

Annotation project	Species	Assembly	Annotation
allMis4	<i>Alligator mississippiensis</i>	GCA_000281125.4	GCF_000281125.3_ASM28112v4_genomic.gff.gz
allSin1	<i>Alligator sinensis</i>	GCA_000455745.1	GCF_000455745.1_ASM45574v1_genomic.gff.gz
anaPla1.5	<i>Anas platyrhynchos</i>	GCA_003850225.1	
andRos1	<i>Anguilla rostrata</i>	GCA_001606085.1	GCF_000146605.3_Turkey_5.1_genomic.gff.gz
anoCar2.0	<i>Anolis carolinensis</i>	GCA_000090745.2	GCF_000090745.1_Anocar2.0_genomic.gff.gz
aptFor1	<i>Aptenodytes forsteri</i>	GCA_000699145.1	
astMex2.0	<i>Astyanax mexicanus</i>	GCA_000372685.2	GCF_000372685.2_Astyanax_mexicanus-2.0_genomic.gff.gz
balBon1	<i>Balaenoptera bonaerensis</i>	GCA_000978805.1	GCA_000978805.1_ASM97880v1_genomic.gff.gz
balReg1	<i>Balearica regulorum</i>	GCA_000709895.1	
bosTau1.2	<i>Bos Taurus</i>	GCA_002263795.2	
braFlo2	<i>Branchiostoma floridae</i>	GCA_000003815.1	
bucRhi1	<i>Buceros rhinoceros</i>	GCA_000710305.1	
calPeg1	<i>Calidris pugnax</i>	GCA_001431845.1	
calMil6.1.3	<i>Callorhinchus milii</i>	GCA_000165045.2	GCF_000165045.1_Callorhinchus_milii-6.1.3_genomic.gff.gz
calAnn1	<i>Calypte anna</i>	GCA_003957555.2	
catAur1	<i>Cathartes aura</i>	GCA_000699945.1	
chiPla1	<i>Chiloscyllium plagiosum</i>	GCA_004010195.1	
chrPic3.0.3	<i>Chrysemys picta</i>	GCA_000241765.2	GCF_000241765.3_Chrysemys_picta_bellii-3.0.3_genomic.gff.gz
ciolntKH	<i>Ciona intestinalis</i>	GCA_000224145.2	
cluHar1	<i>Clupea harengus</i>	GCA_900323705.1	GCF_900700415.1_Ch_v2.0.2_genomic.gff.gz
colLiv1.0	<i>Columba livia</i>	GCA_000337935.1	
ConCri1.0	<i>Condylura cristata</i>	GCA_000260355.1	GCF_000260355.1_ConCri1.0_genomic.gff.gz
corAlt1	<i>Corapipo altera</i>	GCA_003945725.1	
corBra1	<i>Corvus brachyrhynchos</i>	GCA_000691975.1	
croPor1	<i>Crocodylus porosus</i>	GCA_001723895.1	GCF_001723895.1_CroPor_comp1_genomic.gff.gz
cucCan1	<i>Cuculus canorus</i>	GCA_000709325.1	
cynSem1.0	<i>Cynoglossus semilaevis</i>	GCA_000523025.1	GCF_000523025.1_Cse_v1.0_genomic.gff.gz
cypCar	<i>Cyprinus carpio</i>	GCA_000951615.2	GCF_000951615.1_common_carp_genome_genomic.gff.gz
danRer11	<i>Danio rerio</i>	GCA_000002035.4	GCF_000002035.6_GRCz11_genomic.gff.gz
dasNov3	<i>Dasypus novemcinctus</i>	GCA_000208655.2	GCF_000208655.1_Dasnov3.0_genomic.gff.gz
delLeu2	<i>Delphinapterus leucas</i>	GCA_002288925.3	
egrGar1	<i>Egretta garzetta</i>	GCA_000687185.1	

