Fdist(\text{A}_1, \text{B}_2) + Change(\text{A}_1, \text{B}_2), \\ Fdist(\text{A}_1, \text{B}_2) + Delete(\text{A}_1), \\ Fdist(\text{A}_1, \text{B}_2) + Insert(\text{B}_2) \end{array} \right\}$

$Tdist(\text{D}_4, \text{F}_4)$

$\text{A}_1 \quad \text{B}_2 \quad \text{C}_3 \quad \text{A}_1$

	1	2	3	4
1				
2		0	1	
3				
4				



Algorithme d'édition

$Tdist(\text{B}_2, \text{A}_1)$

$\text{A1} \quad \text{B2} \quad \text{E3} \quad \text{F4}$

	0	1	2	3	4
B2	1	1			

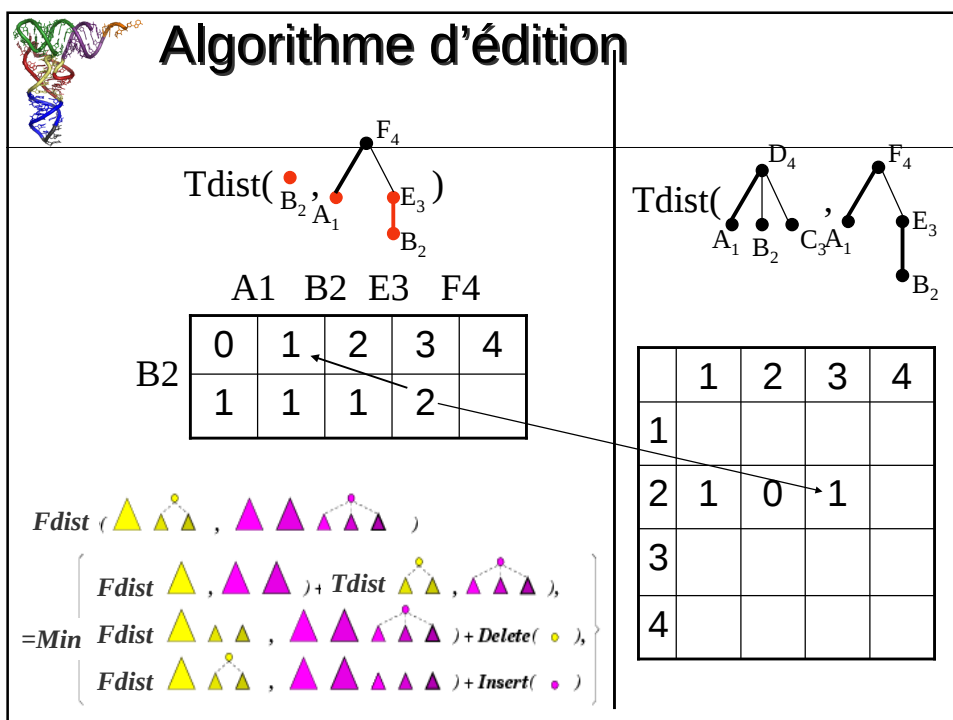
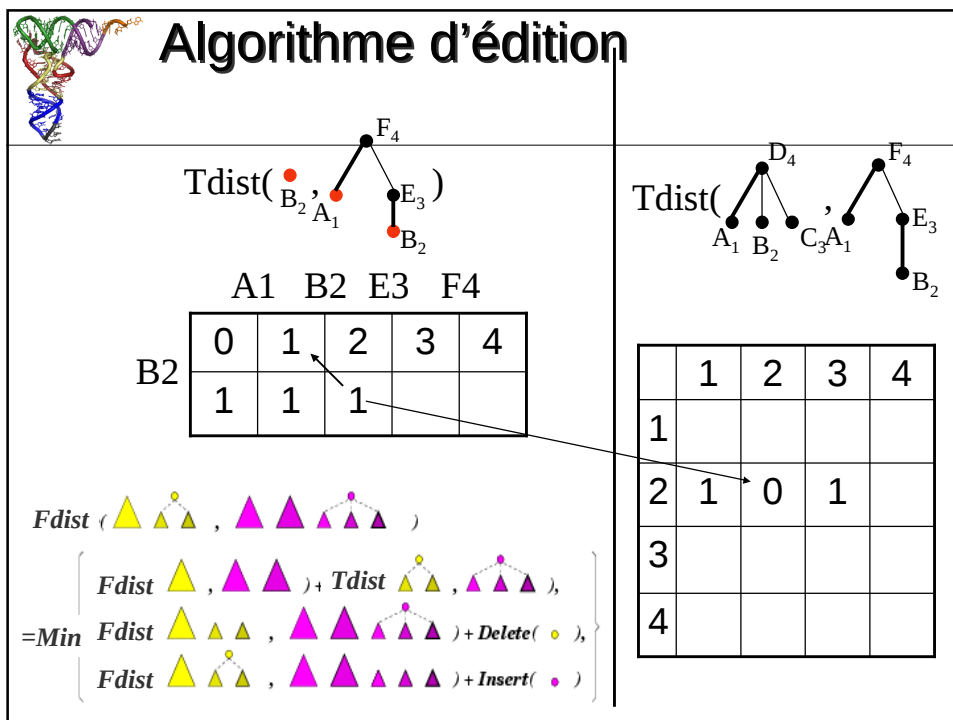
$Tdist(\text{A}_1, \text{B}_2)$

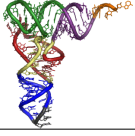
$= \min \left\{ \begin{array}{l} Fdist(\text{A}_1, \text{B}_2) + Change(\text{A}_1, \text{B}_2), \\ Fdist(\text{A}_1, \text{B}_2) + Delete(\text{A}_1), \\ Fdist(\text{A}_1, \text{B}_2) + Insert(\text{B}_2) \end{array} \right\}$

$Tdist(\text{D}_4, \text{F}_4)$

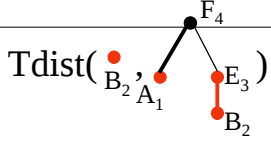
$\text{A}_1 \quad \text{B}_2 \quad \text{C}_3 \quad \text{A}_1$

