

Sequence ID	Start	Alignment	End	Organism
		1 100 200 300 400 500 600 700 800 900 1K 1,100 1,200 1,300 1,400 1,500 1,600 1,700 1,800 1,900 2K 2,100 2,200 2,300 2,400 2,500 2,600 2,700 2,800 2,900 3K 3,100 3,200 3,367		
CTG21514.1 (+)	1		345	Streptococcus pneumoniae
XP_007982279.1 (+)	1		1,135	Chlorocebus sabaeus
XP_006035731.1 (+)	1		1,132	Alligator sinensis
XP_007066464.1 (+)	1		2,831	Chelonia mydas
XP_012401853.1 (+)	1		1,881	Sarcophilus harrisii
XP_006082532.1 (+)	1		1,909	Myotis lucifugus
XP_004427558.1 (+)	1		1,838	Ceratotherium simum simum
XP_004262822.1 (+)	1		1,837	Orcinus orca
XP_001109205.3 (+)	1		1,858	Macaca mulatta
XP_008046808.1 (+)	1		1,822	Carlito syrichta
XP_007909862.1 (+)	1		437	Callorhinus milii
XP_006638354.1 (+)	1		755	Lepisosteus oculatus
XP_005994684.1 (+)	1		533	Latimeria chalumnae
XP_012808014.2 (+)	1		709	Xenopus tropicalis
ETE73308.1 (+)	1		692	Ophiophagus hannah
XP_007482569.1 (+)	1		816	Monodelphis domestica
XP_003222596.2 (+)	1		700	Anolis carolinensis
XP_008160844.1 (+)	1		681	Chrysemys picta bellii
NP_001186122.1 (+)	1		432	Gallus gallus
XP_013153409.1 (+)	1		546	Falco peregrinus
XP_006159184.1 (+)	1		793	Tupaia chinensis
XP_006532473.1 (+)	1		767	Mus musculus
XP_003414352.1 (+)	1		788	Loxodonta africana
XP_013003800.1 (+)	1		448	Cavia porcellus
XP_005858648.1 (+)	1		803	Myotis brandtii
XP_008269868.1 (+)	1		840	Oryctolagus cuniculus
XP_003786275.1 (+)	1		774	Otolemur garnettii
XP_006199331.1 (+)	1		784	Vicugna pacos
XP_012352933.1 (+)	1		776	Nomascus leucogenys
Q5YCV9.4 (+)	1		776	Hylobates lar
NP_001116538.2 (+)	1		776	Homo sapiens
XP_002800600.1 (+)	1		776	Macaca mulatta
XP_003913279.2 (+)	1		776	Papio anubis
XP_008995083.1 (+)	1		497	Callithrix jacchus
XP_010328565.2 (+)	1		852	Saimiri boliviensis boliviensis
XP_007957952.1 (+)	1		780	Orycteropus afer afer
XP_005983781.1 (+)	1		778	Pantholops hodgsonii
EFB25687.1 (+)	1		780	Ailuropoda melanoleuca
XP_013845380.1 (+)	1		451	Sus scrofa