



# 稻曲病菌 UV-2 菌株细菌人工染色体文库构建及分析

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**摘要:** 【目的】稻曲病(Rice false smut)是由稻曲病菌[*Villosiclava virens* (Cooke) Tak.]引起的严重危害水稻的真菌病害。构建稻曲病菌 UV-2 的大片段 DNA 细菌人工染色体(Bacterial artificial chromosome, BAC)文库,为致病相关基因的鉴定及在图位克隆、比较基因组学等方面的研究奠定基础。【方法】以幼嫩菌丝为材料制备大分子基因组 DNA 包埋块,用 *Hind* III 部分酶解后经脉冲凝胶电泳筛选,回收大片段 DNA 并与 pIndigoBAC536-S 载体连接,连接产物转化大肠杆菌菌株 DH10B T1 Phage-Resistant 细胞后进行蓝白斑筛选,白色菌落转入 384 孔板置于 -80 °C 低温保存。【结果】成功构建 UV-2 菌株的高质量、高覆盖度的 BAC 文库,该文库共含 10 368 个克隆,平均插入片段为 124.4 kb,空载率小于 1%,约覆盖该菌基因组的 36.8 倍。【结论】克服了真菌大分子基因组 DNA 制备难控制的技术难题,建立了首个稻曲病菌的 BAC 文库。该文库已作为一种公共基因组资源向研究者开放(<http://GResource.hzau.edu.cn>)。

**关键词:** 大分子 DNA, 细菌人工染色体, 水稻, 稻曲病, 稻曲病菌

# Construction of a bacterial artificial chromosome library of *Villosiclava virens* UV-2 genome

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**Abstract:** [Objective] Rice false smut caused by *Villosiclava virens* (Cooke) Tak. is a serious fungal disease of rice. This study aimed to construct a high molecular weight bacterial artificial chromosome (BAC) library of UV-2 genome. The BAC library will help identify pathogenicity-associated genes and provide genome resource for map-based cloning and comparative genomic analysis. [Methods] High molecular weight genomic DNA was isolated from young mycelia of UV-2 strain, digested with restriction enzyme *Hind* III and size selected by PFGE. Large genomic DNA fragments were recovered and ligated to the pIndigoBAC536-S vector. The ligation product was used to transform the *Escherichia coli* strain DH10B T1 Phage-Resistant cells and the transformants were selected on a blue-white selection medium. White colonies were individually picked into 384-well microtiter plates and stored at -80 °C. [Results] We constructed a high quality deep coverage BAC library of UV-2 strain. The BAC library consisted of 10 368 clones with an average insert size of 124.4 kb and an empty clone rate of lower than 1%, covering 36.8-fold of the UV-2 genome. [Conclusion] We established a method for preparation of fungal high molecular weight genomic DNA, and constructed the first BAC library of *V. virens* genome successfully. The BAC library has been opened to researchers as a public genomic resource (<http://GResource.hzau.edu.cn>).

**Keywords:** High molecular weight DNA, Bacterial artificial chromosome, Rice, Rice false smut, *Villosiclava virens*

[*Villosiclava virens* (Cooke) Tak.]

(Rice false smut)

*Villosiclava virens* (Cooke) Tak.,

[1]

*Ustilaginoidea*

[2]

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[1], [3-4], [5], [6] 2003, UV-2 UV-12, [7], [8], [9], [10] 18S rRNA, 5.8S rRNA, ITS, PCR, 2004, [11], UV-7, 2012, [12] rep-PCR, 5, 48, BOX, REP, ERIC, [13] (Bacterial artificial chromosome, BAC), F, BAC, DNA, [14-15], BAC, (http://GResource.hzau.edu.cn)

