

REFERENCES

ภาษาไทย

- ชาญวิทย์ เกษตรศิริ. 2542. อุบัติปะนิพัทธ์และการเมือง. 112-129.
- รันชัย สังฆ์สุข. 2541. การสะสมโพลีนและน้ำตาลของข้าวที่คีมเมื่อได้รับภาวะที่คีมและแล้ง.
- วิทยานิพนธ์วิทยาศาสตร์ธรรมชาติ จุฬาลงกรณ์มหาวิทยาลัย 2541.
- สุเทพ ลิ่มทองกุล. 2531. การปรับปรุงคุณภาพข้าว. ศูนย์วิจัยข้าวปทุมธานี สถาบันวิจัยข้าว กรมวิชาการเกษตร 4-9.
- สงกรานต์ จิตรากร. 2543. ความหลากหลายทางชีวภาพของข้าวในประเทศไทย: เอกสารประกอบการสัมมนาทางวิชาการข้าวแห่งชาติ: การวิจัยและพัฒนาพันธุ์ข้าวด้วยเทคโนโลยีชีวภาพ. 26-38.
- สำเริง ตัน. สัมภาษณ์, 24 มกราคม 2543.
- อภิชาติ วรรณวิจิตร และคณะ. 2544. เทคโนโลยีชีวภาพกับการปรับปรุงพันธุ์ข้าว: วิทยาศาสตร์และเทคโนโลยีกับข้าวไทย. 79-122.
- เอกสารงาน ชีวสิสุกุล. 2542. พันธุ์ข้าวต้านทานโรค แมลง ໄล์เดือนฝอย ทนดินเบรื้อง ดินเค็ม และทนแล้ง. ฝ่ายถ่ายทอดเทคโนโลยี สถาบันวิจัยข้าว กรมวิชาการเกษตร. 98 หน้า.
- เยาวนุช เวศร์ภาดา และคณะ. 2543. ข้าววัดน้ำรวมแห่งชีวิต. 103-110.

ภาษาอังกฤษ

- Ahmed,J. and Gupta, S. 1991. Germination and growth of some salt-resistant selections in high salt concentration solutions. *Int. Rice Res. News.* 16(5) :15.
- Akbar, M 1986. Breeding for salinity tolerance in rice. .p.39-63. In salt affected soils of Pakistan, India and Thailand. The International Rice Research Institute, Los Banos, The Philippines
- Akbar, M. and Senadhira,D. 1988. Sensitivity of rice seedlings to salinity. International Rice Research Institute, los Banos, The Philippines
- Anishetty,N.M. 1983. Status of rice genetic resources. *Rice Germplasm Genetic Conservation Workshop IRRI*, pp.25.
- Apuya,N.R.,Frazier,B.L.,Keim,P.,Roth,E.J.and Lark,K.G.1988. Restriction fragment length polymorphisms as genetic markers in soybean, *glycine max(L.)merrill*. *Theor. Appl. Genet.* 75: 889-901.

- Arunin, S. 1984. Characteristics and management of salt affected soils in the northeast of Thailand. Ecology and management of problem soils in Asia. 336-351.
- Baishya , S., Sachdev , A., Johari ,RP.and Mehta ,SL. 2000 RAPD analysis of aromatic and non-aromatic rice (*Oryza sativa L.*) J. of Plant Biochem and Biotech. 9 (1) :23-26.
- Bong, B.B., Tobita, S., Bermawie, N. and Senboku, T. 1996. Salt Tolerance of Cultivated Rice Varieties from Vietnam. Jircus J. 3 : 75-83.
- Bohra, J.S. and Doerffling, K. 1993. Potassium nutrition of rice (*Oryza sativa L.*) varieties under NaCl salinity. Plant Soil 152:299-303.
- Chang,T.T. 1976. Manual on genetic conservation of rice germplasm for evaluation and utilization. IRRI Los Banos, Manila Philippines, pp. 77.
- Chang,T.T. 1976a. The origin, evaluation, cultivation, dissemination and diversification of Asian and African rices. Euphytica 25 : 425-441.
- Chang,T.T 1976 b. The rice cultures. Phil. Trans. Raval. Soc. London B275 : 143-157.
- Chang,T.T 1979. Rice..In Simmonds N.W.(ed) Evaluation of crop plants Longmans. London. New York, pp.98-104.
- Chang,T.T., and Bardenas, E.A. 1965. Morphology and varietal characteristics of the rice plant. IRRI Tech. Bull. 4 Los Banos, Philippines, pp. 40.
- Chatterjee,D.1948. A modified key and enumeration of species of *Oryza* India. J.Agric.Sci.18:185-192.
- Claes, B., R. DeKeyser, R. Villarroel, M. Van den Bulche, G. Bauw, M. Van Montagu and A, Caplan. 1990. Characterization of a rice gene showing organic specific expression in response to salt stress and drought. The Plant Cell 2:19-27
- Dawson ,IK., Chalmers , KJ., Waugh , R. and Powell , W. 1993.Detection and analysis of genetic –variati0n in *Hordeum-spontaneum* populations from Israel using RAPD markers. Mol Ecol 2 :(3) 151-159.
- Flowers, T.J. and Reo, A.R. 1981. Variability in the resistance of sodium chloride salinity within rice (*Oraza sativa L.*) varieties. New Phytol. 88:363-373.
- Godwin,I.D., Sangduen, N. and Kunanuvatchaidach R.1997. RAPD polymorphism among variant and phenotypically normal rice somaclonal progenies. Plant cell Rep 16 : (5) 320-324.