	1	2	3	4
1				
2	1	0	1	
3				
4				





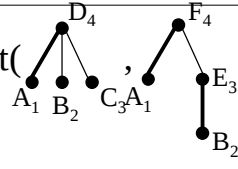
Algorithme d'édition



$$\text{Tdist}(B_2, A_1, E_3)$$

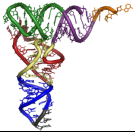
	A1	B2	E3	F4
B2	0	1	2	3
	1	1	1 ← 2	

$\text{Fdist}(\text{triangle}, \text{triangle}, \text{triangle})$
 $= \text{Min} \left\{ \begin{array}{l} \text{Fdist}(\text{triangle}, \text{triangle}, \text{triangle}) + \text{Tdist}(\text{triangle}, \text{triangle}, \text{triangle}), \\ \text{Fdist}(\text{triangle}, \text{triangle}, \text{triangle}) + \text{Delete}(\text{triangle}), \\ \text{Fdist}(\text{triangle}, \text{triangle}, \text{triangle}) + \text{Insert}(\text{triangle}) \end{array} \right\}$

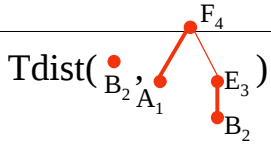


$$\text{Tdist}(A_1, B_2, C_3, A_1, E_3, B_2)$$

	1	2	3	4
1				
2	1	0	1	
3				
4				



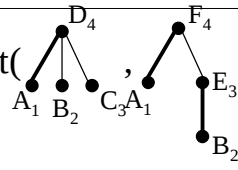
Algorithme d'édition



$$\text{Tdist}(B_2, A_1, E_3)$$

	A1	B2	E3	F4
B2	0	1	2	3
	1	1	1	2 ← 3

$\text{Tdist}(\text{triangle}, \text{triangle})$
 $= \text{Min} \left\{ \begin{array}{l} \text{Fdist}(\text{triangle}, \text{triangle}) + \text{Change}(\text{triangle}, \text{triangle}), \\ \text{Fdist}(\text{triangle}, \text{triangle}) + \text{Delete}(\text{triangle}), \\ \text{Fdist}(\text{triangle}, \text{triangle}) + \text{Insert}(\text{triangle}) \end{array} \right\}$



$$\text{Tdist}(A_1, B_2, C_3, A_1, E_3, B_2)$$

	1	2	3	4
1				
2	1	0	1	3
3				
4				

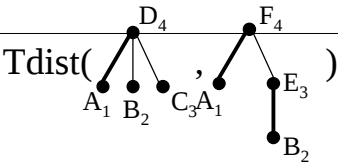


Algorithme d'édition

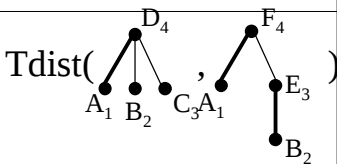
...



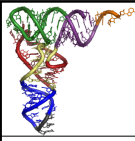
Algorithme d'édition



	A1	B2	E3	F4
A1	0	1	2	3
B2	1	0	1	2
C3	2	1	0	1
D4	3	2	1	2
	4	3	2	3

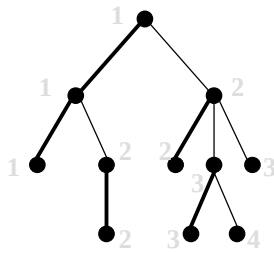


	1	2	3	4
1	0	1	2	3
2	1	0	1	3
3	1	1	2	4
4	3	3	3	3



Complexité de l'algorithme

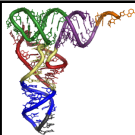
Chaque sommet s intervient autant de fois qu'il est dans un sous-arbre de la décomposition. C'est sa hauteur réduite $HR(s)$.



$$C_{\text{edit}}(T_1, T_2) = HR(T_1) \times HR(T_2)$$

Dans le pire des cas :

$$C_{\text{edit}}(T_1, T_2) = |T_1|^2 \times |T_2|^2$$



Complexité moyenne

[Dulucq, Tichit 2003]

Considérons la série génératrice des arborescences relativement à leur taille et à leur hauteur réduite,

$$f(q, t) = \sum_{n \geq 1, k \geq 1} a_{n,k} q^k t^n$$

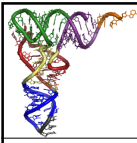
dans laquelle $a_{n,k}$ est le nombre d'arborescences ayant n sommets et une hauteur réduite égale à k .

La hauteur réduite moyenne d'un arbre à $n+1$ sommets est

$$HR(n+1) = \frac{1}{C_n} \sum_k k \cdot a_{n+1,k} = \frac{1}{C_n} \left(\left[\frac{d}{dq} f(q, t) \right]_{q=1}, t^{n+1} \right)$$

Où C_n désigne le nombre d'arbres à $n+1$ sommets :

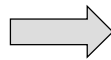
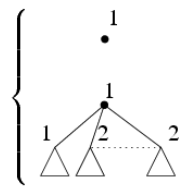
$$C_n = \frac{1}{n+1} \binom{2n}{n}$$



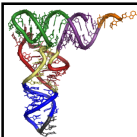
Complexité moyenne

[Dulucq, Tichit 2003]

$$f(q, t) = \sum_{n \geq 1, k \geq 1} a_{n,k} q^k t^n$$



$$f(q, t) = qt + qt \cdot f(q, t) \frac{1}{1 - f(q, qt)}$$



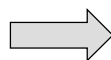
Complexité moyenne

[Dulucq, Tichit 2003]

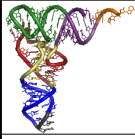
$$\left[\frac{d}{dq} f(q, t) \right]_{q=1} = \frac{t}{2} \frac{1 - 2t + \sqrt{1 - 4t}}{1 - 4t}$$

$$\left(\left[\frac{d}{dq} f(q, t) \right]_{q=1}, t^{n+1} \right) = \begin{cases} 4^{n-1} + \binom{2n-1}{n} & \text{si } n \geq 1 \\ 1 & \text{si } n = 0 \end{cases}$$

$$HR(n+1) = (n+1) \frac{\binom{2n-1}{n} + 4^{n-1}}{\binom{2n}{n}}$$



$$C_{\text{edit}} = HR(n) \times HR(m) \sim n^{3/2} m^{3/2}$$

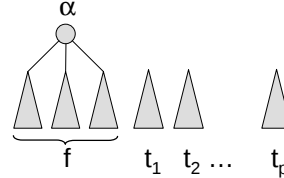


Tree edition algorithm

Zhang, Shasha 1989

$$\text{Score}(\alpha(f), \alpha'(f')) = \text{Max}$$

$$\begin{cases} \text{Subs}(\alpha, \alpha') + \text{Score}(f, f') \\ \text{Ins}(\alpha') + \text{Score}(\alpha(f), f') \\ \text{Del}(\alpha) + \text{Score}(f, \alpha'(f')) \end{cases}$$



$$\text{Score}([\alpha(f) \circ t_1 \circ \dots \circ t_p], [\alpha'(f'), t'_1 \circ \dots \circ t'_q]) = \text{Max}$$

$$\begin{cases} \text{Score}(\alpha(f), \alpha'(f')) + \text{Score}([t_1 \circ \dots \circ t_p], [t'_1 \circ \dots \circ t'_q]) \\ \text{Ins}(\alpha') + \text{Score}([\alpha(f) \circ t_1 \circ \dots \circ t_p], [f', t'_1 \circ \dots \circ t'_q]) \\ \text{Del}(\alpha) + \text{Score}([f \circ t_1 \circ \dots \circ t_p], [\alpha'(f') \circ t'_1, \dots \circ t'_q]) \end{cases}$$

