

( )

## RAPD

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( / / :                / / :                )

(RAPD)

DNA      PCR

UPGMA

/      /

( / )

/

r = /

RAPD

(Mozaffarian, 2005;                    *Cerasus vulgaris*  
.Webster & Looney, 1996)

*Cerasus*

(Section)

*P. cerasus* L.

Prunoideae

*Cerasus*

*P. avium* L.

(Rosaceae)

*Prunus*

*Prunus cerasus* L.

		<i>P. fruticosa</i>
	Eucerasus	
/		
	<i>Prunus</i>	
(FAO, 2007)		
		$2n = 4x = 32$
		$(2n = 2x = 16)$
		(Dirlewanger et al., 2007; Webster & Looney, 1996)
		$(2n = 4x = 32)$
		.2007; Webster & Looney, 1996)
		Morellos
	Amarelles	
		Kentish
(Schmidt et al., 1985)		
		(Webster & Looney, 1996)
		(Duke cherry)
		$(2n=4x=32)$
	DNA	
RAPDs	AFLPs	SSRs
(Gerlach & Stosser, 1998; Struss et al., 2003; Tavaud et al., 2001; Wunch & Hormaza, 2002)		
RAPD		
(Williams et al., 1990)		
		(Dirlewanger et al., 2007; Webster & Looney, 1996)
RAPD		
(Weeden et al., 1992)		
		(Webster & Looney, 1996)
		<i>P. gondouinii</i>
		<i>P. mahaleb</i> L.
		$(2n = 2x = 16)$
		Mahalab
		Eucerasus
		(Webster & Looney, 1996)
		(Besnard et al., 2001)
		(Oraguzie et al., 2001)
		(Zhongping, 2007)
		( )
(Gerlach & Stosser, 1998; Moreno & Trujillo, 2005; Lisek et al., 2006)	RAPD	
		( )
		RAPD
		Cherry
		(Cai et al., 2007)

...  
 nm      nm  
 DNA              )  
 (              ) DNA              (Outgroup  
 TIB              RAPD  
 MOLBIOL

*Taq* DNA

(dNTPs)              polymerase  
 MgCl<sub>2</sub>      (              ) PCR      (              )  
 μl              .(              )      (              )  
 mM   MgCl<sub>2</sub> / mM   X      PCR      DNA  
*Taq*              / μM dNTPs /      (1980) Murray & Thompson  
 .              DNA      ng      DNA

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NBDK<sub>19</sub>  
 NBSC<sub>1</sub>  
 NBSC<sub>2</sub>  
 HMSC<sub>1</sub>  
 HMSC<sub>2</sub>  
 HMSC<sub>3</sub>  
 NVSC<sub>1</sub>  
 NVSC<sub>2</sub>  
 NVSC<sub>3</sub>  
 CHSC<sub>1</sub>  
 CHSC<sub>2</sub>  
 CHSC<sub>3</sub>  
 HTSC<sub>1</sub>  
 HTSC<sub>2</sub>  
 HTSC<sub>3</sub>  
 HTSC<sub>4</sub>  
 HTSC<sub>5</sub>  
 HTSC<sub>6</sub>  
 HTSC<sub>7</sub>  
 HTSC<sub>8</sub>  
 HTSC<sub>9</sub>  
 HTSC<sub>10</sub>  
 NBDKD  
 NBSCH  
 NBDKM  
 NBDKB  
 HDZSC<sub>1</sub>  
 HDZSC<sub>2</sub>  
 HDDK<sub>1</sub>  
 HDML

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TIBM					PCR
			°C		
		BE13			°C
	(      )	TIBM BE4			°C
				Bio-Rad	
				PCR	
	(      )				(i-Cycler)
NBDK <sub>19</sub>	( / )				
			HTSC <sub>5</sub>		
				/	
					TBE
	(      )				
HMSC <sub>2</sub>	( / )				
			HMSC <sub>3</sub>	/ μg/l	
( / ) NBDK <sub>19</sub>					
HTSC <sub>3</sub>	( / )				
			HTSC <sub>5</sub>		
				/	
					UV
					DNA

NTsys (Ver 2.02)

UPGMA

1

NBDK<sub>19</sub>)

DNA

(NBSC<sub>1</sub>

TIB MOL BOIJ

(HDDK<sub>1</sub>)

CHSC<sub>3</sub> . DNA  
(NBDKB)

1

CHSC<sub>a</sub>

1. Tris Boric Acid EDTA
  2. Gel document, UVP
  3. Dice
  4. Unweighted Paired Group Method Using Arithmetic Average

RAPD

			5 →
/	TGC TCG GCT C		TIBMBA-02
/	GTG CGA GAA C		TIBMBA-03
/	GGA CGA CCG T		TIBMBA-06
	TCG GGA GTG G		TIBMBA-14
/	TCA CGT GGC T		TIBMBB-03
/	CTT CGG TGT G		TIBMBB-13
	ACA CCG TGC C		TIBMBB-17
	ACA GTA GCG G		TIBMBC-02
/	GAG GCG ATT G		TIBMBC-05
	GTC ATG CGA C		TIBMBC-09
	AAC GTC GAG G		TIBMBC-10
/	CCT GGC ACA G		TIBMBC-13
/	CTG GTG CTC A		TIBMBC-16
/	CCC AAG GGA A		TIBMBE 04
/	GGG AAG CGT C		TIBMBE 08
/	TCG GTG AGT C		TIBMBE 13
/	GGG AAA AGC C		TIBMBE-17