Efet.01	<i>Eisenia fetida</i>	GCA_003999395.1	
EleEdw1.0	<i>Elephantulus edwardii</i>	GCA_000299155.1	
equCab3.0	<i>Equus caballus</i>	GCA_002863925.1	GCF_002863925.1_EquCab3.0_genomic.gff.gz
eriEur2	<i>Erinaceus europaeus</i>	GCA_000296755.1	GCF_000296755.1_EriEur2.0_genomic.gff.gz
esoLuc3	<i>Esox lucius</i>	GCA_000721915.3	GCF_004634155.1_Eluc_v4_genomic.gff.gz
eurHel1	<i>Eurypyga helias</i>	GCA_000690775.1	
galGal6a	<i>Gallus gallus</i>	GCA_000002315.5	
gavGan1	<i>Gavialis gangeticus</i>	GCA_001723915.1	GCF_001723915.1_GavGan_comp1_genomic.gff.gz
gekJap1.1	<i>Gekko japonicus</i>	GCA_001447785.1	GCF_001447785.1_Gekko_japonicus_V1.1_genomic.gff.gz
halLeu4.0	<i>Haliaeetus leucocephalus</i>	GCA_000737465.1	
hetGla2	<i>Heterocephalus glaber</i>	GCA_000247695.1	
hipCam1	<i>Hippocampus comes</i>	GCA_001891065.1	GCF_001891065.1_H_comes_QL1_v1_genomic.gff.gz
lagObl1	<i>Lagenorhynchus obliquidens</i>	GCA_003676395.1	
lipVex1	<i>Lipotes vexillifer</i>	GCA_000442215.1	
loxAfr3	<i>Loxodonta Africana</i>	GCA_000001905.1	GCF_000001905.1_Loxafr3.0_genomic.gff.gz
melGall5.0	<i>Meleagris gallopavo</i>	GCA_000146605.4	GCF_000146605.3_Turkey_5.1_genomic.gff.gz
melUnd6.3	<i>Melopsittacus undulatus</i>	GCA_000238935.1	Anguilla_rostrata
merNub1	<i>Merops nubicus</i>	GCA_000691845.1	
nsch1	<i>Neomonachus schauinslandi</i>	GCA_002201575.1	GCF_002201575.1_ASM220157v1_genomic.gff.gz
notScu2	<i>Notechis scutatus</i>	GCA_900518725.1	GCF_900518725.1_TS10Xv2-PRI_genomic.gff.gz
numMel1.0	<i>Numida meleagris</i>	GCA_002078875.2	
opiTho1	<i>Opisthocomus hoazin</i>	GCA_000692075.1	
OfaDov1	<i>Orbicella faveolata</i>	GCA_002042975.1	
orcOrc1.1	<i>Orcinus orca</i>	GCA_000331955.2	
ornAna1.p.v1	<i>Ornithorhynchus anatinus</i>	GCA_004115215.2	GCF_004115215.1_mOrnAna1.p.v1_genomic.gff.gz
OryAfe1.0	<i>Orycteropus afer afer</i>	GCA_000298275.1	
oryLat1	<i>Oryzias latipes</i>	GCA_002234675.1	GCF_002234675.1_ASM223467v1_genomic.gff.gz
parMaj1.1	<i>Parus major</i>	GCA_001522545.3	
pelCri1	<i>Pelecanus crispus</i>	GCA_000687375.1	
pelSin1.0	<i>Pelodiscus sinensis</i>	GCA_000230535.1	GCF_000230535.1_PelSin_1.0_genomic.gff.gz
petMar1.0	<i>Petromyzon marinus</i>	GCA_002833325.1	
phaCar1	<i>Phalacrocorax carbo</i>	GCA_000708925.1	
phaCin4.1	<i>Phascolarctos cinereus</i>	GCA_002099425.1	GCF_002099425.1_phaCin_unsw_v4.1_genomic.gff.gz

phoRub1	<i>Phoenicopterus ruber</i>	GCA_000687265.1	
phyCat1	<i>Physeter catodon</i>	GCA_002837175.2	GCF_002837175.2_ASM283717v2_genomic.gff.gz
podCri1	<i>Podiceps cristatus</i>	GCA_000699545.1	
pomCan1	<i>Pomacea canaliculate</i>	GCA_003073045.1	
proCap1	<i>Procapia capensis</i>	GCA_004026925.2	
pseNaj2	<i>Pseudonaja textilis</i>	GCA_900518735.1	GCF_900518735.1_EBS10Xv2-PRI_genomic.gff.gz
pytBiv5.0.2	<i>Python bivittatus</i>	GCA_000186305.2	GCF_000186305.1_Python_molurus_bivittatus-5.0.2_genomic.gff.gz
ramVar4.0	<i>Ramazzottius varieornatus</i>	GCA_001949185.1	
rhiTyp2	<i>Rhincodon typus</i>	GCA_001642345.2	GCF_001642345.1_ASM164234v2_genomic.gff.gz
salSal2	<i>Salmo salar</i>	GCA_000233375.4	GCF_000233375.1_ICSASG_v2_genomic.gff.gz
strCam1	<i>Struthio camelus</i>	GCA_000698965.1	
susScr11.1	<i>Sus scrofa</i>	GCA_000003025.6	
tauEry1	<i>Tauraco erythrolophus</i>	GCA_000709365.1	
thaSir1	<i>Thamnophis sirtalis</i>	GCA_001077635.2	GCF_001077635.1_Thamnophis_sirtalis-6.0_genomic.gff.gz
tinGut2	<i>Tinamus guttatus</i>	GCA_000705375.2	
turTru1	<i>Tursiops truncatus</i>	GCA_001922835.1	
tytAlb1	<i>Tyto alba</i>	GCA_000687205.1	
aRhiBiv1.1	<i>Rhinatrema bivittatum</i>	GCA_901001135.1	GCF_901001135.1_aRhiBiv1.1_genomic.gff.gz
nanPar1	<i>Nanorana parkeri</i>	GCA_000935625.1	GCF_000935625.1_ASM93562v1_genomic.gff.gz
RCv2.1	<i>Lithobates catesbeianus</i>	GCA_002284835.2	GCA_002284835.2_RCv2.1_genomic.gff.gz
xenTro10	<i>Xenopus tropicalis</i>	GCA_000004195.4	GCF_000004195.4_UCB_Xtro_10.0_genomic.gff.gz
TaeGut1	<i>Zebra finch</i>	GCA_003957565.2	GCF_003957565.1_bTaeGut1_v1.p_genomic.gff.gz