

IDENTIFICATION OF GOOD COMBINERS IN EARLY MATURING × HIGH YIELDING CULTIVARS OF INDICA RICE (*ORYZA SATIVA* L.)*

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Key words: Indica rice, Line × tester design, GCA, SCA, Grain yield

Abstract

Line × tester analysis involving seven early maturing lines and four high yielding testers, made to produce 28 F₁s hybrids. The F₁ data revealed that both additive and non-additive gene were important in controlling yield contribution in Indica rice. Variances were significant for GCA and SCA effects for all the characters indicating the significance of both additive and non-additive genes. There was high proportion of non-additive genes for the expression of different traits. Cultivars Anjali, MTU-7029 and BPT-5204 were identified as best general combiners for yield and yield traits among the parents. The most promising specific combiners for grain yield and its contributing traits were Govind × HUR 105, NDR 97 × HUR 4-3, Anjali × HUR 4-3, NDR 97 × MTU 7029, Vandana × BPT 5204, Shanthi × HUR 105, Anjali × MTU 7029 and Shanthi × BPT 5204.

Introduction

Rice (*Oryza sativa* L., 2n = 24), a member of Poaceae (Gramineae) is world's most important staple food crop that feed over half of the global population. It is cultivated in tropical and subtropical regions. Rice is grown in more than 114 countries, over an area of 161.4 m ha in a wide range of ecosystems under varying temperature and water regimes with the production of 466.7 mt (on milled basis) (FAO 2011). The total area under rice in India is about 45 m ha (22% of cropped area) with an annual production of 103.4 mt and contributes 25% to total agricultural GDP. The world population will grow up to 8.5 billion till 2030 for that about 40% more rice will be required to feed growing population (Khush 2006). Therefore, increasing its productivity is of high importance in breeding programmes. Plant breeding strategies should be made for selection of plants to achieve expected level of heterosis as well as the specific combining ability. Reduced plant height, effective tillers per hill, increased kernel number per panicle, increased 1000-grain weight and higher yield are the most important rice traits to be improved in breeding programmes (Paterson *et al.* 2005).

For, high yielding varieties of crop plant for qualitative and quantitative traits, plant breeders often face problems in selecting parents and crosses. Combining ability analysis is one of the most valuable tools to ascertain the combining ability effects and helps in selecting the desirable parents for making crosses and crosses for the exploitation of heterosis (Khatun *et al.* 2010). Line × tester analysis provides information related to general combining ability (GCA) and specific combining ability (SCA) effects of parents. This information is helpful in estimating various types of gene action (Chaudhury and Sasmal 1992, Ganapathy *et al.* 2007 and Akter *et al.* 2010). Therefore in the present investigation, an attempt was made to study combining ability (GCA and SCA) in rice cultivars for different traits to identifying good combiners, desirable cultivars to developing high yielding rice varieties.

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Materials and Methods

The experiment was conducted at Agricultural Farm, Institute of Agricultural Sciences, Banaras Hindu University, Varanasi, UP, India during *Kharif* 2008 - 2010. The material for the present investigation comprised 11 genetically diverse cultivars (Table 1), in which seven lines were used as male parents *viz.* Anjali (P₁), HUR 3022 (P₂), Vandana (P₃), Govind (P₄), Annada (P₅), NDR 97 (P₆) and Shanthi (P₇) and 4 lines (broad based varieties) were used as testers (female parents) *viz.*, HUR 4-3 (P₈), HUR 105 (P₉), MTU 7029 (P₁₀) and BPT 5204 (P₁₁) of rice chosen on the basis of early maturity and high yielding potential, respectively. All the parents were crossed in all possible combinations in line × tester mating to produce 28 F₁s. The experiment comprising 28 F₁s along with their parents were evaluated in randomized block design (RBD) with three replications during *Kharif* seasons of 2008 - 2010. Twenty days old seedlings of each parents and F₁s (one seedling per hill) were transplanted in field. The spacing between row to row 20 cm and plant to plant 15 cm was maintained in 3 m row length in each plot. All recommended cultural practices and plant protection measures were followed.