- Gomez, K.A., Tuazon,D, and Nano, N.E.1979. Germplasm Bank Information Retrieval System. IRRI Research Paper Series No. 45 Los Banos, Philippines, pp 24.
- Gregorio, G.B. 1997. Tagging salinity tolerance genes in rice using amplified fragment length polymorphism(AFLP). Ph.D.thesis, University of the Philippines, Los Banos, Philippines.
- Huh, M.K. and Huh,H.W. 2001. Genetic diversity of *Raphanus sativus* var. hortensis f. raphanis troides in Korea using AFLP markers. Korean J. Genetic 23 : (1) 45-53.
- Iwaki, S. 1956. Salt infury in rice plants. Mem. Ehime, Univ. (Agric.) 2:1-156 (in Japanese with English summary).
- Kaddah, M. T. and Fakhary, S. I. 1961. Tolerance of Egyptian rice to salt : 1, salinity effects when applied continuously and intermittently at different stages of growth after transplanting. Soil Sci. 91:113-120
- Katiyar, S. K., Tan, Y. ,Zhang, Y. ,Huang, B. ,Xu, Y., Zhao, L., Huang, N., Khush, G.S. and Bennett J.1994. Identification of RAPD markers linked to the gene controlling gall midge resistant against all biotypes in China. Rice Genetics Newsletter 11 : 128-131.
- Keim,P., Diers B.W., Olson,T.C. and Shomaker,R.C. 1990. RFLP mapping in soybean: Association between marker loci and variation in quantitative traits. Genetics 126:735-742.
- Khush, G.S.1975. Rice. In Robert C.King (ed) Handbook of Genetics, Vol 2 Plenum Press, pp.31-58.
- Ko, H.L., Cowan, D.C., Henry,R.J., Graham,G.C., Blankeney, A.B. and Lewin,L.G.1994. Random amplified polymorphic DNA analysis of Australian rice varieties. Euphytica. 80 :179-189.
- Kohyama, K., Wichaidit,P., Pramojanee,P., Sukchan,S. and Wada,H.1993. Salinity in the watershed of Northeast Thailand. In: ADRC (ed)" Research Activities of ADRC ontributed to Agricultural Development in Northeast Thailand " Khon Kaen, Thailand.
- Kolchinsky, A., Kolesnikova, M. and Ananiev, E. 1991. Portraying of plant genomes using polymerase chain reaction amplification of ribosomal 5S genes. Genome 34 :1028-1031.

- Limpinuntana, V. 1978. Physiological aspects of adaptation of rice (*Oryza sativa L.*) and barley (*Hordeum vulgare L.*) to low oxygen concentrations in the root environment. Ph.D. thesis, University of Western Australia, Australia.
- Litt, M. and Luty, J.A. 1989. A hypervariable microsatellite revealed by in vitro amplification of dinucleotide repeat within the cardiac muscle actin gene. Am.J. Hum. Genet. 44: 388-396.
- Lynch, M. 1990. The similarity index and DNA fingerprinting. Molecular Biology and Evolution 7: 478-484.
- Mackill, D.J. 1995. Plant Genetic Resources Classifying Japonica Rice Cultivars with RAPD Markers. Crop Sci 35 : 889-894.
- Mitsuchi, M., Widhidit, P. and Jeungnijirund, S. 1989. Soil of the Northeast Plateau, Thailand. Tech. Bull. Trop. Agric. Res. Center, 25. 55pp
- Monna, L., Miyao, A., Inoue, T., Fukuoka, S., Yamazaki, M., Zhong, H-S., Sasaki, T. and Minobe, Y. 1994. Determination of RAPD markers in rice and their conversion into sequence tagged sites (STSs) and STS-specific primers. DNA Res. 1 :139-148.
- Mundy, J. and Chua, N.H. 1988. Abscisic acid and water stress induced expression of a novel rice gene. EMBO J. 7: 2279-2286.
- Narayanan, K.K. and Sree Rangasamy, S.R. 1991. Genetic analysis for salt tolerance in rice. In Rice Genetics II 1:167-173.
- Ng, Q.N. 1978. A biosystematic study of Asian rice. Ph.D. Thesis University of Birmingham U. K.
- Oka, H.I. 1964. Pattern of interspecific relationships and evolutionary dynamics in *Oryza*. Rice Genetics and Cytogenetics (Proc. Symp. held by IRRI) Elsevise, Amsterdam, pp.71-90.
- Oka, H.I. 1974. Experimental studies on the origin of cultivated rice. Genetics 78 :475-486.
- Oka, H.I. 1975. Rice in Asia. Univ. Tokyo Press, Tokyo pp,277-287.
- Porteres, R. 1956. Taxonomic agrobotanique des riz cultives *O.sativa* Linn et. *O. glaberrima* Steudel. J. Agric Trop. Bot. Appl. 3 : 341-384.
- Saghai-Maroof, M.A., Soliman, K.M., Jorgensen, R.A., Allard, R.W. 1984. Ribosomal DNA spacer-length polymorphisms in barley : Mendelian inheritance,chromosomal

- location, and population dynamics.Proc. Natl. Acad. Sci. USA Dec.,81(24):8014-8018.
- Sampath, S.1962. The genus *Oryza* : its taxonomy and species interrelationships.Oryza 1:1-29.
- Sampath, S. 1973. Origins of cultivated rices. Indian J. Genet. Plt. Breed.33 :157-161.
- Sano,Y., Morishima,H ,and Oka, H.I. 1980. Intermediate perennial annual populations of *O. perennis* found in Thailand and their evolutionary significance. Bot. Mag. Tokvo 93 : 291-305.
- Sharma, S.D., and Shastry, S.V.D. 1965. Taxonomic studies in genus *Oryza* VI A. Modification classification. Indian J. Genet. Plt. Breed. 25 :173-178.
- Tancred, S.J., Zeppa,A. and Graham,G.C. 1994.The use of PCR (RAPD) technique in improving plant variety rights description of a new Queensland apple(*Malus domestica*) cultivar. Aus. J. Exp. Agri. 34 :665-667.
- Tanksley, SD.,Young ,N.D.,Paterson,A.H. and Bonierbale, M.W. 1989. BioTech. 7:257.
- Tanksley,S.,Causse,M.,Fulton,T.,Ahn,N.,Wang,Z.,Wu,K.,Xiao,J.,Yu,Z.,Second,G.and McCouch,S. 1992. A high density molecular map of the rice genome. Rice Genet Newsletter. 9 :111-115.
- Tautz,D.1989.Hypervariability of simple sequences as a general source for polymorphic DNA markers. Nucl. Acids Res. 17(16);6463-6471.
- Tautz,D. and Renz, M.1984.Simple sequences are ubiquitous repetitive components of eukaryotic genomes. Nucl. Acids Res. 12(10) : 4127-4138.
- Thomson, D. and Henry, R.1993. Use of DNA from Dry Leaves for PCR and RAPD Analysis. Plant Mol Biol Rep. 11(3): 202-206.
- Valdes,A.M.,Slatkin, M. and Friermer,N.B. 1993. Allele frequencies at microsatellite loci : the step wise mutation model revisited. Genetics 133: 737-749.
- Vajrabhaya, M., Thanapaisal, T. and Vajrabhaya, T. 1989. Development of salt tolerance line of Khao Dawk Mali105 and Leuang Pratew 123 rice cultivars through tissue culture. Plant Cell. Rep. 8: 411-414.