$O(n^3 \log n)$ [Klein 1998]

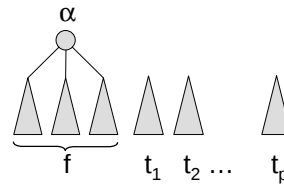


Tree alignment algorithm

Jiang, Wang, Zhang 1995

$$\text{Score}(r(f), r'(f')) = \text{Max}$$

$$\begin{cases} \text{Subs}(\alpha, \alpha') + \text{Score}(f, f') \\ \text{Ins}(\alpha') + \text{Score}(\alpha(f), f') \\ \text{Del}(\alpha) + \text{Score}(f, \alpha'(f')) \end{cases}$$



$$\text{Score}(\alpha(f) \circ t_1 \circ \dots \circ t_p; \alpha'(f') \circ t'_1 \circ \dots \circ t'_q) = \text{Max}$$

$$\begin{cases} \text{Score}(\alpha(f); \alpha'(f')) + \text{Score}(t_1 \circ \dots \circ t_p; t'_1 \circ \dots \circ t'_q) \\ \text{Ins}(\alpha') + \text{Max}_i \{ \text{Score}(\alpha(f) \circ \dots \circ t_i; f') + \text{Score}(t_{i+1} \circ \dots \circ t_p; t'_1 \circ \dots \circ t'_q) \} \\ \text{Del}(\alpha) + \text{Max}_j \{ \text{Score}(f; \alpha'(f') \circ t'_1 \circ \dots \circ t'_j) + \text{Score}(t_1 \circ \dots \circ t_p; t'_{j+1} \circ \dots \circ t'_q) \} \end{cases}$$

$O(n^4 \log n)$



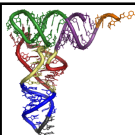
Edition vs Alignment

$$\text{Score}([r(f), t_1, \dots, t_p], [r'(f'), t'_1, \dots, t'_q]) = \text{Max}$$

$$\left\{ \begin{array}{l} \dots \\ \text{Ins}(r') + \text{Score}([r(f), t_1, \dots, t_p], [f', t'_1, \dots, t'_q]) \\ \dots \end{array} \right.$$

$$\text{Score}([r(f), t_1, \dots, t_p], [r'(f'), t'_1, \dots, t'_q]) = \text{Max}$$

$$\left\{ \begin{array}{l} \dots \\ \text{Ins}(r') + \text{Max} \{ \text{Score}([r(f), \dots, t_i], f') + \text{Score}([t_{i+1}, \dots, t_p], [t'_1, \dots, t'_q]) \} \\ \dots \end{array} \right.$$



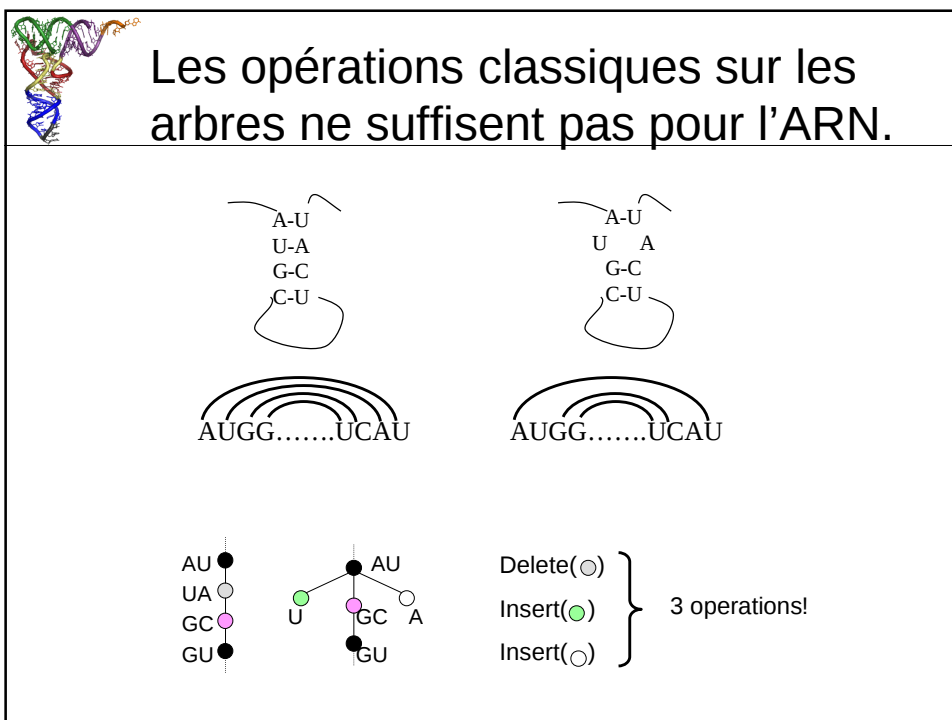
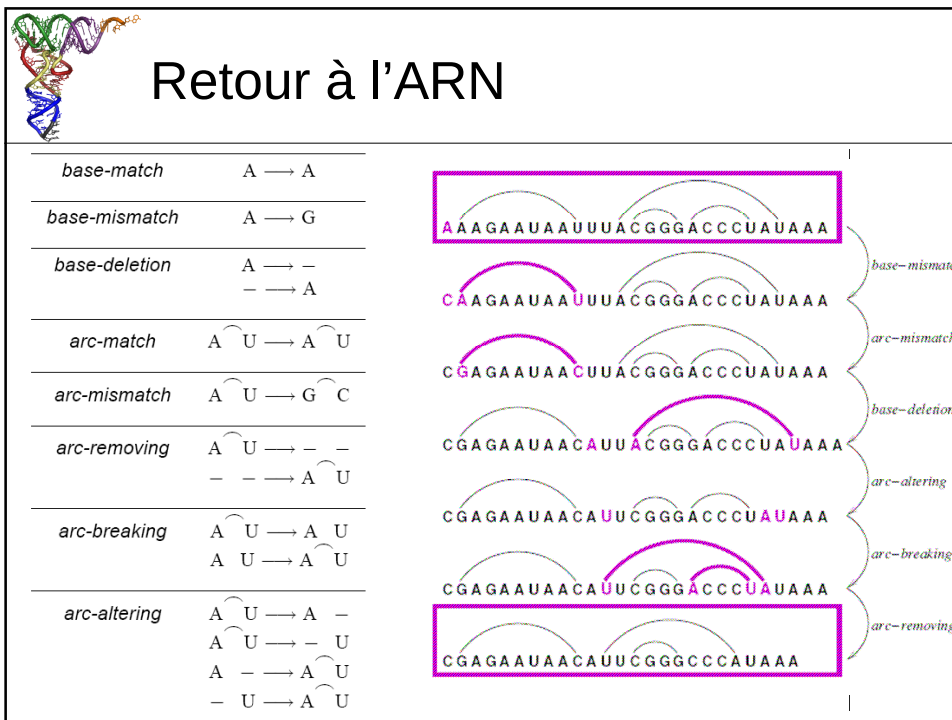
Edition vs Alignment