Data were recorded for nine quantitative characters *viz.*, panicle initiation, days to 50% flowering, days to maturity, plant height (cm), effective tiller/plant, 1000-grain weight (g), spikelet fertility (%) and grain yield/plant (g) on five randomly selected plant from all F₁s and parents in each replication. The randomly selected plants were tagged before maturity. Days to 50 per cent flowering and days to maturity were recorded on plot basis. The average of the five plants was taken as the mean value for each treatment and the mean values were used for statistical analysis. The pooled data were analyzed according to ANOVA (Steel and Torrie 1980). Combining ability (GCA and SCA) analysis was done using line × tester method (Kempthorne 1957) and the analysis was done using statistical software WINDOSTAT Ver. 8.3.

Results and Discussion

In the study, mean squares of parents and crosses showed significant difference ($p < 0.01$) between cultivars for all the traits under study (Table 2). The difference between parents indicated that the experimental materials possessed considerable variability and they are suitable for genetic studies. The ANOVA for combining ability indicated that variances due to lines, testers and interaction effects (line × tester) were highly significant for all the characters. These trends indicated the presence of both additive and non-additive gene action. These results conform the findings of Sarker *et al.* (2002).

Further analysis of GCA/SCA variance showed that the nature of gene action as non-additive for all the traits. The SCA variances was higher for all the characters indicating the role of non-additive gene, which results from dominance, epistasis and various other interaction effects with non fixable genetic variation. The presence of greater non-additive gene offers scope for exploiting hybrid vigour through heterosis breeding and hence these parents can be further exploited for production of good commercial variety. Khatun *et al.* (2010) and Amudha and Thiyagarajan (2011) also reported that non-additive gene action was greater than additive gene action for yield and most of the yield contributing traits.

The proportional contribution of lines, testers and their interaction for eight different traits are shown in Table 3. It is evident that lines played important role towards plant height (72.43%), panicle initiation (60.99%), days to 50% flowering (60.13%), days to maturity (59.02%) and grain yield per plant (37.98%) indicating predominant maternal influence on these traits. Testers were more important for 1000-grain weight (56.33%), effective tiller per plant (30.07) and grain yield per plant (13.57%). It revealed that preponderance influence for the traits. The contribution of paternal and maternal interaction (line × tester) were high for spikelet fertility (62.78), grain yield

Table 1. Variety wise pedigree and features of Indica rice cultivars.

Name of cultivars	Parentage and year of release	Source of cultivars	Specific features of Indica rice cultivars
Anjali (P1)	RR-19-2 × RR-149-1129 (2001)	Central Rice Research Institute, Cuttack, Odisha, India	Semi dwarf, 90-95 days for maturity, grains - short bold, white; tolerant to drought, resistant to blast and SB; yield: 30-35 q/ ha.
HUR 3022(P2)	IR 36 × HR 137 (2005)	Banaras Hindu University, Varanasi, Uttar Pradesh, India	Semi dwarf- 100 cm; 105 days for maturing, grains – long slender, fine, tolerant to BLB, SB, leaf and neck blast; yield: 48-50 q/ha.
Vandana (P3)	C-22 × Kalakeri (1992)	Central Rice Research Institute, Cuttack, Odisha, India	Tall (95-155 cm), early maturing, grains – long bold, white, mod. resistance to termitite and SB , blast and BS; yield: 30-35 q/ha.
Govind (P4)	IR 20 × IR 2 (1983)	Tamil Nadu Agricultural University, Tamil Nadu, India	Dwarf, early maturing, long Slender grain, flowering in 75 days, yield: 35-40.
Annada (P5)	MTU-15 × Yaikaku Nantoku (1988)	Central Rice Research Institute, Cuttack, Odisha, India	110-112 days to maturing, short statured grains, mod. resistant to blast and stem borer, GM, BPH; yield: 45- 50 q/ha.
NDR 97 (P6)	Nagina 22 × Ratna (1992)	Narendra Deva Univ. of Agric. & Tech., Faizabad, Uttar Pradesh, India	Medium, early maturing, medium slender, flowering in 70 days, yield: 40- 45 q/ha.
Shanthi (P7)	Ratna × IR-36 (2002)	Acharya N G Ranga Agriculture University, Andhra Pradesh, India	Semi dwarf, 120-130 days for maturity, grains – long slender, resist. to blast, brown spot and, WBPH; yield : 50 q/ha.
HUR 4-3 (P8)	Mutant of Lanjhi; (2009)	Banaras Hindu University, Varanasi, Uttar Pradesh, India	Semi dwarf 90-100 cm, 135-140 days for maturity, grains- LS; resist. to leaf roller and BPH, yield: 52-54 q/ ha
HUR 105 (P9)	Mutant of MPR 7-2 (2009)	Banaras Hindu University, Varanasi, Uttar Pradesh, India	Semi dwarf (100-102 cm), 130-135 days for maturity, grains - LS; tolerant to leaf and neck blast, BS, SB; yield: 58-60 q/ ha.
MTU7029 (P10)	Vasista × Mashuri (1982)	Acharya N G Ranga Agriculture University, Andhra Pradesh, India	Dwarf, 140 days for maturity, medium slender grain shape, resistant to BLB and tolerant to many diseases, yield: 55-60 q/ ha
BPTS204 (P11)	GEB 24 × T N-1 + Mahsuri (1986)	Acharya N G Ranga Agriculture University, Andhra Pradesh, India	Dwarf, low grain weight, medium grain length, fine and medium slender decorticated grain shape, yield: 55-65 q/ ha, 140-150 days for maturity

