

Individuals			Illumina				Oxford Nanopore Sequencing					Assembly statistics					
Species	Subspecies	Voucher	Raw reads (M)	Cleaned reads	Nbr of gigabases	Estimated coverage	Nbr of flowcells	Nbr of bases (Gb)	N50	Average size	Estimated coverage	Genome size (Gb)	Nbr of scaff.	N50 (Kb)	Busco score	OMM genes	Missing data (%)
<i>Proteles cristata</i>	cristata	TS307	716.7	522.8	129.50	51.8	16	27.3	9,175	5,555	10.9	2.39	5,669	1,309	92.8	12,062	22.43
<i>Proteles cristata</i>	cristata	TS491	663.8	526.1	140.73	56.3	NA					NA				NA	
<i>Proteles cristata</i>	septentrionalis	NMSZ201854	750.9	516.2	132.44	53.0	NA					NA				12,050	22.96
<i>Otocyon megalotis</i>	megalotis	TS305	710.2	584.4	154.81	61.6	13	33	4,393	3,092	13.2	2.75	11,081	728	92.9	11,981	22.02
<i>Otocyon megalotis</i>	megalotis	TS306	861.2	820	240.71	96.3	NA					NA					
<i>Otocyon megalotis</i>	virgatus	FMNH158128	661.7	554.1	100.30	40.1	NA					NA					