

STUDIES ON SEMINAL ATTRIBUTES IN RELATION TO  
CHROMOSOMAL PROFILE IN MURRAH BUFFALO BULLSVinod Kumar Shakya<sup>1</sup>, Shrikant Joshi<sup>2</sup>, Madhu Shivhare<sup>3,\*</sup>, S.P. Nema<sup>1</sup> and Vishnu Gupta<sup>1</sup>

## ABSTRACT

Seminal attributes and karyological study was conducted on twenty five Murrah buffalo breeding bulls belonging to Central Semen Station, Bhopal. Five semen samples from each buffalo bull were collected at 7 days interval. Fresh semen was evaluated for volume, mass motility, sperm concentration, progressive motility, live sperm count and morphological abnormalities. The mean volume, mass motility, progressive motility, sperm concentration, live sperm count and total abnormal sperms were  $2.71 \pm 0.112\%$ ,  $3.34 \pm 0.073\%$ ,  $68.4 \pm 1.30\%$ ,  $978.9 \pm 34.37\%$ ,  $89.52 \pm 0.38\%$ , and  $8.82 \pm 0.873\%$ . Significant ( $P < 0.05$ ) variation was observed between the bulls with regard to sperm abnormalities. However, non-significant difference was observed for volume, sperm concentration, mass motility, progressive motility, and live sperm count. The overall mean sperm abnormalities found for abnormal head, middle piece and tail were  $1.68 \pm 0.160\%$ ,  $2.0 \pm 0.238\%$ , and  $5.12 \pm 0.475\%$ , respectively. Tail

abnormalities differed significantly between bulls. Karyological screening of conventionally stained slides and G and C banded slides of breeding bulls did not reveal any deviation from normal diploid chromosome number and morphology of water buffaloes. The presence of structural abnormalities like gaps and breaks were observed at very low frequency. In general the centromeric region was G-band negative. The centromeric heterochromatin of submetacentric autosomes stained faintly than that of acrocentric autosomes, the X-Chromosome had a prominent C-band.

**Keyword:** *Bubalus bubalis*, buffaloes, karyological, progressive motility, centromeric region

## INTRODUCTION

Selection of young bulls at an early age is crucial for commercial semen producers (Hafez and Hafez, 2000). A deficiency in bull reproductive traits has a larger impact on herd productivity

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and fertility problems than in a single female (Rodriguez-Martinez, 2008). Evaluation of semen quality has been based on routine semen analysis (motility, morphology and acrosome integrity, Selvaraju *et al.*, 2008). Chromosomal abnormalities account for a substantial loss in animal production and problem related to fertility which can be screened even at calf hood stage and the abnormal animal may be removed from breeding programme to avoid transmission of such abnormalities to their progenies. About 8% of reproductive deficient animals have shown chromosomal abnormalities and hence timely culling of such bulls will improve fertility of the herd (Pattanayak *et al.*, 2007).

## MATERIALS AND METHODS

The present study was conducted on Twenty five Murrah buffalo breeding bulls belonging to Central Semen Station, Bhopal. Semen ejaculates were harvested from bulls using Swedish pattern, artificial vagina. From each bull five samples were collected, at an interval of 7 days. Semen evaluation was done immediately after collection. Volume of the semen was noted directly from the graduations of semen collection glass tube. Mass motility was observed as per the procedure described by Herman

and Madden (1953). The individual motility of spermatozoa was expressed in terms of percentage of progressively motile spermatozoa (Zemjanis, 1970). Spermatozoa concentration (million/ml) was determined using photo colorimeter (IMV), standardized at 530 n (Willet and Buckner, 1951). Vital count of spermatozoa (percentage) was estimated using Eosin-Nigrosin staining technique (Campbell *et al.*, 1956). Sperm morphological abnormalities in head, middle piece and tail were studied as per the procedure described by Rao (1971). For karyological studies short term lymphocyte culture technique using RPMI 1640 culture medium was adopted (Moorhead *et al.*, 1960). Pokeweed was used as mitogen. G and C banding were done by standard procedure. A total of 20 to 30 metaphase plates were screened for each bull. General morphology and number of chromosomes along with abnormalities in morphology and numbers if any was recorded.

## RESULT AND DISCUSSION

### Seminal attributes volume

The overall mean semen volume recorded was  $2.71 \pm 0.112$  ml (Table 1) ranging from  $2 \pm 0.332$  to  $4.4 \pm 0.87$  ml. Analysis of variance revealed non-

Table 1. Seminal attributes of buffalo breeding bulls (mean $\pm$ SE).