$$\text{Score}(\text{Diagram 1}, \text{Diagram 2}) = \text{Max}$$

$$\left\{ \begin{array}{l} \dots \\ \text{Ins}(\bullet) + \text{Score}(\text{Diagram 1}, \text{Diagram 2}) \\ \dots \end{array} \right.$$

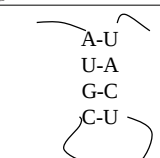
$$\text{Score}(\text{Diagram 1}, \text{Diagram 2}) = \text{Max}$$

$$\left\{ \begin{array}{l} \dots \\ \text{Ins}(\bullet) + \text{Max} \{ \text{Score}(\text{Diagram 1}, \text{Diagram 2}) + \text{Score}(\text{Diagram 1}, \text{Diagram 2}) \} \\ \dots \end{array} \right.$$

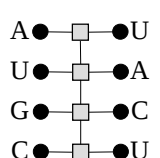


A first solution

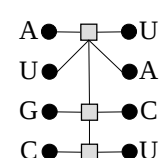
Höchsmann, Töller, Gierich, Kurtz 2003
(RNAforester)



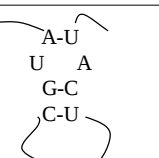
A-U
U-A
G-C
C-U



A • □ • U
U • □ • A
G • □ • C
C • □ • U



A • □ • U
U • □ • A
G • □ • C
C • □ • U



A-U
U A
G-C
C-U



AUGG.....UCAU



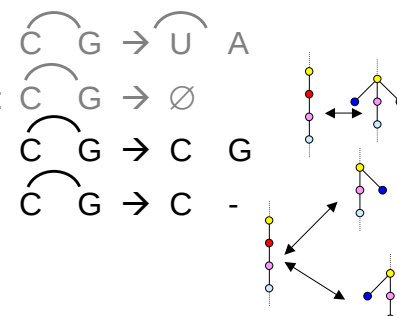
AUGG.....UCAU

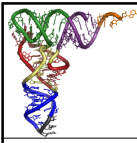
But this implies some constraints on the scores. For example:

Arc-deletion = Arc-Breaking + 2 Base-Deletion

New edition operations on trees

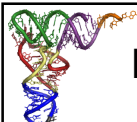
- Operations on bases:
 - Substitution: $A \rightarrow C$
 - Deletion / Insertion: $A \rightarrow \emptyset$
- Operations on arcs:
 - Arc-substitution: $\overset{\curvearrowright}{C} G \rightarrow \overset{\curvearrowright}{U} A$
 - Arc-deletion / Arc-insertion: $\overset{\curvearrowright}{C} G \rightarrow \emptyset$
 - Arc-breaking / :
 - Arc-altering / :





Tree edition, tree alignment

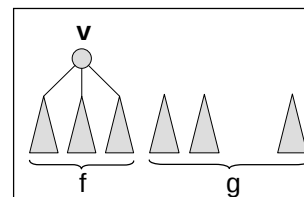
	Tree operations	RNA operations
Edition	$O(n^3 \log n)$ [Zhang-Shasha 1989, Klein 1998]	NP-complete [Blin, Fertin, Sinoquet, Rusu 2003]
Alignment	$O(n^4)$ [Jiang, Wang, Zhang 1995]	?



RNA structure alignment algorithm

$$A(v(f) \circ g, v'(f') \circ g') =$$

$\min \left\{ \begin{array}{l} \text{basedel}(v) + A(g, v'(f') \circ g') \\ \quad \text{if } v \text{ is a leaf} \\ \text{baseins}(v') + A(v(f) \circ g, g') \\ \quad \text{if } v' \text{ is a leaf} \\ \text{basesub}(v, v') + A(g, g') \quad [\text{Jiang, Wang, Zhang 1995}] \\ \quad \text{if } v, v' \text{ are leaves} \\ \text{pairedel}(v) + \min\{A(f, p') + A(g, s') \mid p' \circ s' = v'(f') \circ g'\} \\ \quad \text{if } v \text{ is an internal node} \\ \text{pairins}(v') + \min\{A(p, f') + A(s, g') \mid p \circ s = v(f) \circ g\} \\ \quad \text{if } v' \text{ is an internal node} \\ \text{pairs}(v, v') + A(f, f') + A(g, g') \\ \quad \text{if } v, v' \text{ are internal nodes} \\ \text{scission}(v, v') + \min\{A(f, p') + A(g, s') \mid p' \circ v'_c \circ s' = g'\} \\ \quad \text{if } v \text{ is an internal node, } v', v'_c \text{ are leaves} \\ \text{fusion}(v, v_c, v') + \min\{A(p, f') + A(s, g') \mid p \circ v_c \circ s = g\} \\ \quad \text{if } v, v_c \text{ are leaves, } v' \text{ is an internal node} \\ \text{leftalt}(v, v'_c) + \min\{A(f, p') + A(g, s') \mid p' \circ v'_c \circ s' = v'(f') \circ g'\} \\ \quad \text{if } v \text{ is an internal node, } v'_c \text{ is a leaf} \\ \text{rightalt}(v, v') + \min\{A(f, p') + A(g, s') \mid p' \circ s' = g'\} \\ \quad \text{if } v \text{ is an internal node, } v' \text{ is a leaf} \\ \text{leftcompl}(v_c, v') + \min\{A(p, f') + A(s, g') \mid p \circ v_c \circ s = v(f) \circ g\} \\ \quad \text{if } v_c \text{ is a leaf, } v' \text{ is an internal node} \\ \text{rightcompl}(v, v') + \min\{A(p, f') + A(s, g') \mid p \circ s = g\} \\ \quad \text{if } v \text{ is a leaf, } v' \text{ is an internal node} \end{array} \right.$