, BAC, BAC, DNA, DNA, DNA, BAC, UV-2, BAC, 2003, BAC, 18S,

# 1 材料与amp;方法

## 1.1 稻曲病菌的培养

UV-2 (PSA) UV-2, PSA, 4 d, , 28 °C 3 d, 100 mL [16-17], 180 r/min 28 °C 2 d, DNA

## 1.2 UV-2 菌株大分子基因组 DNA 包埋块的制备

BAC [14-15,18], 1.1 50 g, 2, 0.7 mmol/L NaCl 2, 50 mL 0.7 mmol/L NaCl, (Driselase, Sigma) 10 g/L, 100 r/min 31 °C 4 h,

Miracloth (Calbiochem) , ,  
 1 500×g 15 min 40 mL 18  
 0.7 mmol/L NaCl , 1 mL , λ ladder PFG  
 1.2 mol/L , 1% marker (NEB), ,  
 (1.2 mol/L ) , , 0.5 cm  
 , 30 min; (Plug) , λ ladder PFG marker  
 40 mL 1 g/L K (Sigma) ( 1% (0.5 mg/L) , ,  
 N-lauroylsarcosine ) 50 mL , 50 °C  
 2×24 h ( K 1 )  
 Plug, 1 mmol/L PMSF ( ,  
 ) T<sub>10</sub>E<sub>10</sub> (10 mmol/L Tris-HCl  
 10 mmol/L EDTA, pH 8.0) 2 , TE  
 (10 mmol/L Tris-HCl 1 mmol/L EDTA, pH 8.0)  
 2 , 1 h, ; ,  
 TE , 4 °C

### 1.3 部分酶解及大片段 DNA 的第一次筛选

DNA *Hind*  
 ,  
 DNA ,  
 , UV-2  
 ,  
 , DNA ,  
 DNA , ,  
 : 0.2  
 0.3 U/18 min 0.4 0.6 U/15 min  
 25 μL H<sub>2</sub>O 10 μL 10×Buffer  
 10 μL 40 mmol/L Spermidine (Sigma) 5 μL  
*Hind* 1/2 Plug, 37 °C  
 ,  
 0.5×TBE  
 14 °C 1–50 s 120° 6 V/cm 18 h

### 1.4 大片段 DNA 的第 2 次筛选和洗脱回收

,  
 1  
 110–220 kb 220–300 kb  
 2 0.5×TBE 14 °C 4–4 s  
 120° 6 V/cm 18 h 1  
 ,  
 ,  
 a1 a2 (a2>a1), b1 b2 (b2>b1),  
 DNA, DNA,

### 1.5 克隆载体的制备

,  
 ,  
 pIndigoBAC536-S  
 pIndigoBAC536<sup>[18]</sup> ,  
 pHZAUBAC1<sup>[15]</sup>,  
 pHZAUBAC1 Qiagen  
 , pHZAUBAC1 *Hind*  
 FastAP (Fermentas)  
 , pIndigoBAC536-S,  
 [14–15] DNA  
 λDNA ,  
 6 mg/L 30%–40% , -80 °C ,  
 λDNA/*Hind* III ,  
 UV-2 BAC

### 1.6 BAC 文库构建

2 DNA 84  $\mu$ L DNA  
 4  $\mu$ L 2  $\mu$ L 5 U/ $\mu$ L T4 Ligase (Fermentas)  
 10  $\mu$ L T4 Buffer, 16  $^{\circ}$ C 65  $^{\circ}$ C  
 (1% +2% ) 1 h,  
 4  $^{\circ}$ C 2  $\mu$ L  
 20  $\mu$ L DH10B T1 Phage-Resistant  
 (Invitrogen) , 325 V  
 , SOC 1 h ,  
 12.5 mg/L , 80 mg/L X-gal,  
 100 mg/L IPTG LB , 37  $^{\circ}$ C  
 ,  
 4  $^{\circ}$ C ,  
 384 ,  
 -80  $^{\circ}$ C

### 1.7 文库质量分析

384 4-5 ,  
 , -Sce (NEB) ,  
 :  
 0.5 $\times$ TBE 14  $^{\circ}$ C 5-15 s 120 $^{\circ}$  6 V/cm 17 h