Table 2. Analysis of variance for combining ability for eight different characters.

Source of variation	d.f.	Panicle initiation	Days to 50 % flowering	Days to maturity	Plant height (cm)	Effective tiller / plant	1000-grain wt. (g)	Spikelet fertility (%)	Grain yield/ plant (g)
Replication	2	0.228	0.642	0.399	0.503	0.217	1.175	0.869	0.107
Genotypes	38	767.443**	722.311**	726.008**	938.836**	9.116**	13.345**	116.275**	76.543**
Parents	10	1088.511**	956.920**	977.303**	1338.681**	3.423**	9.675**	69.912**	82.184**
Parents vs Crosses	1	1653.930**	1341.372**	1256.894**	694.625**	36.307**	30.1502*	266.506**	160.215**
Crosses	27	615.696**	612.489**	613.274**	799.789**	10.218**	14.082**	127.883**	71.351**
Line	6	204.272**	224.069**	245.847**	1707.385**	2.717**	8.102**	79.942**	37.471**
Testers	3	229.195**	226.988**	225.342**	612.691**	4.020**	10.407**	26.155**	8.063**
L x T effects	18	276.810**	280.163**	294.445**	254.034**	7.089**	7.012**	120.434**	51.853**
Error	76	1.081	1.225	1.166	1.767	0.549	0.908	1.550	1.493
σ^2 GCA		66.310	65.811	64.290	92.831	1.131	2.313	7.372	6.251
σ^2 SCA		91.903	92.980	97.760	84.082	2.181	2.031	39.620	16.791
σ^2 GCA/ σ^2 SCA		0.722	0.713	0.662	1.103	0.524	1.132	0.193	0.371

*, **p < 0.05 and 0.01, respectively.

Table 3. Proportional contribution (%) of lines, testers and their interactions to total variance.

Source	Panicle initiation	Days to 50% flowering	Days to maturity	Plant height (cm)	Effective tiller/ plant	1000-grain wt. (g)	Spikelet fertility (%)	Grain yield/ plant (g)
Due to lines	60.99	60.12	59.02	72.43	23.67	10.48	31.63	37.98
Due to testers	9.03	9.38	8.97	6.40	30.07	56.33	5.59	13.57
Due to line x testers	29.97	30.49	32.01	21.18	46.26	33.19	62.78	48.45

per plant (48.45%) and effective tiller per plant (46.26%) and were low for rest of all the characters. These results are similar to those reported by Sarker *et al.* (2002) and Rashid *et al.* (2007).

The estimates of general combining ability for parents (Table 4) revealed that none of the parents was a consistently good general combiner except Anjali (male parent) for all the traits studied. However, the female parents MTU 7029 and BPT 5204 were found to be good general combiner for grain yield per plant. High GCA effect of BPT 5204 for grain yield per plant was associated with in high GCA effect for effective tiller per plant, days to maturity, days to 50% flowering and panicle initiation, and significant negative GCA effect for 1000-grain weight and plant height. Better combining ability of MTU 7029 for grain yield per plant was due to its high GCA effect for spikelet fertility, days to maturity, days to 50% flowering and panicle initiation, and negative significant GCA for effective tiller per plant. Likewise, male parent HUR 105 appeared to be good general combiner for days to maturity, plant height and effective tiller per plant and 1000-grain weight while, male parent HUR 4-3 was good general combiner for spikelet fertility and negative for rest of the traits.

