

gp120

scening

Finished

HOME







TABLE

READ

TREE

Summary

Site	Cover	Aderine	Thymine	Ganine	Cytosine	Deletion	Con. Nuc	% Non C.	Mean Non Cor Qua...	HPS	Entropy	Codons
528	171	0	158	0	0	13	T	0.0000	0.0000	6	0.10539...	
529	171	171	0	0	0	0	A	0.0000	0.0000	6	-0.0	ATA:I (166), ATT:I (3), ATG:M (2),
530	171	0	171	0	0	0	T	0.0000	0.0000	0	-0.0	
531	171	165	3	2	0	0	A	2.9240	0.0000	0	0.21095...	
532	171	4	0	167	0	0	G	2.3302	59.3114	1	0.160082...	CCC:A (164), AGC:G (3), GTC:V (2), ACC:T (1),
533	171	1	2	3	165	0	C	3.5088	61.9000	2	0.720495...	
534	177	0	0	0	177	0	C	0.0000	0.0000	2	-0.0	
535	181	0	181	0	0	0	T	0.0000	0.0000	0	-0.0	TTG:L (180), TCG:S (.),
536	181	0	180	0	1	0	T	0.5525	0.0000	1	0.049384...	
537	181	0	0	181	0	0	G	0.0000	0.0000	1	-0.0	
538	182	182	0	0	0	0	A	0.0000	0.0000	0	-0.0	ATA:I (181), ATG:M (1),
539	182	0	183	0	0	0	T	0.0000	0.0000	0	-0.0	
540	196	195	0	1	0	0	A	0.5102	0.0000	0	0.046192...	
541	197	0	197	0	0	0	T	0.0000	0.0000	0	-0.0	TAG:* (196), TAA:* (.),
542	197	197	0	0	0	0	A	0.0000	0.0000	0	-0.0	
543	197	1	0	196	0	0	G	0.5076	59.3067	1	0.045395...	
544	202	0	203	0	0	0	T	0.0000	0.0000	1	-0.0	TAC:Y (203),
545	208	208	0	0	0	0	A	0.0000	0.0000	0	-0.0	
546	212	0	0	0	2.2	0	C	0.0000	0.0000	0	-0.0	
547	214	13	1	0	200	0	C	6.5421	61.1429	1	0.372883...	CAA:Q (195), AAA:K (13), TAA:* (1),
548	216	215	0	0	0	0	A	0.0000	0.0000	1	-0.0	
549	216	215	0	1	0	0	A	0.4630	0.0000	1	0.042565...	
550	216	0	218	0	0	0	T	0.0000	0.0000	1	-0.0	TAG:* (218),
551	216	215	0	0	0	0	A	0.0000	0.0000	0	-0.0	

REFERENCE: TAACAAATAGATAATGATACTACCAAGCTAATAGTGTGACCAAGTTGTAACACCTCAGTCAATACACAGGCTGTCCAAAAGGTATCCFTT5AG6CAATTCCCATACAT7AT1GTGCCCGGCTG5PTTFTGCGA
COVER: 
NON-CONSENSUS: 
ENTROPY: 
INSERTIONS (len.1): 
DELETIONS (len.1): 
530 540 550 560 570 580 590 600 610 620 630 640 650 660 670
0  100%

Tree Scan

Unique Tips: 28

Window  100Mag.  174% Depth  20

Create Full Pairwise Tree

Dataset Summary

No. Reads Mapped: 3953

Forward Mapped: 2110

Reverse Mapped: 1843

Mean Read Length: 194

Table Scan

Frame  1Non Con.  0.05Quality  10HPS Len.  3

SAVE TABLE

Incel len.  1