## 2 结果与分析

### 2.1 基因组 DNA 包埋块(Plug)的制备及部分酶解

UV-2 ,  
 , DNA Plug  
 , Plug DNA ,  
 , DNA

DNA 1A UV-2  
 DNA Hind ,  
 , UV-2 DNA Plug DNA  
 , DNA  
 DNA , 0.3 U/18 min  
 100-150 kb DNA ,  
 0.6 U/15 min ,  
 0.4 U/15 min (37  $^{\circ}$ C)  
 DNA

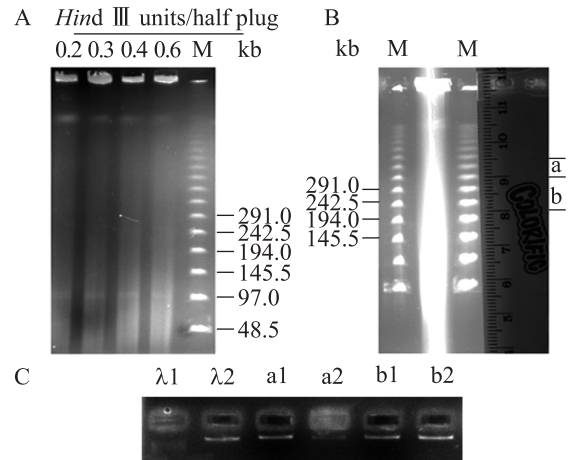


图1 稻曲病菌 UV-2 基因组 DNA 部分酶解条件比较 (A)、大片段 DNA 第 1 次筛选(B)和第 2 次筛选后回收 DNA 浓度检测(C)

Fig. 1 Comparison of partial digestion conditions of the *Villosiclava virens* UV-2 genomic DNA (A), the first screening of large fragment DNA (B) and concentration of recovered large fragment DNA after the second screening (C)

: 0.2 U 0.3 U: *Hind* III 37  $^{\circ}$ C 18 min; 0.4 U 0.6 U:  
*Hind* III 37  $^{\circ}$ C 15 min; M:  $\lambda$  ladder PFG  
 (NEB).  $\lambda$ 1  $\lambda$ 2 1 2 ng DNA ; a1 a2 b1 b2  
 1  $\mu$ L DNA.

Note: 0.2 U, 0.3 U: *Hind* III digest at 37  $^{\circ}$ C for 18 min; 0.4 U, 0.6 U: *Hind* III digest at 37  $^{\circ}$ C for 15 min; M:  $\lambda$  ladder PFG DNA marker (NEB);  $\lambda$ 1,  $\lambda$ 2 are the DNA standards of 1 and 2 ng respectively; a1, a2, b1, b2: 1  $\mu$ L DNA recovered after the second screening of large fragment DNA.

2.2 大片段 DNA 的两次筛选和回收

*Hind* 37 °C 15 min  
 18 Plug  
 DNA 1  
 1 ( 1B),  
 Plug DNA *Hind*  
 110–220 kb (a) 220–300 kb (b)  
 2 DNA  
 a b  
 a1 a2 b1 b2  
 DNA  
 a1 a2 b1 b2  
 λDNA ( 1C), a1 a2  
 b1 b2 1–2 mg/L , a2  
 1 mg/L , DNA

2.3 载体制备

pIndigoBAC536-S 10 mg/L ( 2),  
 6 mg/L,  
 30%–40% , -80 °C  
 pIndigoBAC536-S (*Hind* )

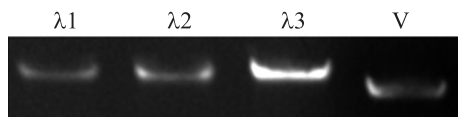


图2 载体 pIndigoBAC536-S 浓度检测  
 Fig. 2 Concentration estimation for the vector pIndigoBAC536-S

: λ1 λ2 λ3 8 16 32 ng DNA ; V: 1 μL pIndigoBAC536-S

Note: λ1, λ2, λ3 are the DNA standards of 8, 16 and 32 ng respectively; V: 1 μL pIndigoBAC536-S vector.