- Verma , SK. , Khanna ,V. and Singh ,N. 1999. Random amplified polymorphic DNA analysis of Indian scented basmati rice germplasm for identification of variability and duplicate accessions , if any Electrophoresis. 20 (8) :1786 -1789.
- Vos,P.,Hoger,R.,Bleeker,M.,Reijans,M.,VandeLee,T.,Hornes,M.,Fritters,A.,Pot,J.,Peleman ,J.,Kuiper,M and Zabeau,M.1995. AFLP: A new concept for DNA fingerprinting. Nucl.Acids. Res.23 : 4407-4414.
- Wang, Z.Y. and Tanksley, S.D. 1989. Restriction fragment length polymorphism in *Oryza sativa L.* Genome32 : 1113-1118.
- Weber, J.L. and May, P.E. 1989. Abundant class of human DNA polymorphisms which can be typed using the polymerase chain reaction. Am.J. Hum. Genet.44:388-396.
- Weng, M.L., Chen, Y., Zhang, D., Jia, J., Li, C., Jin,D., Hguyen, H.T., Wang,B. 2000. Rice seed identification by computerized DNA fingerprinting techniques. Plant & Animal genome VIII Conference. 1:9-12.
- Williams, J.G.K.,Kubelik,A.R.,Livak, K.J.,Rafalski,J.A and Tingey,S.V. 1990. DNA polymorphism amplified by arbitrary primers are useful as genetic markers. Nucl. Acids Res. 18 : 6531-6535.
- Williams, J.G.K., Hanafey, M.K., Rafalski, J.A. and Tingey, S.V. 1993. Genetic analysis using random amplified polymorphic DNA markers Methods in enzymology 218 : 709-740.
- Wolff, K., Schoen, E.D.and Peter-Van Rijn J.1993. Optimizing the generation of random amplified polymorphic DNA in chrysanthemum. Theor. Appl. Genet. 86 :1033-1037.
- Yamaguchi-Shinozaki, K., Mundy, J. and Chua, N.H. 1989. Four tightly linked *rab* genes are differentially expressed in rice. Plant Mol. Biol. 14: 29-39.
- Yeo, A.R.,Yeo, M.E., Flowers, S.A. and Flowers,T.J. 1990. Screening of rice(*Oryza sativa L.*) genotypes for physiological characters contributing to salinity resistance, and their relationship to overall performance. Theor. Appl. Genet. 79: 377-384.
- Yu, L X. and Nguyen, H.T. 1994. Genetic variation detected with RAPD markers among upland and lowland rice cultivars. Theor. Appl. Genet. 87 :668-672.

Zheng , K.L., Shen,B. and Qian, H.R. 1991.DNA polymorphisms generated by arbitrary primed PCR in rice.Rice Genetics Newsletter. 8 :134-136.

APPENDICES

Appendix A

Analysis of amplification products by RAPD

1. Scoring of polymorphic products of RAPD analysis in 6 rice cultivars by random primer

1.1 primer X6

size (bp)	POK	KDML	KTH	LPT	LDP	IR
1517	0	0	1	1	1	1
1400	0	0	0	0	0	0
1300	0	0	1	0	1	0
1200	0	0	0	0	0	0
1100	0	0	1	0	1	0
1000	0	1	0	1	0	1
900	0	0	0	0	0	1
850	0	0	0	0	0	0
800	0	0	1	0	1	1
750	0	0	0	0	0	0
700	0	1	0	1	1	1
650	1	0	0	0	0	0
600	0	0	0	0	0	1
550	0	0	0	0	1	0
500	0	0	0	0	0	0
450	0	0	0	0	0	0
400	0	0	0	0	0	0
350	0	0	0	0	0	0
300	0	0	0	0	0	0
275	0	0	0	0	0	0
250	0	0	0	0	0	0
200	0	0	0	0	0	0

Pokkali = POK Khao Dawk Mali 105 = KDML

Khao Tah Haeng 17 = KTH Leuang Pratew 123 = LPT

Look Daeng Pattani = LDP IR 28 = IR

1.2 primer X8

size (bp)	POK	KDML	KTH	LPT	LDP	IR
1517	0	0	0	0	0	0
1400	0	0	0	0	0	0
1300	0	1	0	0	0	0
1200	1	0	0	1	1	1
1100	0	0	0	0	0	0
1000	0	0	0	1	0	0
900	0	0	1	0	1	0
850	0	0	0	0	0	0
800	0	1	1	1	1	1
750	0	0	0	0	0	0
700	1	1	0	1	0	1
650	0	0	0	0	0	0
600	1	0	0	1	0	0
550	0	0	0	0	0	0
500	1	0	0	0	0	0
450	0	0	0	0	0	0
400	1	0	0	1	0	1
350	0	0	0	0	0	1
300	1	0	0	0	0	1
275	0	0	0	0	0	0
250	1	0	0	1	0	1
200	0	0	0	0	0	0

Pokkali = POK Khao Dawk Mali 105 = KDML

Khao Tah Haeng 17 = KTH Leuang Pratew 123 = LPT

Look Daeng Pattani = LDP IR 28 = IR

1.3 primer X9

size (bp)	POK	KDML	KTH	LPT	LDP	IR
1517	1	0	0	0	0	0
1400	0	0	0	0	0	0
1300	0	0	0	0	0	0
1200	0	0	0	0	0	1
1100	0	0	0	0	0	0
1000	1	1	0	0	0	0
900	0	0	0	0	0	0
850	0	0	0	0	0	0
800	0	1	1	1	1	1
750	0	0	0	0	0	0
700	1	0	0	0	0	0
650	0	0	1	1	1	1
600	1	1	0	0	0	0
550	0	0	0	0	0	0
500	1	0	0	0	1	0
450	0	1	1	1	0	0
400	0	0	0	0	1	1
350	1	1	1	1	0	0
300	0	0	0	0	0	1
275	1	0	0	0	0	0
250	1	0	0	1	0	0
200	0	0	0	0	0	0

