

No	Species	Assembly Method	Sequencing Technology	Genome Coverage
1	<i>Homo sapiens</i>	Celera	Sanger	20x
2	<i>Pogona Vitticeps</i>	SOAP deNovo	Illumina HiSeq 2000	85.5x
3	<i>Anolis Carolinensis</i>	Arachne v.3.0.0	ABI	7.10x
4	<i>Gallus gallus</i>	Celera Assemblerv.5.4	Sanger; 454	12x
5	<i>Monodelphis domestica</i>	ARACHNE2+	Sanger	6.8x
6	<i>Ornithorhynchus anatinus</i>	PCAP	WGS plasmid, fosmid end and BAC end sequences	6x