λDNA (30 mg/L) ,  
 λDNA ,  
 100 μL ,  
 5 ; λDNA  
 100 μL , λDNA  
 30 ng:60 ng ,  
 100 μL 800 ,  
 BAC  
 2.4 BAC 文库构建和质量检测  
 UV-2 *Hind* III DNA a1 a2  
 b1 b2 16 °C ,  
 4 °C , a1, a2  
 (a1>a2>1 200), ,  
 20 ,  
 a1 100 kb ,  
 a2 130 kb b1  
 b2  
 (b2<b1<400), b1 b2  
 a2  
 8 , 1 500  
 384  
 27 384 10 368 ,  
 -80 °C  
 168 , -*Sc*  
 168  
 1% 3  
 3  
 4  
 168  
 120–130 kb ,

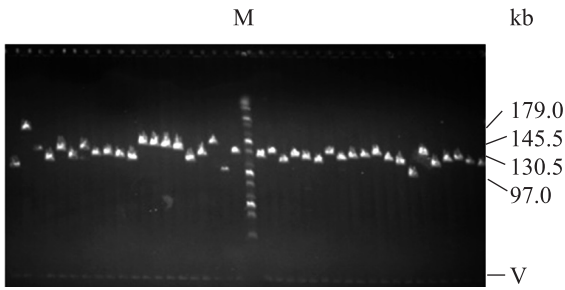


图3 UV-2 BAC 文库随机 BAC 克隆的插入片段检测  
 Fig. 3 Insert size estimation of randomly selected BAC clones of the UV-2 BAC library

Note: M: Middle range PFG marker; V: pIndigoBAC536-S vector (-7 kb).

### 3 讨论

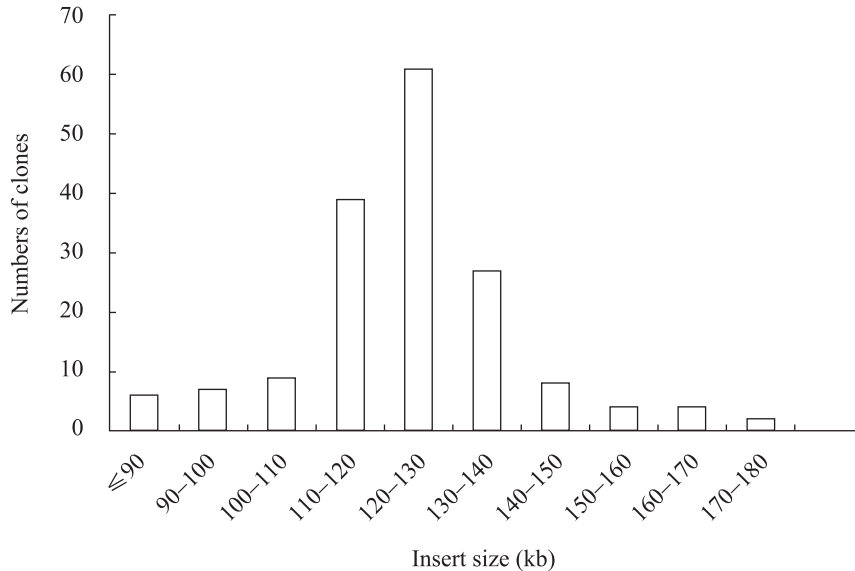


图4 UV-2 BAC 文库插入片段分析(共 168 克隆)  
 Fig. 4 Insert size analysis of the UV-2 BAC library (168 clones)

- DNA  
(BAC)  
(124.4 kb),  
UV-2 BAC  
36.8  
BAC
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