



Homo\_sapiens\_GH-N G 199  
Homo\_sapiens\_nean . 200  
Pan\_troglodytes\_G . 200  
Gorilla\_gorilla\_g . 200  
Pongo\_abelii\_GH-N . 200  
Homo\_sapiens\_GH-V . 200  
Homo\_sapiens\_nean . 200  
Pan\_troglodytes\_G . 200  
Gorilla\_gorilla\_g . 200  
Pongo\_abelii\_GH-V . 201  
Homo\_sapiens\_CSH- . 200  
Homo\_sapiens\_CSH- . 201  
Homo\_sapiens\_CSH- . 201  
Homo\_sapiens\_nean . 200  
Homo\_sapiens\_nean . 201  
Homo\_sapiens\_nean . 201  
Pan\_troglodytes\_C . 201  
Pan\_troglodytes\_C . 200  
Pan\_troglodytes\_C . 201  
Gorilla\_gorilla\_g . 200  
Gorilla\_gorilla\_g . 201  
Gorilla\_gorilla\_g . 201  
Gorilla\_gorilla\_g . 201  
Pongo\_abelii\_CSH- . 201  
Pongo\_abelii\_CSH- . 201

Sp-1 distal Pit-1 CRE Proximal Pit-1 TRE TATA InrE

Figure S1 alignments of promoters' nucleotide sequences.