Seminal attributes	Mean $\pm$ SE	Morphological abnormalities	Mean $\pm$ SE (%)
Volume(ml)	2.71 $\pm$ 0.11	Head abnormalities	1.68 $\pm$ 0.16
Sperm concentration (mill/ml)	978.9 $\pm$ 34.37	Mid piece abnormalities	2.0 $\pm$ 0.23
Mass activity(score)	3.34 $\pm$ 0.07	Tail abnormalities	5.12 $\pm$ 0.47
Progressive motility (%)	68.4 $\pm$ 1.30	Total abnormalities	8.82 $\pm$ 0.87
Live sperm count (%)	89.52 $\pm$ 0.38		
Sperm Morphology (%)	11.12 $\pm$ 0.41		

significant difference ( $P < 0.05$ ) in the volume of semen between bulls. This is in agreement with the findings of Tiwari *et al.* (2009) who reported almost similar volume of semen in Tarai and Surti buffalo bulls. Patel *et al.* (2012) reported higher values in Jafarabadi, Mehsana and crossbred (HF x Kankrej, F1). These variations may be attributed to breed variations.

### Mass motility

Based on 0 to 5 scale, the mean mass motility ( $3.34 \pm 0.073$ ) of spermatozoa in the semen of breeding bull ranged from 2.6 to 4.0 ( $P > 0.05$ ). These observations are in close resemblance to the findings of previous workers (Shukla and Misra, 2005).

### Progressive motility

The overall mean progressive motility of spermatozoa in present study was found to be  $68.4 \pm 1.30\%$ , ranging between 55 to 77%. Between bull variation in the progressive motility of spermatozoa was found to be statistically non significant. The present findings are in close agreement with the observations of Pandey (2001); Narayan *et al.* (1999), though, Selvaraju *et al.* (2008) reported lower progressive motility in Murrah bulls.

### Sperm concentration

The mean value of sperm concentration ( $978.9 \pm 34.37$  million/ml) in semen of buffalo breeding bulls, ranged from 632.8 to 1335.8 million/ml. The present results corroborate with the findings of Tiwari *et al.* (2009) in Tarai buffalo bulls. Selvaraju *et al.* (2008) reported a very higher concentration of sperm in Murrah buffalo bulls.

### Live sperm count

The overall mean live spermatozoa count was  $89.52 \pm 0.38\%$ , with a range of 86.4 to 92.6%. Statistically it did not reveal significant difference between the bulls. The present study is in agreement with the finding of Shukla (2002), who also recorded similar mean percent live spermatozoa in the neat semen of the Murrah bulls.

### Sperm abnormalities

The mean percentage of abnormal spermatozoa in the semen of different bulls was  $8.82 \pm 0.873\%$ , which is fairly well comparable with the findings of Pandey (2001); Patel (2011). Some other workers reported higher percent of total sperm abnormalities (Nath *et al.*, 1991; Shabd, 1998). A few workers reported lower percent total abnormal sperm abnormalities (Singh *et al.*, 2000; Perumal *et al.*, 2009) however, Shukla and Misra (2005) observed statistically non-significant difference between bulls. The average head and mid-piece abnormalities of spermatozoa in the present study were  $1.68 \pm 0.160$  and  $1.71 \pm 0.285$  respectively. Fairly well comparable finding were recorded for sperm head (Gunarajasingham *et al.*, 1996, Veerabramhaiah *et al.*, 2010; Patel, 2011) and for mid-piece abnormalities (Shukla, 2002; Patel, 2011). There was no significant variation between bull and between replicate variations in the sperm head abnormalities in the present study. Similar findings were recorded by Shukla and Misra (2005); Siddiquee *et al.* (2011); Patel (2011). Significant between bull variation ( $P < 0.05$ ) was recorded in the mid-piece abnormalities of the Murrah bulls in the present study. This was in compliance to the findings of Shukla and Misra (2005). As regards tail abnormalities ( $2.09 \pm 0.21\%$ ) in the present study are concerned they were considerably low as compared to several other workers (Gunarajisingam

*et al.*, 1996; Patel, 2011).

### Cytogenetic studies

In the present study the observed diploid chromosomes number was  $2n=50$ . The karyotype comprised of 5 pairs of submetacentric, 19 pairs of acrocentric autosomes and 1 pairs of acrocentric sex chromosomes. The observed G-band and C-band patterns were in closed agreement with the features reported by earlier workers in river type buffalo (Bongso and Hilmi, 1982; Thiagrajan, 1987; Yadav, 1981; Yadav and Balkrishan, 1982). In the present study structural abnormalities were recorded in the form of gaps and breaks at very low frequency. This might be associated with mechanical losses during harvesting and preparation of slides. Many of the chromosomal abnormalities in cattle breeding bulls have been reported by various workers (Kieffer and Cartwright, 1968; Mandal *et al.*, 2003; Ahmad *et al.*, 2004). However, availability of such reports in buffalo breeding bulls is very scanty. In the present study no abnormality could be traced out in any of the breeding bulls. This could be due to the fact that all the bulls maintained at central semen station have been procured from selected stock and also the buffaloes have less prominence and aesthetic values compared to sacred cattle consistently they are slaughtered more frequently with any physical or reproductive abnormalities. In the present study karyological screening of few animals revealed the presence of structural abnormalities like gaps and breaks. However, the frequency of such abnormalities was very low. None of the bull showed presence of any numerical chromosomal abnormalities.

### CONCLUSION

Variation in the mean volume, mass activity, individual motility, sperm concentration and live sperm percent between bulls were non-significant. However, statistical variation for morphological abnormalities between breeding bulls was highly significant.

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