Pokkali = POK Khao Dawk Mali 105 = KDML

Khao Tah Haeng 17 = KTH Leuang Pratew 123 = LPT

Look Daeng Pattani = LDP IR 28 = IR

1.4 primer X10

size (bp)	POK	KDML	KTH	LPT	LDP	IR
1517	0	0	0	0	0	0
1400	0	0	0	0	0	0
1300	0	0	0	0	0	0
1200	0	0	0	0	0	0
1100	0	0	0	0	0	0
1000	0	0	0	0	0	0
900	0	0	0	0	0	0
850	0	0	0	0	0	0
800	0	0	0	1	0	0
750	0	0	0	0	0	0
700	0	0	0	0	0	0
650	0	0	0	0	0	0
600	1	0	0	1	0	0
550	0	0	1	0	0	0
500	1	1	0	1	0	1
450	0	0	0	0	1	0
400	1	0	0	1	0	0
350	0	0	0	0	0	0
300	1	0	0	1	0	0
275	0	0	0	0	0	0
250	1	0	0	1	0	0
200	1	0	0	0	0	0

Pokkali = POK Khao Dawk Mali 105 = KDML

Khao Tah Haeng 17 = KTH Leuang Pratew 123 = LPT

Look Daeng Pattani = LDP IR 28 = IR

1.5 primer C1

size (bp)	POK	KDML	KTH	LPT	LDP	IR
1517	0	0	0	0	0	0
1400	0	0	0	0	0	0
1300	0	0	0	0	0	0
1200	0	0	1	1	0	0
1100	0	0	1	0	0	0
1000	0	0	0	1	1	1
900	1	0	0	1	0	0
850	0	0	0	0	0	0
800	0	0	0	1	0	0
750	0	0	0	0	0	0
700	0	0	1	0	0	1
650	0	0	0	0	0	0
600	0	0	1	1	0	1
550	0	0	0	0	0	1
500	1	0	0	1	0	0
450	0	0	0	1	0	1
400	1	0	0	1	0	0
350	1	0	1	0	0	0
300	0	0	0	0	0	1
275	0	0	0	0	0	0
250	1	0	0	1	0	0
200	0	0	0	0	0	0

Pokkali = POK Khao Dawk Mali 105 = KDML

Khao Tah Haeng 17 = KTH Leuang Pratew 123 = LPT

Look Daeng Pattani = LDP IR 28 = IR

2. Scoring of polymorphic products of RAPD analysis in 12 rice cultivars by random primer X9

size (bp)	POK	KDML	KTH	LPT	LDP	GR	IR	MN	NP	YY	FT	LNP
1517	0	0	1	1	1	1	0	1	0	0	0	0
1000	0	1	1	1	1	1	0	1	1	0	1	0
900	0	0	0	0	0	0	0	1	0	0	0	0
800	0	1	1	1	1	1	0	1	0	0	0	0
700	0	0	0	0	0	0	0	1	0	0	0	0
650	0	0	1	1	1	1	1	1	1	1	1	1
600	1	1	0	0	0	0	0	0	0	0	0	0
550	0	0	1	0	0	0	1	1	0	0	0	0
500	1	1	0	1	1	1	0	0	1	1	1	1
450	0	1	1	1	0	0	0	0	0	0	1	1
400	0	0	0	0	1	1	1	0	0	0	0	0
350	1	1	1	1	1	1	0	1	1	1	0	0
300	0	0	0	0	0	0	1	0	0	0	0	0
275	1	0	0	0	0	0	0	0	0	0	0	0
250	0	1	0	1	0	1	0	1	1	0	1	0
200	0	0	1	0	0	0	0	0	0	0	0	0

Pokkali	= POK	Khao Dawk Mali 105	= KDML
Khao Tah Haeng 17	= KTH	Leuang Pratew 123	= LPT
Look Daeng Pattani	= LDP	Gow Ruang 88	= GR
IR 28	= IR	Muey Nawng 62 M	= MN
Nahng Pa-yah 132	= NP	Yah Yaw	= YY
Foi Tawng	= FT	Leb Nok Pattani	= LNP

APPENDIX B

Analysis of variance for salt tolerance criteria

Completely randomized design

Replication(Rep.) = 7

Treatment: Variance(Var.) = 12

Rice cultivars (12)

Pokkali	=	POK	Khao Dawk Mali 105	=	KDML
Khao Tah Haeng 17	=	KTH	Leuang Pratew 123	=	LPT
Look Daeng Pattani	=	LDP	Gow Ruang 88	=	GR
IR 28	=	IR	Muey Nawng 62 M	=	MN
Nahng Pa-yah 132	=	NP	Yah Yaw	=	YY
Foi Tawng	=	FT	Leb Nok Pattani	=	LNP

1. Leaf damage (%) data of Table 3.9

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	20.00	5.26	19.05	27.73	28.57	37.50	15.00
KDML	19.05	8.00	10.53	21.05	18.18	14.29	15.79
KTH	5.88	5.26	16.67	16.67	0.00	13.33	17.39
LPT	9.52	9.52	18.18	13.04	9.09	19.05	16.00
LDP	14.29	14.29	16.67	19.05	16.67	15.00	11.11
GR	22.22	18.18	26.09	12.50	11.11	6.67	14.29
IR	47.37	52.17	45.83	52.94	50.00	50.00	45.45
MN	59.09	60.00	57.14	45.00	38.89	40.91	45.00
NP	64.29	53.33	41.18	68.75	46.15	35.29	28.57
YY	42.86	40.00	40.00	58.38	41.18	40.91	40.00
FT	41.18	31.58	30.00	31.82	44.44	35.00	61.54
LNP	38.10	41.67	45.45	41.18	34.78	50.00	52.38
Rep. totals	383.85	339.26	366.79	408.11	339.06	357.95	362.52
Rep. means	31.99	28.27	30.57	34.01	28.25	29.83	30.21

Analysis of variance for leaf damage (%) – Table 3.9

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	20013.75	1819.43	28.43**
Error	72	4607.55	63.99	
Total	83	24621.30		

cv=26.3%, **=significant at 1% level

Table of variance means for leaf damage (%) (average over 7 replications)

Var.	Ranks	Means
POK	6	21.87 ^b
KDML	3	15.27 ^{ab}
KTH	1	10.74 ^a
LPT	2	13.49 ^{ab}
LDP	4	15.30 ^{ab}
GR	5	15.87 ^{ab}
IR	11	49.11 ^d
MN	12	49.43 ^d
NP	10	48.22 ^{cd}
YY	8	43.33 ^{cd}
FT	7	39.37 ^c
LNP	9	43.37 ^{cd}
Mean		30.45

Means followed by a common letter are not significantly different at the 5% level by DMRT.