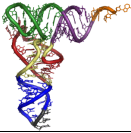
Spécifique à l'ARN

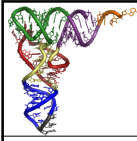
[Blin, Denise, Dulucq, Herrbach, Touzet 2008]

Complexité au pire : $O(n^4)$

66

 Complexité de la comparaison		
	Arbres	ARN
Edition	Pire : $O(n^4)$ / $O(n^3 \log n)$ [Zhang, Shasha 1989 / Klein 1998] Moyenne : $O(n^3)$ [Dulucq, Tichit 2003]	NP-complet [Blin, Fertin, Sinoquet, Rusu 2003]
Alignement	Pire : $O(n^4)$ [Jiang, Wang, Zhang 1995] Moyenne : $O(n^2)$ [Herrbach, Denise, Dulucq 2007]	Pire : $O(n^4)$ [Blin, Denise, Dulucq, Herrbach, Touzet 2008] Moyenne : $O(n^2)$ [Herrbach, Denise, Dulucq 2007]
17/01/2010 Projet Brasero - ANR Blanc 2006 67		

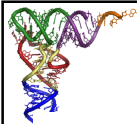
 Une approche multi-échelles		



Algorithmme d'édition

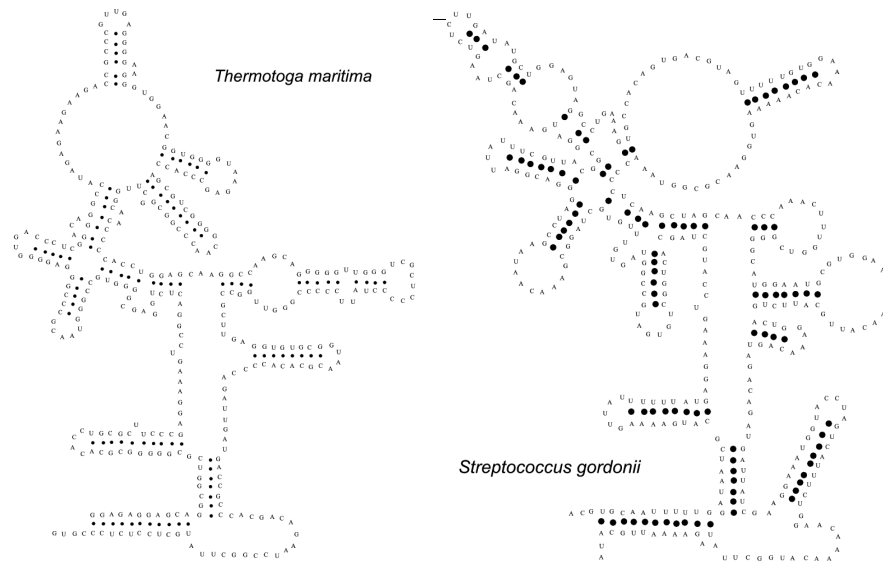
Zhang, Shasha 1989

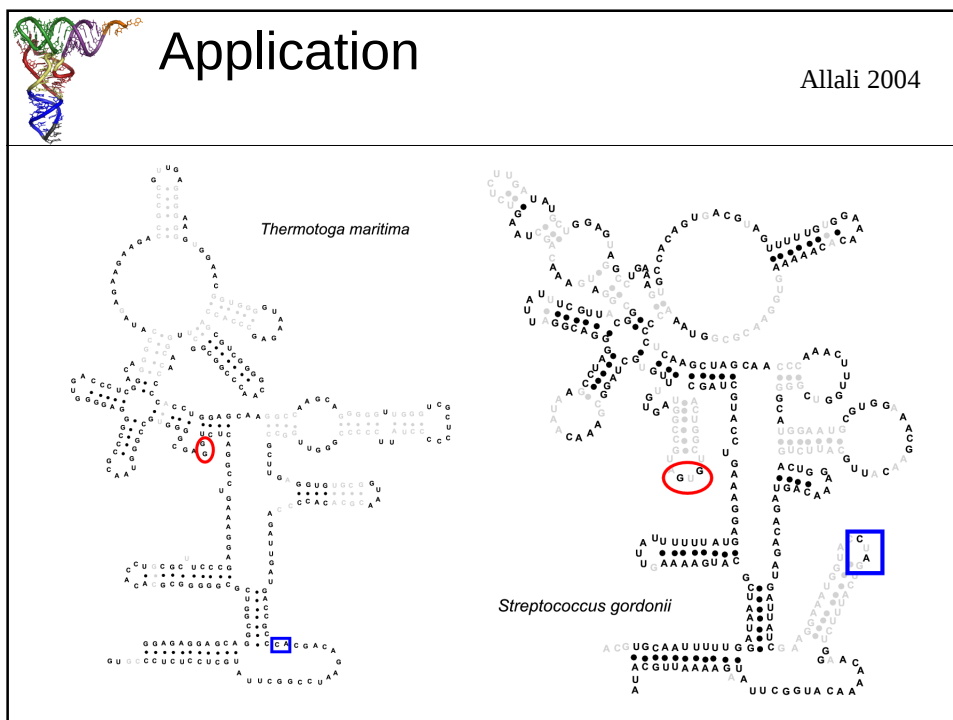
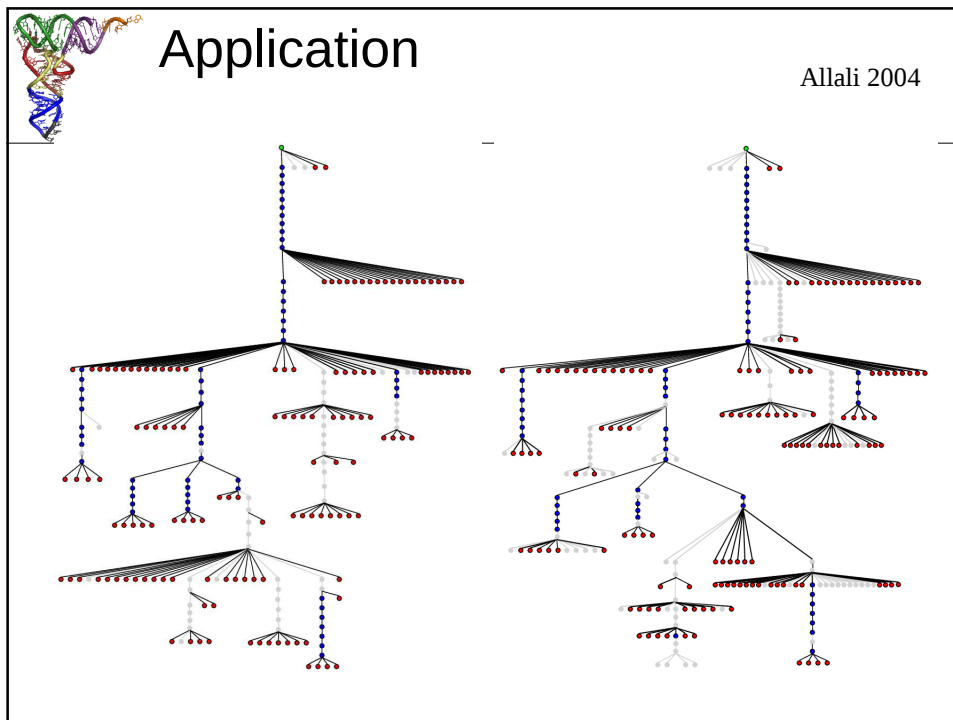
$$\begin{aligned}
 & Tdist (\triangle \triangle , \triangle \triangle \triangle) \\
 & = \text{Min} \left\{ \begin{aligned} & Fdist (\triangle \triangle , \triangle \triangle \triangle) + \text{Change} (\triangle , \triangle), \\ & Fdist (\triangle \triangle , \triangle \triangle \triangle) + \text{Delete} (\triangle), \\ & Fdist (\triangle \triangle , \triangle \triangle \triangle) + \text{Insert} (\triangle) \end{aligned} \right\} \\
 \\
 & Fdist (\triangle \triangle \triangle , \triangle \triangle \triangle \triangle) \\
 & = \text{Min} \left\{ \begin{aligned} & Fdist (\triangle \triangle \triangle , \triangle \triangle \triangle \triangle) + Tdist (\triangle \triangle , \triangle \triangle \triangle), \\ & Fdist (\triangle \triangle \triangle , \triangle \triangle \triangle \triangle) + \text{Delete} (\triangle), \\ & Fdist (\triangle \triangle \triangle , \triangle \triangle \triangle \triangle) + \text{Insert} (\triangle) \end{aligned} \right\}
 \end{aligned}$$

