

# Gibbs Sampling

ttacctt**aac**

**gata**tctgtc

**acg**gcgttcg

ccct**aaa**gag

cgtc**aga**ggt

**ttac**cttaac

gat**atc**tgtc

acggcg**ttc**g

ccctaa**agag**

**cgt**cagaggt

ttacctt**aac**

**gata**tctgtc

**acg**gcgttcg

ccct**aaa**gag

cgtc**aga**ggt

ttacctt**aac**

gatatc**tgt**c

**acg**gcgttcg

ccct**aaa**gag

cgtc**aga**ggt

RandomizedMotifSearch

(may change all k-mers in one step)

GibbsSampler

(changes one k-mer in one step)

# Algoritmo GibbsSampler

```
GibbsSampler(Dna, k, t, N)
    randomly select k-mers Motifs = (Motif1, ..., Motift) in each string from Dna
    BestMotifs ← Motifs
    for j ← 1 to N
        i ← Random(t)
        Profile ← profile matrix constructed from all strings in Motifs except for Motifi
        Motifi ← Profile-randomly generated k-mer in the i-th sequence
        if Score(Motifs) < Score(BestMotifs)
            BestMotifs ← Motifs
    return BestMotifs
```

## Un ejemplo de GibbsSampling

	ttACCT <b>taac</b>	ttACCT <b>taac</b>
	gAT <b>GTct</b> gtc	gAT <b>GTct</b> gtc
Dna	<b>ccgG</b> CGTtag	→ -----
	<b>cacta</b> ACGAg	<b>cacta</b> ACGAg
	cgtcag <b>AGGT</b>	cgtcag <b>AGGT</b>

	Motifs	Profile(Motifs)
Count(Motifs)	t a a c G T c t a c t a A G G T	A: 2/4 1/4 1/4 1/4 C: 0 1/4 1/4 1/4 G: 1/4 1/4 1/4 0 T: 1/4 1/4 1/4 2/4
A:	2 1 1 1	A: 2/4 1/4 1/4 1/4
C:	0 1 1 1	C: 0 1/4 1/4 1/4
G:	1 1 1 0	G: 1/4 1/4 1/4 0
T:	1 1 1 2	T: 1/4 1/4 1/4 2/4

	Count(Motifs)	Profile(Motifs)
A:	3 2 2 2	3/8 2/8 2/8 2/8
C:	1 2 2 2	1/8 2/8 2/8 2/8
G:	2 2 2 1	2/8 2/8 2/8 1/8
T:	2 2 2 3	2/8 2/8 2/8 3/8

Dna    ttACCT**taac**  
 Dna    gAT**GTct**gtc  
 Dna    **ccgG**CGTtag    →    -----  
 Dna    **cacta**ACGAg  
 Dna    cgtcag**AGGT**

ttACCT**taac**  
 gAT**GTct**gtc  
 -----  
 ccg**GCGT**tag  
**cacta**ACGAg  
 cgtcag**AGGT**

ttACCT**taac**  
 gAT**GTct**gtc  
 ccg**GCGT**tag  
**cacta**ACGAg  
 cgtcag**AGGT**

ccgG	cgGC	gGCG	GC GT	CG Tt	GT ta	T tag
$4/8^4$	$8/8^4$	$8/8^4$	$24/8^4$	$12/8^4$	$16/8^4$	$8/8^4$

Dna    ttACCT**taac**                    -----  
          gAT**GTct**gtc                    gAT**GTct**gtc  
          ccg**GCGT**tag    →    ccg**GCGT**tag  
          **cacta**ACGAg                    **cacta**ACGAg  
          cgtcag**AGGT**                    cgtcag**AGGT**

	ttACCT <b>taac</b>	-----	
	gAT <b>GTct</b> gtc	gAT <b>GTct</b> gtc	
Dna	ccg <b>GCGT</b> tag	→	ccg <b>GCGT</b> tag
	c <b>acta</b> ACGAg	c <b>acta</b> ACGAg	
	cgtcag <b>AGGT</b>	cgtcag <b>AGGT</b>	
Motifs	G T c t	A: 2/4 0 0 1/4	
	G C G T	C: 0 2/4 1/4 0	
	a c t a	G: 2/4 1/4 2/4 0	
	A G G T	T: 0 1/4 1/4 3/4	
Profile(Motifs)			

ttACCT**taac**

gAT**GTct**gtc

ccg**GCGT**tag

**cacta**ACGAg

cgtcag**AGGT**

Count(Motifs)  
A: 3 1 1 2  
C: 1 3 2 1  
G: 3 2 3 1  
T: 1 2 2 4

Profile(Motifs)  
A: 3/8 1/8 1/8 2/8  
C: 1/8 3/8 2/8 1/8  
G: 3/8 2/8 3/8 1/8  
T: 1/8 2/8 2/8 4/8

ttAC	tACC	ACCT	CCTt	CTta	Ttaa	taac
$2/8^4$	$2/8^4$	$72/8^4$	$24/8^4$	$8/8^4$	$4/8^4$	$1/8^4$

tt**ACCT**taac

gAT**GTct**gtc

ccg**GCGT**tag

cacta**ACGA**g

cgtcag**AGGT**