2. Plant height (cm) data of Table 3.9

2.1 Plant height (cm) under 0 dS/m (control)

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	130.67	127.67	127.00	118.00	116.33	129.67	127.00
KDML	107.00	92.67	103.33	99.00	86.67	101.33	103.00
KTH	101.33	108.33	107.33	112.00	103.33	101.00	108.00
LPT	109.00	121.00	117.00	95.67	96.33	105.67	110.00
LDP	108.00	104.33	110.67	102.33	99.67	96.67	104.33
GR	97.67	107.33	102.67	106.33	98.67	112.67	100.33
IR	69.33	69.67	64.00	72.00	69.00	72.00	69.67
MN	112.00	106.33	114.00	114.00	114.33	95.00	111.00
NP	94.33	94.33	98.67	90.33	100.33	94.00	98.33
YY	96.00	99.33	92.00	105.67	108.67	104.00	108.00
FT	107.67	114.00	100.00	101.33	98.00	104.00	108.00
LNP	102.67	89.67	92.00	99.33	105.67	96.67	100.33
Rep. totals	1235.67	1234.66	1228.67	1215.99	1197.00	1213.01	1247.66
Rep. means	102.97	102.89	102.39	101.33	99.75	101.08	103.97

Analysis of variance for plant height (control)

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	12487.98	1135.27	33.02**
Error	72	2475.66	34.38	
Total	83	14963.64		

cv = 5.7%, * * = significant at 1% level

Table of variance means for plant height (cm) (average over 7 replications)

Var.	Ranks	Means
POK	12	125.19 ^g
KDML	4	99.00 ^{bcd}
KTH	9	105.90 ^{def}
LPT	10	107.81 ^{ef}
LDP	7	103.71 ^{c-f}
GR	6	103.67 ^{c-f}
IR	1	69.38 ^a
MN	11	109.52 ^f
NP	2	95.76 ^b
YY	5	101.95 ^{b-e}
FT	8	104.71 ^{c-f}
LNP	3	98.05 ^{bc}
Mean		102.06

Means followed by a common letter are not significantly different at the 5% level by DMRT.

2.2 Plant height (cm) under 8 dS/m

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	116.67	117.33	125.67	119.00	122.00	117.33	120.00
KDML	92.00	96.00	86.67	96.33	86.67	98.00	100.00
KTH	101.00	108.00	93.67	93.33	110.00	101.00	98.33
LPT	97.67	105.33	83.67	101.67	103.33	102.33	100.00
LDP	104.00	98.67	105.33	94.00	102.33	95.00	88.33
GR	100.33	96.00	107.00	96.33	91.33	92.67	101.67
IR	51.00	55.33	60.00	49.33	57.00	60.00	57.67
MN	90.67	76.00	74.67	86.00	81.00	76.67	86.33
NP	72.33	67.00	65.33	61.67	60.00	69.33	75.00
YY	78.33	85.00	93.67	88.00	82.00	90.67	80.67
FT	86.33	90.67	80.67	87.00	80.33	78.33	72.67
LNP	72.67	70.67	66.67	77.33	71.67	57.00	66.00
Rep. totals	1063.00	1066.00	1043.02	1049.99	1047.66	1038.33	1046.67
Rep. means	88.58	88.83	86.92	87.50	87.31	86.53	87.22

Analysis of variance for plant height

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	24093.94	2190.36	66.35**
Error	72	2377.02	33.01	
Total	83	26470.96		

cv=6.6%, * * =significant at 1% level

Table of variance means for plant height (cm) (average over 7 replications)

Var.	Ranks	Means
POK	12	119.71 ^f
KDML	7	93.67 ^d
KTH	11	100.76 ^e
LPT	10	99.14 ^{de}
LDP	9	98.24 ^{de}
GR	8	97.90 ^{de}
IR	1	55.76 ^a
MN	4	81.62 ^c
NP	2	67.24 ^b
YY	6	85.48 ^c
FT	5	82.29 ^c
LNP	3	82.29 ^b
Mean		87.56

Means followed by a common letter are not significantly different at the 5% level by DMRT.

3. Tillering data of Table 3.9

3.1 Tillering under 0 dS/m (control)

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	6.33	7.33	6.00	10.67	8.67	9.67	10.67
KDML	11.33	15.67	13.67	10.33	10.00	10.00	14.67
KTH	15.33	6.67	6.33	11.67	10.33	7.67	10.00
LPT	10.67	9.00	8.00	10.67	7.00	11.33	8.67
LDP	8.67	9.67	12.67	12.33	11.67	10.67	12.33
GR	7.33	9.00	10.00	9.00	7.67	9.67	11.00
IR	17.00	14.00	16.00	13.67	14.33	15.33	16.33
MN	7.00	6.67	7.00	7.33	6.67	9.67	9.67
NP	13.00	10.33	13.67	10.33	13.00	15.33	11.67
YY	14.67	11.33	9.33	6.67	9.00	9.67	10.00
FT	12.00	8.33	9.33	11.33	10.33	11.33	13.33
LNP	11.00	15.33	15.67	14.67	11.67	11.33	12.33
Rep. totals	134.33	123.33	127.67	128.67	120.34	131.67	140.67
Rep. means	11.19	10.28	10.64	10.72	10.03	10.97	11.72

Analysis of variance for tillering(control)

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	362.6835	32.9712	8.63**
Error	72	275.2171	3.8225	
Total	83	637.9006		

cv=18.1% **=significant at 1% level

Table of variance means for tillering (average over 7 replications)

Var.	Ranks	Means
POK	2	8.48 ^{ab}
KDML	9	12.24 ^{de}
KTH	5	9.71 ^{abc}
LPT	4	9.33 ^{abc}
LDP	8	11.14 ^{cde}
GR	3	9.10 ^{abc}
IR	12	15.24 ^f
MN	1	7.72 ^a
NP	10	12.48 ^e
YY	6	10.10 ^{bcd}
FT	7	10.85 ^{cde}
LNP	11	13.14 ^e
Mean		10.79

Means followed by a common letter are not significantly different at the 5% level by DMRT.