NVSC<sub>3</sub> ( / ) NBDKD  
 NVSC<sub>1</sub> / ( NVSC<sub>2</sub>  
 NVSC<sub>2</sub> NVSC<sub>1</sub>  
 NVSC<sub>3</sub>  
 . .  
 HMSC<sub>3</sub> HMSC<sub>2</sub> /  
 ( / )  
 . .  
 NBDKD<sub>19</sub>  
 . .  
 HMSC<sub>1</sub>  
 HMSC<sub>3</sub> HMSC<sub>2</sub>  
 HMSC<sub>1</sub>  
 . .  
 NVSC<sub>1</sub> CHSC<sub>2</sub> CHSC<sub>1</sub> HMSC<sub>1</sub>  
 NVSC<sub>2</sub>  
 . .  
 HTSC<sub>7</sub> HTSC<sub>6</sub> ( / ) CHSC<sub>2</sub> CHSC<sub>1</sub>

( )

A large grid of vertical lines forming a stepped pattern. The grid consists of approximately 100 columns and 50 rows of lines. The lines are black and set against a white background. They create a series of diagonal steps that slope upwards from left to right. The spacing between the lines varies slightly, with more space between groups of lines than between individual lines within a group. The overall effect is a dense, geometric texture.

NBDKD ( / ) . HTSC<sub>9</sub> HTSC<sub>8</sub>

NBDKM	.	HTSC <sub>10</sub> HTSC <sub>6</sub>
HTSC <sub>1</sub>	.	HTSC <sub>10</sub> HTSC <sub>7</sub>
HTSC <sub>4</sub> HTSC <sub>3</sub> HTSC <sub>2</sub>	.	HDZSC <sub>2</sub> HDZSC <sub>1</sub>
.	HTSC <sub>5</sub>	( / )