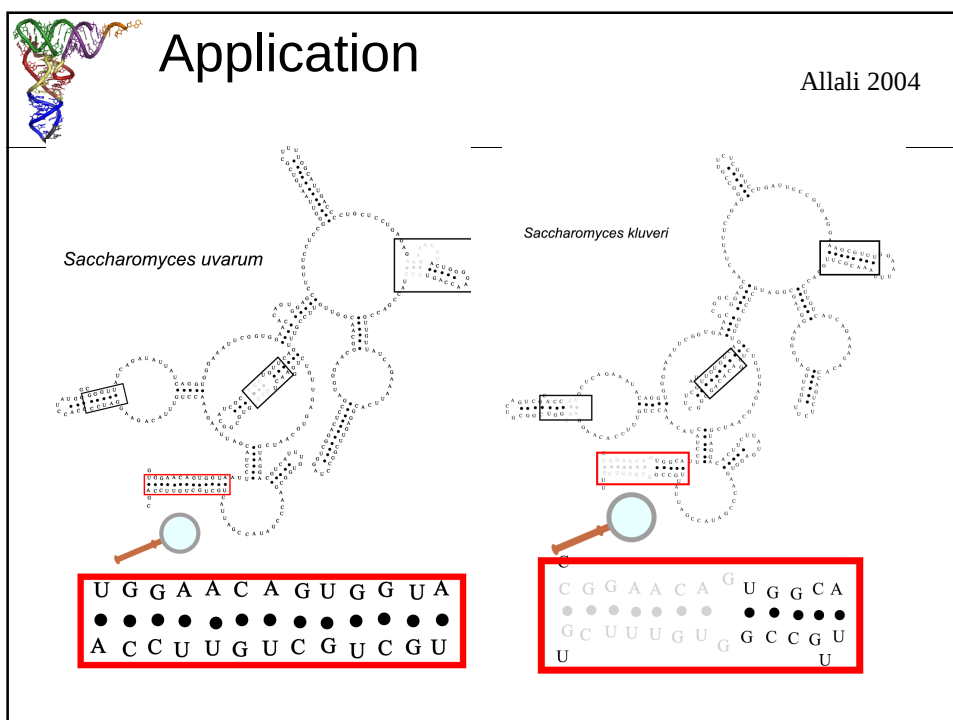
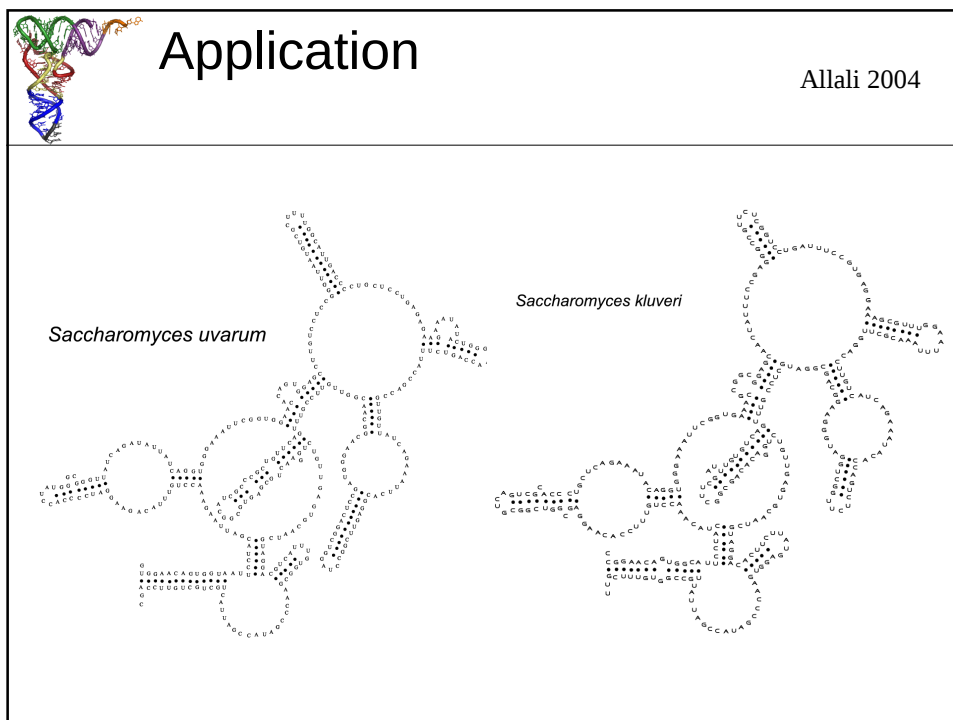