3.2 Tillering under 8 dS/m

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	7.33	6.67	5.67	5.67	7.67	6.00	8.33
KDML	8.00	7.33	8.67	7.00	8.00	6.67	8.33
KTH	8.33	9.00	8.33	9.33	7.67	7.00	10.00
LPT	9.33	9.67	10.67	9.67	6.67	9.00	6.67
LDP	7.00	6.67	5.33	9.33	5.67	10.67	12.67
GR	8.00	8.00	6.67	9.33	7.00	7.67	7.67
IR	6.33	5.33	5.67	3.67	5.00	6.33	8.33
MN	3.33	2.67	4.00	4.33	2.50	3.67	3.67
NP	3.67	4.00	2.33	2.00	3.00	2.00	4.33
YY	4.33	3.67	5.67	4.00	4.67	5.00	3.67
FT	5.00	5.67	5.33	4.67	4.00	8.33	6.33
LNP	4.67	5.00	4.33	3.67	4.33	3.67	6.00
Rep. totals	75.32	73.68	72.67	72.67	66.18	76.01	86.00
Rep. means	6.28	6.14	6.06	6.06	5.52	6.33	7.17

Analysis of variance for tillering

Source of variation	Degrees of Freedom	Sum of squares	Mean square	F
Var.	11	316.0347	28.7304	17.19**
Error	72	120.3128	1.6710	
Total	83	436.3475		

cv=20.8%. ** =significant at 1% level

Table of variance means for tillering (average over 7 replications)

Var.	Ranks	Means
POK	7	6.76 ^{cd}
KDML	8	7.71 ^{de}
KTH	11	8.52 ^e
LPT	12	8.81 ^e
LDP	10	8.19 ^{de}
GR	9	7.76 ^{de}
IR	6	5.81 ^{bc}
MN	2	3.45 ^a
NP	1	3.05 ^a
YY	3	4.43 ^{ab}
FT	5	5.62 ^{bc}
LNP	4	4.52 ^{ab}
Mean		6.22

Means followed by a common letter are not significantly different at the 5% level by DMRT

4. Shoot/ root ratio data of Table 3.10

4.1 Shoot/ root ratio under 0 dS/m (control)

Var.	Rep.1	Rep.2	Rep.3	Rep. 4	Rep. 5	Rep. 6	Rep. 7
POK	6.76	5.98	3.09	4.39	4.25	3.48	6.49
KDML	5.39	7.49	4.83	4.62	4.08	4.51	7.08
KTH	4.77	4.17	4.02	3.20	4.66	4.67	4.01
LPT	3.86	4.05	3.85	3.66	7.99	2.66	4.43
LDP	1.92	2.57	4.78	4.53	1.43	1.10	5.91
GR	3.70	3.77	5.05	1.92	5.43	3.78	7.02
IR	9.18	6.08	5.19	2.86	6.13	4.88	7.76
MN	4.66	4.78	3.78	5.49	4.14	3.99	5.56
NP	2.60	4.97	3.63	3.06	2.00	7.70	4.16
YY	3.77	2.72	1.58	2.44	3.52	1.77	5.25
FT	6.22	4.58	3.08	1.86	3.53	4.98	8.60
LNP	4.05	4.08	4.16	4.10	3.88	4.09	4.06
Rep. totals	56.88	55.24	47.04	42.13	51.04	47.61	70.33
Rep. means	4.74	4.60	3.92	3.51	4.25	3.97	5.86

Analysis of variance for shoot/root ratio

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	54.5761	4.9615	2.11*
Error	72	168.9248	2.3462	
Total	83	223.5010		

cv=34.7% , * =significant at 5% level

Table of variance means for shoot/root ratio (average over 7 replications)

Var.	Ranks	Means
POK	10	4.92
KDML	11	5.43
KTH	5	4.21
LPT	6	4.36
LDP	2	3.18
GR	7	4.38
IR	12	6.01
MN	8	4.63
NP	3	4.02
YY	1	3.01
FT	9	4.69
LNP	4	4.06
Mean		4.41

Means followed by a common letter are not significantly different at the 5% level by DMRT

4.2 Shoot/ root ratio under 8 dS/m

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	4.46	4.89	3.93	3.67	3.79	4.01	3.85
KDML	5.49	4.92	4.89	3.57	4.51	5.56	4.33
KTH	4.05	4.17	4.00	3.71	3.48	3.32	3.19
LPT	2.10	2.72	3.06	3.84	2.86	3.13	2.97
LDP	3.45	2.02	2.84	3.24	2.25	2.98	4.08
GR	3.75	3.73	3.40	4.52	3.16	3.12	3.69
IR	6.00	5.84	4.31	5.13	4.76	5.49	3.49
MN	3.64	2.38	4.51	5.22	6.30	4.96	5.24
NP	3.71	3.52	3.39	2.62	3.32	5.27	3.42
YY	3.21	3.02	2.82	1.78	3.21	3.26	2.18
FT	2.06	2.89	3.04	3.14	1.55	2.59	3.54
LNP	3.64	3.33	3.14	2.76	5.93	1.96	4.78
Rep. totals	45.56	43.43	43.33	43.20	45.12	45.65	44.76
Rep. means	3.80	3.62	3.61	3.60	3.76	3.80	3.73

Analysis of variance for shoot/root ratio

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	47.15236	4.28658	6.93**
Error	72	44.51380	0.61825	
Total	83	91.66616		

cv = 21.2% , ** = significant at 1% level

Table of variance means for shoot/root ratio (average over 7 replications)

Var.	Ranks	Means
POK	9	4.09 ^{cd}
KDML	11	4.75 ^{de}
KTH	8	3.70 ^{bc}
LPT	3	2.95 ^{ab}
LDP	4	2.98 ^{ab}
GR	6	3.62 ^{abc}
IR	12	5.00 ^e
MN	10	4.61 ^{de}
NP	5	3.61 ^{abc}
YY	2	2.78 ^{ab}
FT	1	2.69 ^a
LNP	7	3.65 ^{abc}
Mean		3.70

Means followed by a common letter are not significantly different at the 5% level by DMRT.