Among the male parents, Anjali was the better general combiner for all the traits under study and other male parents Annada and Vandana were the best general combiner for grain yield. High GCA effect of Annada for grain yield per plant may be associated with its high GCA effects for effective tiller per plant, days to maturity, days to 50% flowering and panicle initiation and plant height. In Vandana, high GCA effects for grain yield per plant, spikelet fertility and plant height were due to its negative GCA effects for days to maturity, days to 50% flowering and panicle initiation; Shanthi for effective tiller per plant, plant height, days to maturity, days to 50% flowering and panicle initiation and HUR 3022 for panicle initiation, days to 50% flowering and days to maturity.

Negative GCA effects for plant height and positive GCA effect for other traits are desirable. Dwarf plants are required to protect the crop from lodging. Earliness is a desirable trait for aerobic and non aerobic condition of rice cultivation. Parents showing good combining ability were HUR 4-3, Vandana, Govind, NDR 97 (Table 4); these genotypes can be utilized for developing short duration variety.

On the basis of mean performance and GCA effects with respect to yield contributing traits, the lines Anjali, Vandana, Annada, Shanthi was a good general combiners for all yield contributing traits and tester MTU 7029 and BPT 5204 are good for grain yield per plant, panicle initiation, days to 50% flowering, days to maturity, effective tiller per plant and spikelet fertility. Therefore, these genotypes with desirable genes for yield and yield contributing traits can be used as potential donors for improvement of yield potential or yield with early maturity.

In contrast to general combining ability effects, the specific combining ability effects represents dominance and epistatic component of variation, that is not fixable in nature. But if crosses showing high SCA effect involved either both or one good general combining parents, they can be successfully exploited for varietal improvement and expected to throw stable performing transgressive segregants carrying fixable gene effects (Vegeed *et al.* 2011). The estimates of specific combining ability effects of 28 cross combinations for different yield contributing characters are presented in Table 5. It is observed that out of 28 crosses 14 crosses exhibited positive and significant SCA effects for grain yield per plant.

The promising specific combinations for grain yield and other traits were found in Govind × HUR 105, NDR 97 × HUR 4-3, Anjali × HUR 4-3, NDR 97 × MTU 7029, Vandana × BPT 5204, Shanthi × HUR 105, Anjali × MTU 7029 and Shanthi × BPT 5204. It is observed that majority of

Table 4. *Per se* performance and GCA effects associated with each parental line for eight morphological traits.

