

About the followings picked up, please check again.

For example,

- 1) the abbreviations should be written with full names when it appeared as first time.
- 2) some mistake in inputting.
- 3) the abbreviations should be written with full names when it appeared as first time.
- 4) and 10) the abbreviations should be written with full names under the table or refer to M &M, and so on.
- 5) the abbreviations can be used directly when they appeared from second time.
- 6) the references in the text should be refer to the instruction of the manuscript making to authors.
- 7) ,8) and 9)

Primer	No of alleles (A)	Monomorphic	Polymorphic bands (B)	B/A(%)	PIC	RP
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- 11) enhances not only the monitoring of genetic changes in improved gene pools, but also the effort of germplasm conservation and Utilization.

Anyway, the manuscript should be carefully checked again before publication.

1) ISSR, SRAP and RAPD

2) ~~(This technique depends on combination between two types of primers. Table 3-a shows the sequence of primers that were producible, and Table 3-b~~ shows the combinations of the SRAP primers (Pedraza-Garcia, 2010) that produced polymorphic bands.

3) **Keywords:** Wheat, genetic diversity, ISSR, SRAP, RAPD

4)

Symbol of combination	me	Em	Symbol of combination	Me	Em
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5) Resolving power (RP) was calculated according to the following equation;

$$P = \sum 1 - (2 * (0.5 - f))$$

Where f is the frequencies and P is polymorphism

Polymorphism information content (PIC) values were obtained using the following formula:

$$PIC = 1 - \sum P_{ij}^2.$$

6) These results are in agreement with those of El-Assal and Gaber, (2012); Akkaya et al. (1992); Sofalian et al. (2008) who confirmed that di-nucleotide repeat primers were found at lower polymorphic levels.

7)

Primer	No of alleles	Monomorphic	Polymorphic bands	Polymorphic%	PIC	RP
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8)

Markers	No of alleles	Monomorphic	Polymorphic	%	RP	PIC
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9)

Primers	No of Allels	Monomorphic	Polymorphic bands	Pol.%	RP	PIC
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10)

V	ISSR	SRAP	RAPD	Total Bands	Average Bands
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11) Developing effective indicators for genetic diversity of cultivated plants not only enhances the monitoring of genetic changes in improved gene pools, but also the effort of germplasm conservation and Utilization.