5. Panicles/plant data of table 3.10

5.1 Panicles/plant under 0 dS/m (control)

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	5.33	6.33	4.33	6.33	6.67	10.67	9.33
KDML	10.33	14.00	11.00	11.33	14.33	10.00	10.33
KTH	9.67	5.00	5.33	7.33	7.00	6.33	9.67
LPT	10.33	7.33	7.00	6.67	5.00	9.00	5.67
LDP	6.67	6.00	7.33	8.33	7.33	2.67	8.33
GR	6.67	5.67	9.33	4.67	6.00	8.33	8.33
IR	12.00	10.33	13.00	8.67	10.67	9.00	11.00
MN	6.67	5.00	5.33	6.67	5.00	4.00	7.33
NP	8.00	7.00	8.00	7.33	5.67	8.67	9.67
YY	8.33	9.00	6.67	5.67	6.33	7.67	3.67
FT	8.00	5.67	6.00	8.00	6.33	4.00	8.33
LNP	6.33	7.00	8.33	8.00	8.33	7.00	7.00
Rep. totals	98.33	88.83	91.65	89.00	88.66	87.34	98.66
Rep. means	8.19	7.40	7.64	7.42	7.39	7.28	8.22

Analysis of variance for panicles/plant

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	231.6840	21.0622	7.53**
Error	72	201.4843	2.7984	
Total	83	433.1683		

cv=21.9%, **=significant at 1% level

Table of variance means for panicles / plant (average over 7 replications)

Var.	Ranks	Means
POK	5	7.00 ^a
KDML	12	11.62 ^b
KTH	7	7.19 ^a
LPT	8	7.29 ^a
LDP	3	6.67 ^a
GR	6	7.00 ^a
IR	11	10.74 ^b
MN	1	5.71 ^a
NP	10	7.76 ^a
YY	4	6.76 ^a
FT	2	6.62 ^a
LNP	9	7.43 ^a
Mean		7.65

Means followed by a common letter are not significantly different at the 5% level by DMRT.

5.2 Panicles/plant under 8 dS/m

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	5.00	5.33	5.33	4.33	5.33	6.00	6.00
KDML	5.33	5.33	5.33	4.00	2.33	5.00	7.33
KTH	6.00	4.67	3.67	3.67	5.67	5.33	4.67
LPT	6.33	5.00	3.67	6.33	6.33	5.33	6.67
LDP	4.33	5.67	3.67	5.67	4.33	5.67	4.67
GR	6.00	4.00	6.33	4.33	5.00	5.00	4.00
IR	6.00	8.67	7.00	5.00	6.67	6.67	4.00
MN	4.33	5.67	5.00	5.67	5.00	4.00	4.84
NP	4.33	6.67	4.67	5.00	7.00	5.00	5.67
YY	4.50	5.33	8.00	4.33	6.33	5.00	5.67
FT	3.67	5.33	5.00	4.00	5.00	4.00	7.33
LNP	6.33	6.33	5.33	8.00	7.33	4.33	8.67
Rep. totals	62.15	68.00	63.00	60.33	66.32	61.33	69.525
Rep. means	5.18	5.67	5.25	5.03	5.53	5.11	5.79

Analysis of variance for panicles/plant –Table 3.10

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	27.1780	2.4707	1.94*
Error	72	91.4657	1.2704	
Total	83	118.6437		

cv =21.0% ,* = Significant at 5 % level

Table of variance means for panicles / plant (average over 7 replications)

Var.	Ranks	Means
POK	7	5.33 ^{abc}
KDML	5	4.95 ^{ab}
KTH	1	4.81 ^a
LPT	10	5.67 ^{abc}
LDP	2	4.8 ^a
GR	6	4.95 ^{ab}
IR	11	6.29 ^{bc}
MN	4	4.93 ^{ab}
NP	8	5.48 ^{abc}
YY	9	5.59 ^{abc}
FT	3	4.90 ^{bc}
LNP	12	6.62 ^c
Mean		5.36

Means followed by a common letter are not significantly different at the 5% level by DMRT.

6. Filled grains/panicle data of Table 3.10

6.1 Filled grains/panicle under 0 dS/m (control)

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	65.00	69.84	67.86	38.90	70.37	49.35	53.67
KDML	25.28	27.51	29.73	23.40	17.07	25.13	21.10
KTH	35.00	55.93	75.58	38.15	39.90	56.77	41.81
LPT	56.15	74.11	35.30	49.81	59.89	34.78	44.81
LDP	51.14	39.00	47.85	48.45	26.93	58.39	45.24
GR	55.03	46.08	65.53	84.94	57.27	25.82	48.77
IR	25.37	30.91	36.44	32.11	31.93	22.11	28.14
MN	59.48	56.89	29.21	46.92	55.20	65.14	27.62
NP	62.24	46.83	39.77	38.63	48.90	25.07	36.29
YY	36.65	56.18	54.37	99.93	80.93	73.25	78.35
FT	68.98	64.88	60.79	44.62	52.02	59.41	53.36
LNP	97.99	72.86	47.73	87.42	53.80	37.86	98.86
Rep. totals	638.31	641.02	590.16	633.28	594.21	533.08	577.82
Rep. means	53.19	53.42	49.18	52.77	49.52	44.42	48.15

Analysis of variance for grains

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	14807.45	1346.13	6.44**
Error	72	15054.40	209.09	
Total	83	29861.85		

cv =28.9% ,**= significant at 1% level

Table of variance means for grains (average over 7 replications)

Var.	Ranks	Means
POK	10	59.28cd
KDML	1	24.17a
KTH	6	48.99c
LPT	7	50.69c
LDP	4	45.29bc
GR	8	54.78cd
IR	2	29.57ab
MN	5	48.64c
NP	3	42.53bc
YY	11	68.52d
FT	9	57.72cd
LNP	12	70.93d
Mean		50.09

Means followed by a common letter are not significantly different at the 5% level by DMRT.