Parents	Panicle initiation	Days to 50% flowering	Days to maturity	Plant height (cm)	Effective tiller / plant	1000-grain wt. (g)	Spikelet fertility(%)	Grain yield/plant (g)
Lines								
Anjali (P 1)	66.800	82.040	109.170	139.443	9.890	21.747	82.767	20.554
HUR-3022 (P 2)	5.979**	6.788**	6.005**	16.402**	0.502*	1.449**	4.230**	3.70**
Vandana (P 3)	78.480	92.627	120.280	123.200	10.050	21.340	86.237	26.383
Govind (P 4)	14.774**	13.531**	12.749**	-5.283**	-1.086**	-0.651*	-7.036**	-1.65**
Annada (P 5)	67.870	82.223	110.530	142.887	8.560	24.497	83.467	21.245
NDR 97 (P 6)	-8.968**	-9.059**	-9.036**	23.698**	0.012	-0.628*	1.155**	1.52**
Shanthi (P 7)	63.293	78.213	107.007	90.633	7.770	19.440	91.613	18.474
SE	-14.356**	-14.109**	-14.908**	-14.023**	-0.835**	0.487	3.147**	-5.07**
CD at 0.05	67.990	92.483	121.540	97.833	10.267	22.667	80.490	27.356
0.01	5.731**	6.000**	6.882**	-10.693**	0.742**	-0.141	-0.951**	3.57**
Testers	58.433	73.270	101.993	81.593	8.660	20.437	91.120	19.184
HUR 4-3 (P 8)	-13.127**	-13.351**	-12.312**	-11.492**	-0.795**	-0.336	2.106**	-0.13
HUR 105 (P 9)	81.953	96.607	126.923	110.870	8.560	22.533	77.947	24.427
MTU 7029 (P 10)	9.967**	10.199**	10.575**	1.396**	1.460**	-0.180	-2.651**	-1.95**
BPT 5204 (P 11)	0.3001	0.319	0.3117	0.384	0.2139	0.2751	0.3594	0.3527
SE	0.602	0.640	0.6250	0.769	0.4289	0.5516	0.7206	0.7071
CD at 0.05	0.801	0.852	0.8324	1.024	0.5712	0.7345	0.9596	0.9417
HUR 4-3 (P 8)	96.127	110.310	139.973	111.287	8.750	20.607	87.787	29.637
HUR 105 (P 9)	-6.771**	-6.976**	-6.874**	-1.415**	-0.698**	-0.216	1.657**	-2.758**
MTU 7029 (P 10)	95.837	108.267	136.297	111.770	10.677	22.703	87.290	30.453
BPT 5204 (P 11)	0.116	0.434	0.878**	6.268**	0.760**	2.220**	-2.313**	-0.318
SE	111.740	124.630	152.250	89.550	11.223	19.720	93.517	31.167
CD at 0.05	2.013**	1.922**	1.463**	0.093	-1.232**	0.266	0.990**	1.610**
0.01	110.443	123.923	153.533	84.520	9.290	18.250	87.817	33.450
Testers	4.642**	4.620**	4.533**	-4.946**	1.169**	-2.270**	-0.334	1.466**
SE	0.227	0.241	0.235	0.290	0.1617	0.208	0.271	0.266
CD at 0.05	0.455	0.484	0.472	0.581	0.3242	0.416	0.544	0.534
0.01	0.606	0.644	0.629	0.774	0.4318	0.555	0.725	0.712

*, **p < 0.05 and 0.01%. Upper and lower values indicate *per se* performance and GCA effects, respectively.

Table 5. SCA effects of 28 F₁s for eight different traits.

F ₁ s	Panicle initiation	Days to 50 % flowering	Days to maturity	Plant height (cm)	Effective tiller / plant	1000-grain wt. (g)	Spikelet fertility (%)	Grain yield/ plant (g)
Anjali × HUR 4-3	16.81**	16.58**	17.25**	3.65*	0.37*	-1.64**	2.60**	3.03**
Anjali × HUR 105	1.24**	2.25**	2.92**	-2.82**	-1.71**	1.64**	5.22**	-4.28**
Anjali × MTU 7029	-16.53**	-16.88**	-18.03**	2.57**	3.27**	0.47*	-1.56**	1.13**
Anjali × BPT 5204	-1.52**	-1.95**	-2.13**	-3.39**	-1.93**	-0.48*	-6.27**	0.12
HUR 3022 × HUR 4-3	-18.36**	-17.79**	-17.03**	-18.78**	-2.10**	0.28	7.14**	-6.22**
HUR 3022 × HUR 105	5.78**	6.29**	5.53**	8.55**	0.13	0.44	3.86**	2.28**
HUR 3022×MTU 7029	7.96**	7.85**	8.23**	12.36**	0.40*	-1.15**	-7.26**	1.73**
HUR 3022 ×BPT 5204	4.63**	3.65**	3.26**	-2.13**	1.57**	0.43*	-3.38**	2.21**
Vandana × HUR 4-3	11.69**	12.61**	12.67**	10.73**	1.33**	0.64**	-0.49	3.59**
Vandana × HUR 105	-6.51**	-7.94**	-8.09**	-5.70**	1.27**	-2.71**	3.58**	-3.85**
Vandana × MTU 7029	-3.02**	-2.79**	-1.64**	5.89**	-1.59**	0.73**	0.34	-1.21**
Vandana × BPT 5204	-2.15**	-1.87**	-2.94**	-10.92**	-1.02**	1.34**	-3.43**	1.46**
Govind × HUR 4-3	4.55**	4.09**	3.76**	7.81**	-0.78**	-0.14	2.64**	-0.20
Govind × HUR 105	-1.15**	-1.10**	-0.92**	-4.73**	0.98**	-1.54**	-9.08**	6.50**
Govind × MTU 7029	0.49*	0.52*	-0.02	-7.69**	-0.63**	1.45**	4.40**	-6.10**
Govind × BPT 5204	-2.91**	-3.50**	-2.82**	4.62**	0.43**	0.23	2.03**	-0.18
Annada × HUR 4-3	-16.35**	-16.56**	-17.42**	-4.21**	0.33*	0.13	-5.16**	-4.12**
Annada × HUR 105	2.91**	2.34**	1.45**	-5.39**	-0.44**	0.96**	2.56**	-0.23
Annada × MTU 7029	9.54**	8.97**	9.40**	5.55**	-1.09**	-0.37	1.21**	2.26**
Annada × BPT 5204	3.95**	5.25**	6.52**	4.05**	1.20**	-0.72**	1.38**	2.08**
NDR 97 × HUR 4-3	3.48**	3.01**	3.62**	1.59**	-0.16**	2.50**	3.69**	6.26**
NDR 97 × HUR 105	-2.73**	-1.51**	-1.01**	0.89**	0.57**	-0.64**	-8.43**	-1.77**
NDR 97 × MTU 7029	2.40**	1.55**	0.260	-4.99**	-1.22**	-1.90**	5.30**	1.78**
NDR 97 × BPT 5204	-3.61**	-3.06**	-2.88**	5.69**	0.81**	0.04	-0.56*	-6.28**
Shanthi × HUR 4-3	-1.81**	-1.93**	-2.86**	2.39**	1.01**	-1.77**	-10.43**	-2.34**
Shanthi × HUR 105	0.46*	-0.32	0.07**	9.20**	-0.81**	1.84**	2.27**	1.35**
Shanthi × MTU 7029	0.13	0.79**	1.79**	-13.68**	0.86**	0.77**	-2.06**	0.40
Shanthi × BPT 5204	1.25**	1.48**	0.99**	2.08**	-1.06**	-0.84**	10.23**	0.58*
SE ±	0.60	0.64	0.62	0.76	0.43	0.55	0.71	0.71
CD at 0.05	1.20	1.28	1.25	1.53	0.86	1.10	1.44	1.41

