Ref_index	161	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189	190	191	192
Target_seq						G	G	С	Α	С	С	G	G	С	Т	G	G	Α	G	G	Т	G	G	Ν	G	G						
Ref_seq	G	G	G	С	G	G	G	С	Α	С	Α	G	G	С	Т	G	G	G	G	Α	Т	G	G	Α	G	G	Α	G	С	Т	G	G
Α	0	0	0	0	0	0	0	0	27	1	30	0	0	0	0	0	0	0	0	34	0	0	0	31	0	0	33	0	1	0	0	0
G	30	26	29	0	30	27	29	0	2	0	2	33	33	0	0	28	32	30	30	0	0	33	34	1	31	32	1	34	1	0	33	35
С	0	2	0	29	0	0	0	30	1	30	0	0	0	33	0	1	1	1	0	0	0	0	0	0	1	1	0	0	32	1	0	0
Т	0	0	0	0	0	0	0	0	0	0	0	0	0	0	33	1	0	0	0	0	34	1	0	0	0	0	0	0	0	32	1	0
Del	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Ins	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Α%	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	90.0	3.2258	93.75	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	100.0	0.0	0.0	0.0	96.875	0.0	0.0	97.0588	0.0	2.9412	0.0	0.0	0.0
G %	100.0	92.8571	100.0	0.0	100.0	100.0	100.0	0.0	6.6667	0.0	6.25	100.0	100.0	0.0	0.0	93.3333	96.9697	96.7742	100.0	0.0	0.0	97.0588	100.0	3.125	96.875	96.9697	2.9412	100.0	2.9412	0.0	97.0588	100.0
C %	0.0	7.1429	0.0	100.0	0.0	0.0	0.0	100.0	3.3333	96.7742	0.0	0.0	0.0	100.0	0.0	3.3333	3.0303	3.2258	0.0	0.0	0.0	0.0	0.0	0.0	3.125	3.0303	0.0	0.0	94.1176	3.0303	0.0	0.0
T %	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	100.0	3.3333	0.0	0.0	0.0	0.0	100.0	2.9412	0.0	0.0	0.0	0.0	0.0	0.0	0.0	96.9697	2.9412	0.0
Del %	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Ins %	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.125	0.0	0.0	0.0	0.0	0.0	3.2258	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0