6.2 Filled grains/panicle under 8 dS/m

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	67.60	53.94	62.38	50.94	32.04	66.42	35.56
KDML	15.22	14.44	15.75	17.13	18.51	31.68	5.34
KTH	46.59	51.13	28.78	28.95	29.12	63.74	33.91
LPT	22.10	53.82	8.13	34.46	51.64	9.17	22.14
LDP	77.42	21.67	36.67	40.84	16.95	42.47	32.07
GR	8.43	16.34	30.96	7.42	48.72	28.07	47.83
IR	16.62	12.27	7.92	5.93	10.69	7.84	10.92
MN	6.28	6.26	5.26	2.31	10.86	8.99	6.59
NP	7.53	24.17	16.24	36.47	40.82	37.71	24.91
YY	70.98	56.78	41.11	24.96	23.82	35.93	77.00
FT	52.49	55.39	47.99	59.37	31.79	76.13	38.69
LNP	33.77	55.39	47.99	59.37	31.79	76.13	38.69
Rep. totals	425.03	433.88	354.60	357.72	375.14	479.57	386.32
Rep. means	35.42	36.16	29.55	29.81	31.26	39.96	32.19

Analysis of variance for grains

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	21538.56	1958.05	9.47**
Error	72	14883.05	206.71	
Total	83	36421.61		

cv = 42.9%, ** = significant at 1% level

Table of variance means for grains (average over 7 replications)

Var.	Ranks	Means
POK	11	52.70 ^e
KDML	3	16.87 ^{abc}
KTH	8	40.32 ^{de}
LPT	6	28.78 ^{cde}
LDP	7	38.30 ^{de}
GR	4	26.82 ^{bcd}
IR	2	10.31 ^{ab}
MN	1	6.65 ^a
NP	5	26.84 ^{bcd}
YY	9	47.23 ^e
FT	10	51.69 ^e
LNP	12	55.25 ^e
Mean		33.48

Means followed by a common letter are not significantly different at the 5% level by DMRT.

7. 100 Grain weight data of Table 3.11

7.1 100 Grain weight under 0 dS/m (control)

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	2.51	2.75	2.70	2.75	2.75	2.70	2.50
KDML	2.05	1.95	2.20	2.05	2.00	2.10	2.05
KTH	2.26	2.15	2.25	2.05	2.25	2.15	2.15
LPT	2.26	2.05	2.05	2.15	2.25	2.20	2.20
LDP	1.74	1.70	1.60	1.55	1.50	1.65	1.55
GR	2.29	2.65	2.30	2.50	2.45	2.20	2.20
IR	1.76	1.50	1.60	1.55	1.50	1.75	1.80
MN	3.10	2.80	3.25	2.95	3.10	3.00	3.05
NP	1.85	1.80	1.75	1.75	1.80	1.60	1.70
YY	1.71	1.80	1.60	1.60	1.75	1.60	1.55
FT	1.72	1.70	1.75	1.60	1.60	1.15	1.50
LNP	1.25	1.30	1.20	1.15	1.20	1.30	1.20
Rep. totals	24.50	24.15	24.25	23.65	24.15	23.40	23.45
Rep. means	2.04	2.01	2.02	1.97	2.01	1.95	1.95

Analysis of variance for 100 grain weight

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	20.66760357	1.87887305	135.67**
Error	72	0.9708571	0.01384841	
Total	83	21.66468929		

cv = 5.9% , ** = significant at 1% level

Table of variance means for 100 grain weight (gm) (average over 7 replications)

Var.	Ranks	Means
POK	2	2.67 ^b
KDML	6	2.06 ^d
KTH	4	2.18 ^d
LPT	5 ¹	2.17 ^d
LDP	10	1.61 ^f
GR	3	2.37 ^c
IR	9	1.64 ^{ef}
MN	1	3.04 ^a
NP	7	1.75 ^e
YY	8	1.70 ^{ef}
FT	11	1.57 ^f
LNP	12	1.30 ^g
Mean		2.00

Means followed by a common letter are not significantly different at the 5% level by DMRT.

7.2 100 Grain weight under 8 dS/m

Var.	Rep.1	Rep.2	Rep.3	Rep.4	Rep.5	Rep.6	Rep.7
POK	2.65	2.80	2.65	2.35	2.40	2.65	2.45
KDML	1.90	1.75	1.90	1.80	2.15	2.00	1.95
KTH	2.05	2.00	2.20	2.10	2.00	2.00	2.03
LPT	2.20	2.05	2.40	2.15	2.20	2.20	2.20
LDP	1.55	1.65	1.50	1.70	1.55	1.60	1.66
GR	2.10	2.25	2.20	2.30	2.30	2.15	2.24
IR	1.80	1.75	1.60	1.60	1.65	1.65	1.68
MN	2.60	2.55	2.60	2.65	2.65	2.65	2.63
NP	1.60	1.65	1.65	1.80	1.70	1.60	1.70
YY	1.40	1.50	1.50	1.45	1.60	1.65	1.62
FT	1.45	1.60	1.60	1.50	1.50	1.55	1.50
LNP	1.30	1.25	1.30	1.15	1.15	1.15	1.18
Rep. totals	22.60	22.80	23.10	22.55	22.85	22.85	22.84
Rep. means	1.88	1.90	1.93	1.88	1.90	1.90	1.90

Analysis of variance for 100 grain weight (gm)

Source of variation	Degrees of freedom	Sum of squares	Mean square	F
Var.	11	14.79738452	1.34521677	160.54**
Error	72	0.60331429	0.0837937	
Total	83	15.40069881		

cv = 4.8% , ** = significant at 1% level

Table of variance means for 100 grain weight (gm) (average over 7 replications)

Var.	Ranks	Means
POK	2	2.56 ^a
KDML	6	1.92 ^d
KTH	5	2.05 ^c
LPT	4	2.20 ^b
LDP	9	1.60 ^{e,f}
GR	3	2.22 ^b
IR	7	1.68 ^e
MN	1	2.62 ^a
NP	8	1.67 ^e
YY	10	1.53 ^f
FT	11	1.53 ^f
LNP	12	1.21 ^g
Mean		1.90

Means followed by a common letter are not significantly different at the 5% level by DMRT.

Biography

Mr. Chukiat Khotanakul was born on January 24, 1956 in Chonburi. He graduated with the degree of Bachelor of Science (B.Sc. in Chemistry) from Ramkhamhaeng University in 1978 and Master of Science (M.Sc. in Biochemistry) from Chulalongkorn University in 1983. He has been working in the Department of Pharmaceutical Chemistry, Faculty of Pharmaceutical Sciences, Prince of Songkhla University since 1983.

