

SUPPLEMENTARY TABLE

Supplementary Table 1. The AT-content (%) of mono- to hexanucleotide P-SSRs in the different genomic regions of the forest musk deer genome.

Type	5'UTRs	CDSs	Introns	3'UTRs	TEs	Intergenics
Mono-	98.57	61.05	97.80	96.29	99.29	98.16
Di-	44.31	48.17	60.84	64.67	69.23	65.51
Tri-	7.70	26.55	37.55	55.48	53.58	39.33
Tetra-	40.16	27.69	68.31	60.85	81.61	74.58
Penta-	47.32	27.61	59.52	47.73	60.87	60.04
Hexa-	16.67	30.53	41.35	33.33	49.86	36.49
Total	28.35	27.23	67.92	81.51	83.95	69.02