*, **, p < 0.05 and 0.01%, respectively.

the crosses with high SCA effects for grain yield were involved with high \times low combiners indicating additive \times dominance type of gene interactions for expression of traits. Similarly crosses involving high \times high general combiners showed high GCA effects which could be due to predominance of additive \times additive gene action. Some crosses having low \times low general combiners showed high SCA effect suggesting the epistatic gene action may be operating due to genetic diversity in the form of heterozygous loci. Thus in majority of the crosses, high SCA effects for grain yield were attributed to additive and dominance gene actions and a few cross attributed to epistatic interactions. These results were conformity of the finding of Pradhan and Singh (2008) and Amudha and Thiyagarajan (2011).

Negative estimates for SCA may be desirable for few traits like earliness and dwarf plant type, cross combination showing desirable SCA for earliness are Anjali \times MTU 7029, HUR 3022 \times HUR 4-3 and Annada \times HUR 4-3. The desirable cross combinations for dwarf plant type are HUR 3022 \times HUR 4-3 and Vandana \times BPT 5204. These crosses are showing best negative and significant specific combining ability for plant height and earliness. The results are in conformity with the findings of Sarker *et al.* (2002) and Akter *et al.* (2010). Effective tiller per plant, spikelet fertility and 1000-grain weight are positively correlated with grain yield per plant; these traits may be used as selection criteria for screening of high yielding cultivars per variety. The Crosses showing high SCA effects for yield contributing traits were Anjali \times HUR 105, Anjali \times HUR 4-3, NDR 97 \times HUR 4-3 and Shanthi \times HUR 105. NDR 97 \times HUR 4-3 cross possesses favourable alleles for yield contributing traits to exhibit significant SCA effect for morphological characters like panicle initiation, 1000-grain weight and spikelet fertility (Table 5).

It is concluded from the present study that the importance of good \times good general combiners exhibiting high SCA effect can be utilized for improvement through single plant selection in segregating generations. But in crosses having high SCA effects due to good \times poor general combiners may be improved through population improvement. The cross showing high SCA effects involving poor \times poor general combiners may be utilized for heterosis breeding programme.

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(Manuscript received on 8 September, 2012; revised on 30 October, 2013)