( / ) HTSC2 HTSC3 HDZSC<sub>2</sub>

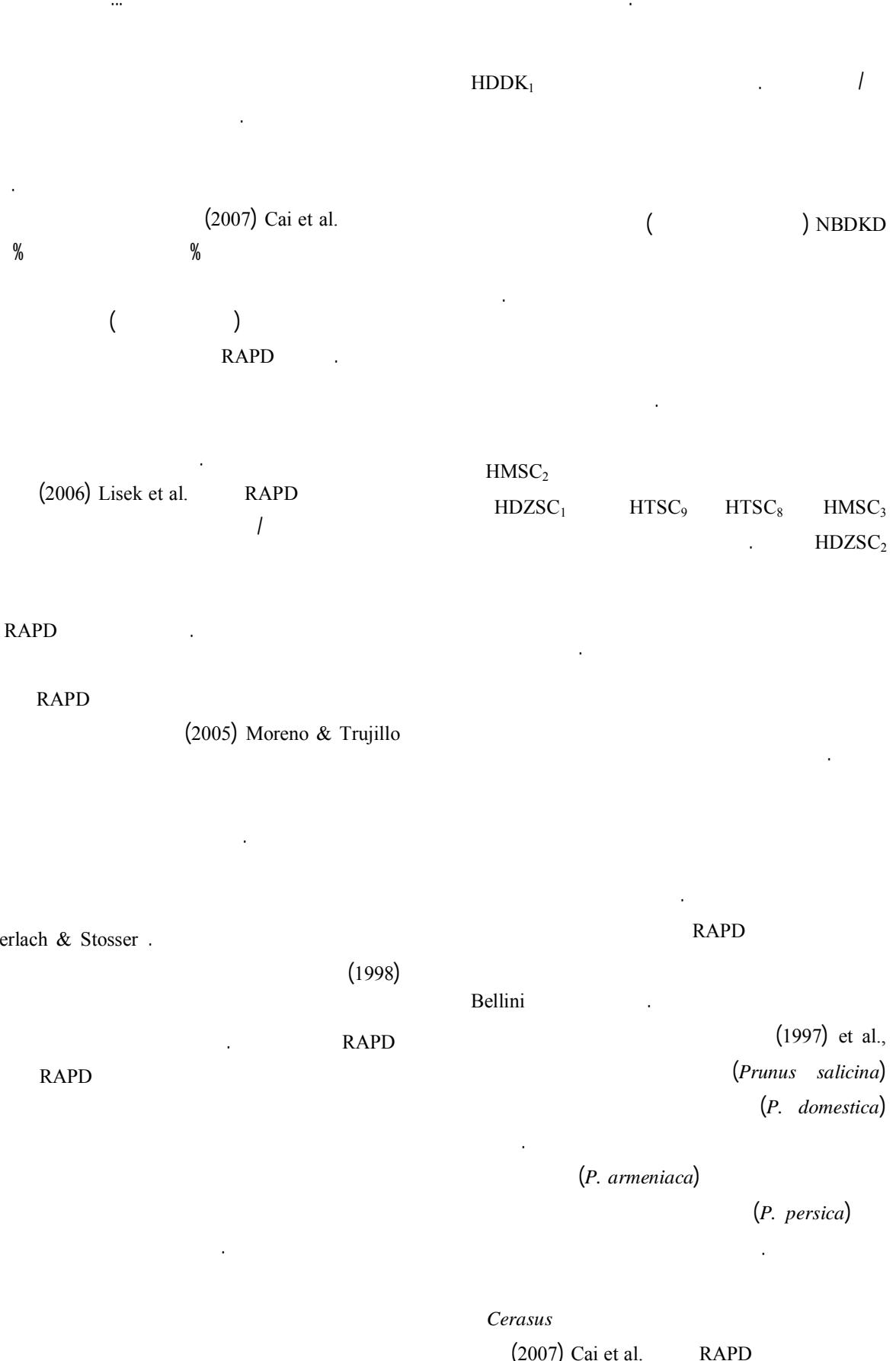
NBDKB

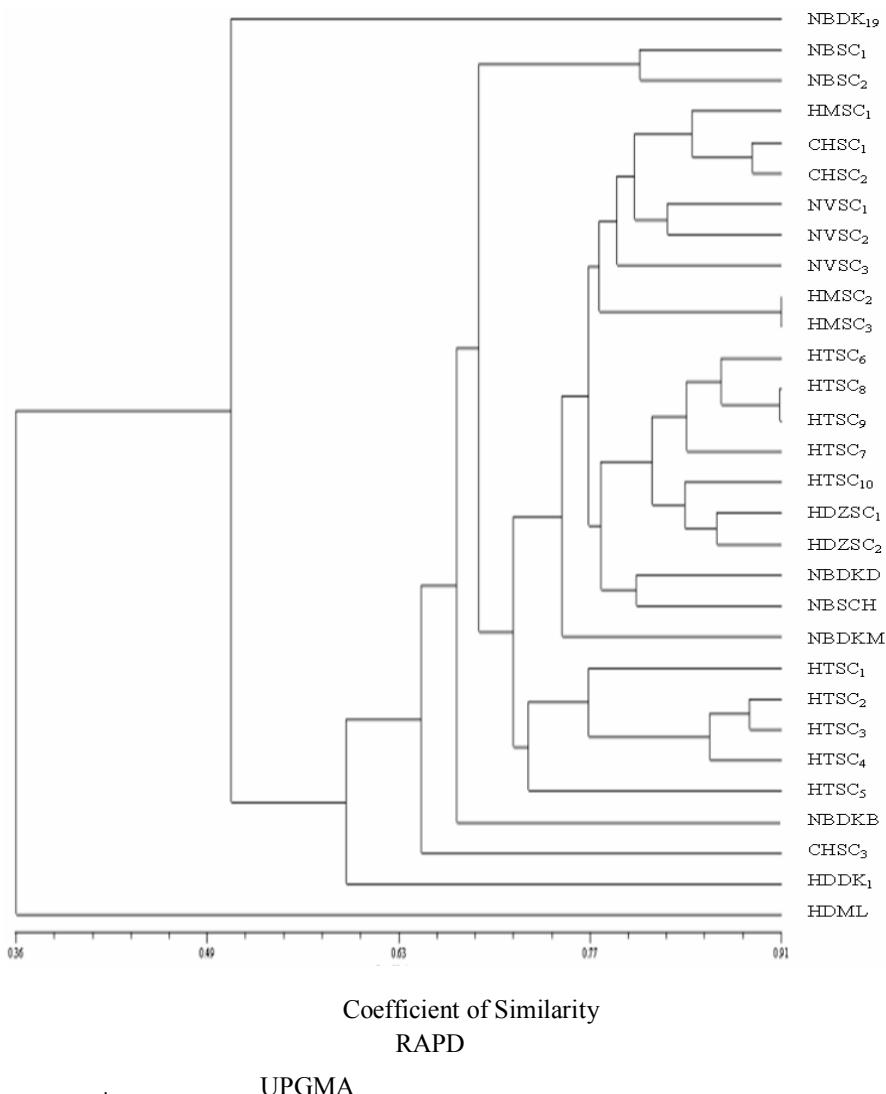
CHSC<sub>3</sub>

HDZSC<sub>1</sub>

NBSCH NBDKD

$$\text{CHSC}_2 \quad / \quad \text{CHSC}_1 \quad \text{CHSC}_3 \quad . \quad /$$





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