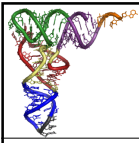
Application

Allali 2004



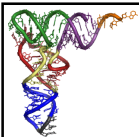
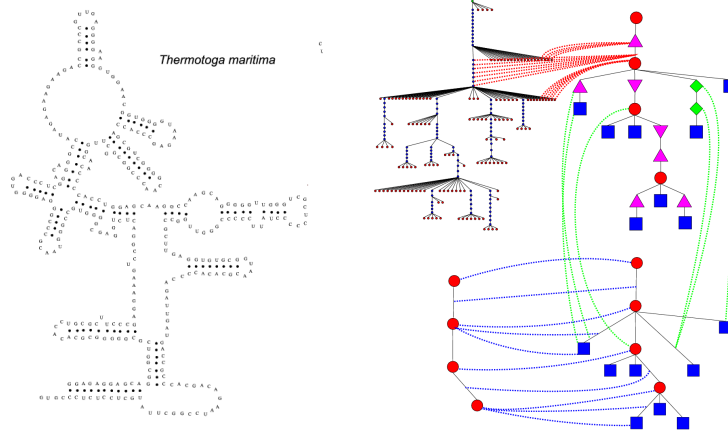






Une approche multi-échelles

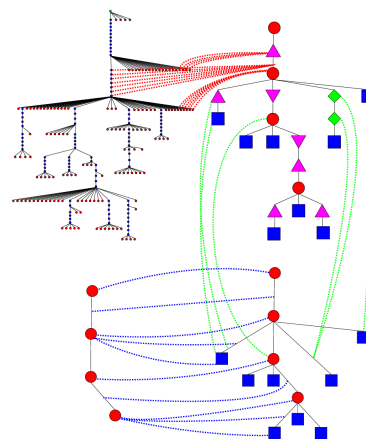
Allali, Sagot 2004

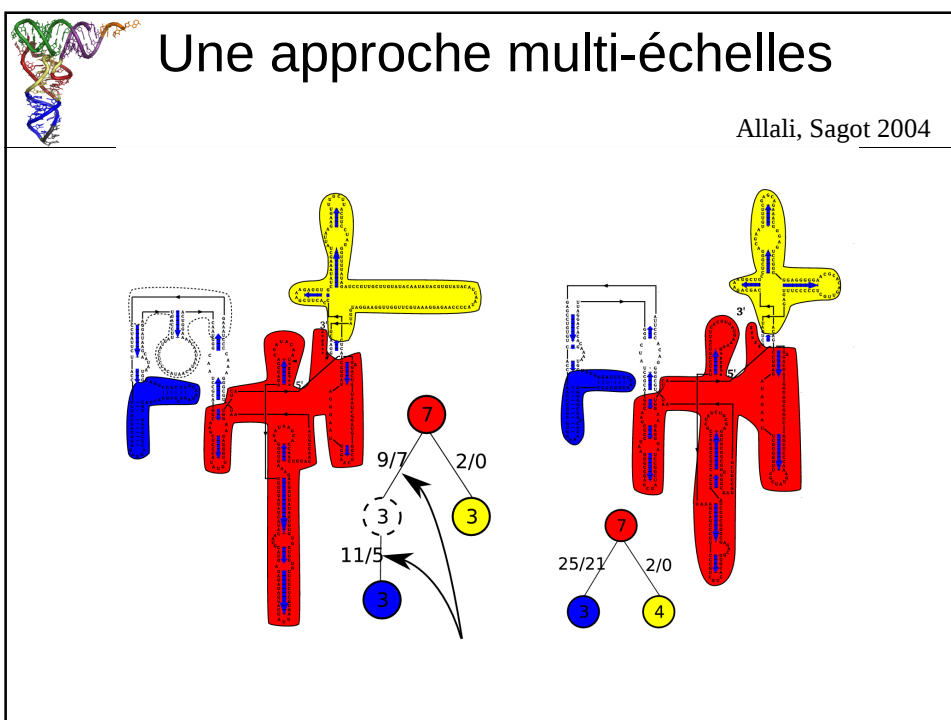
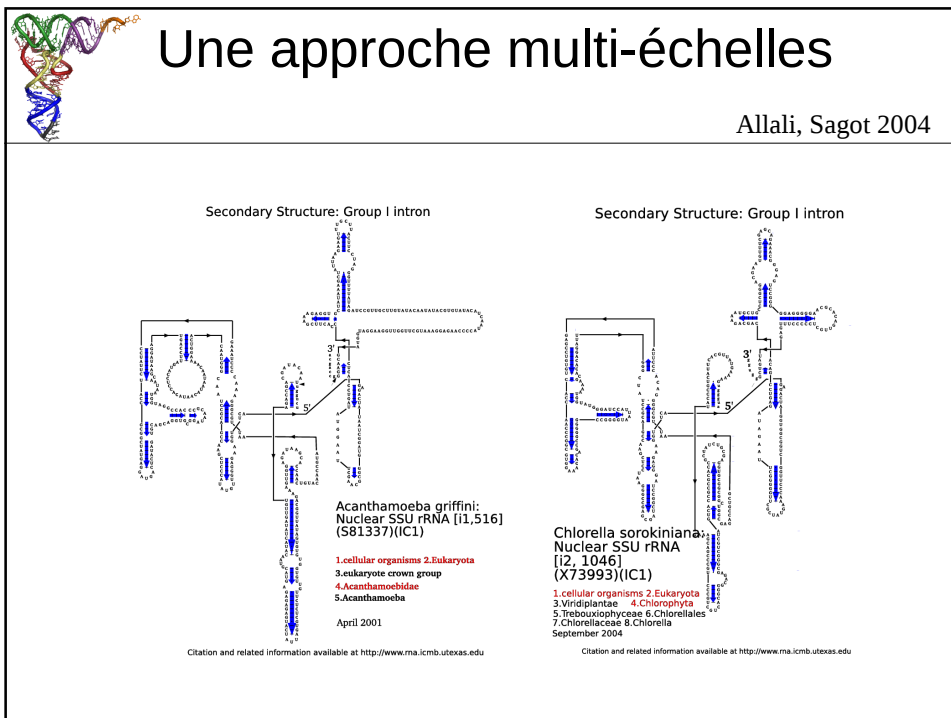


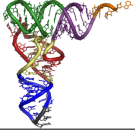
Une approche multi-échelles

Allali, Sagot 2004

Opérations supplémentaires :
Fusion de nœuds
Fusion d'arêtes

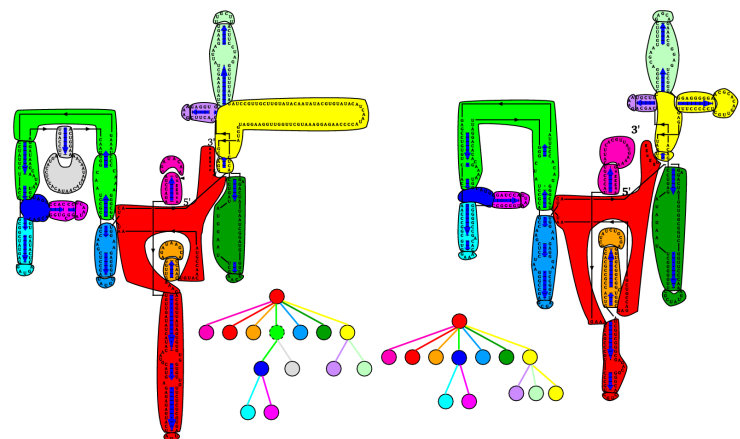


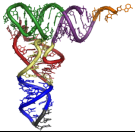




Une approche multi-échelles

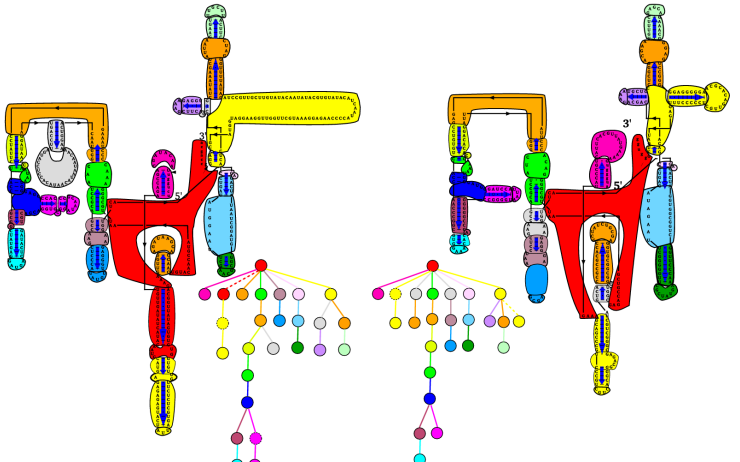
Allali, Sagot 2004

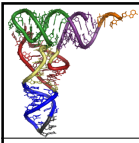




Une approche multi-échelles

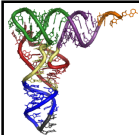
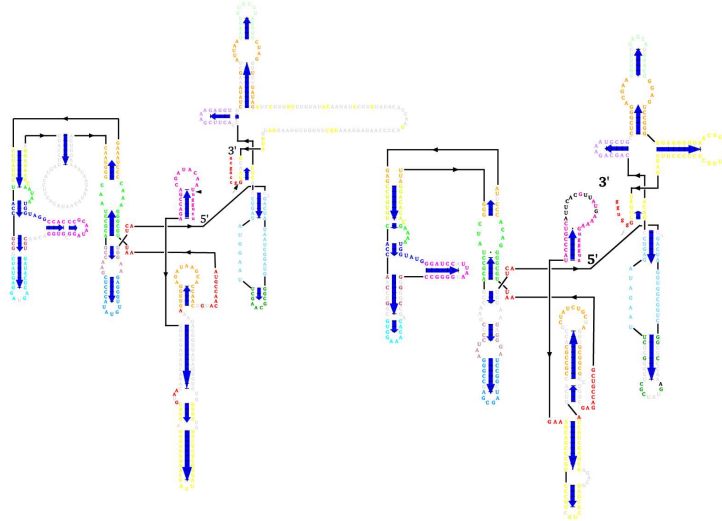
Allali, Sagot 2004





Une approche multi-échelles

Allali, Sagot 2004



Logiciels

NestedAlign

Global alignment score : 4

Alignment of telomere1 (sequence, structure) and telomere2 (sequence, structure)

```

telomere1  = - - - - - ( ( ( ( ( - - - - - ) ) ) ) ) - - - - - ( ( ( ( (
telomere1  = A-DG - - - - - AAAATCATACTGCGT-DG -CAATCAACAGCATCATCAACCCCAAAACG-DAGAG-CA-AATATGTCGTCGTC
telomere1  = AGT-DG - - - - - GGA-TT-A-ATGAGATCGTAGAATCTGCAAGCA - - - - - AAAAGCCG -AAAACCTGACATGAGAGCA-TT-TT-TAG
telomere1  = - - - - - ( ( ( ( ( - - - - - ( ( ( ( ( - - - - - ) ) ) ) ) - - - - -

telomere2  = - - - - - ( ( ( ( ( - - - - - ) ) ) ) ) - - - - - ( ( ( ( ( - - - - - ) ) ) ) ) - - - - - ( ( ( ( (
telomere2  = - - - - - CCGGCGAGAC - - - - - AAA-CAGAGAG - - - - - GAG - - - - - GGA-D-A-A - - - - - GCG - - - - - DCG - - - - - AGG-A-
telomere2  = - - - - - CCGGATG -ACGCTGTAAGAAACAAAGAGGCAAGGCAATAGAG - - - - - ACCTGGAGGAGGAGAAACCCGACAGATGCGAGAAATGAT
telomere2  = - - - - - ( ( ( ( ( - - - - - ) ) ) ) ) - - - - - ( ( ( ( ( - - - - - ) ) ) ) ) - - - - - ( ( ( ( (

```

<http://nestedalign.lri.fr>

[illegible]

[illegible][illegible]



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