

<b>Color Code Description</b>	White = Identical Calls (based on the corresponding reported data)
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Features & Coordinates				Illumina's Platinum Genomes Variants <a href="http://www.illumina.com/platinumgenomes">www.illumina.com/platinumgenomes</a>						Crystal Genetics' Variants <a href="http://www.crystalgenetics.com">www.crystalgenetics.com</a>					
idx	gene	chr	position	varcall	ref	alt	var	filter	varcall	ref	alt	var	filter		
1	BRCA1	17	41,196,408	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
2	BRCA1	17	41,196,821	1	CTT	C	0/1	Silver	1	CTT	C	0/1	PASS		
3	BRCA1	17	41,197,274	1	C	A	0 1	PASS	1	C	A	0/1	PASS		
4	BRCA1	17	41,198,621	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
5	BRCA1	17	41,198,774	1	C	CA	0/1	Silver	1	C	CA	0/1	PASS		
6	BRCA1	17	41,199,913	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
7	BRCA1	17	41,200,109	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
8	BRCA1	17	41,200,537	1	T	C	0 1	PASS	1	T	C,TT	1/2	PASS		
9	BRCA1	17	41,200,704	1	GT	G	0 1	PASS	1	GT	G	0/1	PASS		
10	BRCA1	17	41,201,702	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
11	BRCA1	17	41,202,635	1	G	A	0 1	PASS	0	-	-	-	-		
12	BRCA1	17	41,202,688	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
13	BRCA1	17	41,203,325	1	T	A	0 1	PASS	1	T	A	0/1	PASS		
14	BRCA1	17	41,203,591	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
15	BRCA1	17	41,204,377	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
16	BRCA1	17	41,204,383	0	-	-	-	-	1	G	T	0/1	PASS		
17	BRCA1	17	41,204,390	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
18	BRCA1	17	41,204,836	0	-	-	-	-	1	ACACACT	A	1/1	PASS		
19	BRCA1	17	41,204,842	1	T	A	1 1	Silver	1	T	A	1/1	PASS		
20	BRCA1	17	41,204,845	0	-	-	-	-	1	CTT	T	1/1	PASS		
21	BRCA1	17	41,204,856	1	G	A	1 0	PASS	1	G	A	0/1	PASS		
22	BRCA1	17	41,205,772	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
23	BRCA1	17	41,205,941	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
24	BRCA1	17	41,206,056	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
25	BRCA1	17	41,206,761	1	AT	A	0 1	PASS	0	-	-	-	-		
26	BRCA1	17	41,206,777	0	-	-	-	-	1	TG	G	0/1	PASS		
27	BRCA1	17	41,209,578	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
28	BRCA1	17	41,210,396	1	A	C	0 1	PASS	1	A	C	0/1	PASS		
29	BRCA1	17	41,211,486	1	CACA	C,CACAACA	1 2	PASS	0	-	-	-	-		
30	BRCA1	17	41,211,488	0	-	-	-	-	1	CAAC	CAACAAC,C	1/2	PASS		
31	BRCA1	17	41,211,653	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
32	BRCA1	17	41,212,169	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
33	BRCA1	17	41,212,338	1	A	C	0 1	PASS	1	A	C	0/1	PASS		
34	BRCA1	17	41,212,547	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
35	BRCA1	17	41,212,805	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
36	BRCA1	17	41,213,602	1	CT	C	0 1	PASS	0	-	-	-	-		
37	BRCA1	17	41,213,617	0	-	-	-	-	1	TA	A	0/1	PASS		
38	BRCA1	17	41,213,620	0	-	-	-	-	1	G	A	0/1	PASS		

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idx	gene	chr	position	varcall	ref	alt	var	filter	varcall	ref	alt	var	filter		
39	BRCA1	17	41,213,626	1	G	T	0 1	PASS	1	G	T	0/1	PASS		
40	BRCA1	17	41,213,660	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
41	BRCA1	17	41,213,748	1	T	C	1 0	PASS	1	T	C	0/1	PASS		
42	BRCA1	17	41,213,893	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
43	BRCA1	17	41,213,996	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
44	BRCA1	17	41,214,209	1	CAA	C	0 1	PASS	0	-	-	-	-		
45	BRCA1	17	41,214,210	0	-	-	-	-	1	AAG	G	0/1	PASS		
46	BRCA1	17	41,215,825	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
47	BRCA1	17	41,216,205	1	G	T	0 1	PASS	1	G	T	0/1	PASS		
48	BRCA1	17	41,216,865	0	-	-	-	-	1	TG	G	0/1	PASS		
49	BRCA1	17	41,216,933	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
50	BRCA1	17	41,217,874	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
51	BRCA1	17	41,218,333	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
52	BRCA1	17	41,218,572	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
53	BRCA1	17	41,219,341	1	G	T	0 1	PASS	1	G	T	0/1	PASS		
54	BRCA1	17	41,219,560	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
55	BRCA1	17	41,219,780	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
56	BRCA1	17	41,219,804	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
57	BRCA1	17	41,220,223	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
58	BRCA1	17	41,222,462	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
59	BRCA1	17	41,222,723	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
60	BRCA1	17	41,223,094	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
61	BRCA1	17	41,223,538	1	G	GAATGTTCACTGTAACAATGCTTGT	0 1	PASS	1	G	GAATGTTCACTGTAACAATGCTTGT	0/1	PASS		
62	BRCA1	17	41,224,833	1	G	C	0 1	PASS	1	G	C	0/1	PASS		
63	BRCA1	17	41,224,892	1	G	GT	1 0	PASS	0	-	-	-	-		
64	BRCA1	17	41,225,781	1	A	T	0 1	PASS	1	A	T	0/1	PASS		
65	BRCA1	17	41,225,783	1	A	T	0 1	PASS	1	A	T	0/1	PASS		
66	BRCA1	17	41,225,839	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
67	BRCA1	17	41,226,601	1	G	C	0 1	PASS	1	G	C	0/1	PASS		
68	BRCA1	17	41,226,675	1	A	T	0 1	PASS	1	A	T	0/1	PASS		
69	BRCA1	17	41,226,736	1	GGGGTT	G	0 1	PASS	1	GGGGTT	G	0/1	PASS		
70	BRCA1	17	41,227,083	1	C	CGGAA	0 1	PASS	1	C	CGGAA	0/1	PASS		
71	BRCA1	17	41,227,593	0	-	-	-	-	1	G	TG	0/1	PASS		
72	BRCA1	17	41,229,352	1	C	CT	0 1	PASS	0	-	-	-	-		
73	BRCA1	17	41,229,365	0	-	-	-	-	1	G	GT	0/1	PASS		
74	BRCA1	17	41,229,386	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
75	BRCA1	17	41,229,760	1	AT	A	0 1	PASS	0	-	-	-	-		
76	BRCA1	17	41,229,762	0	-	-	-	-	1	TT	A	0/1	PASS		

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idx	gene	chr	position	varcall	ref	alt	var	filter	varcall	ref	alt	var	filter
77	BRCA1	17	41,229,763	1	T	A	0 1	PASS	0	-	-	-	-
78	BRCA1	17	41,229,773	1	T	C	0 1	PASS	1	T	C	0/1	PASS
79	BRCA1	17	41,229,777	1	G	GT	1 1	PASS	0	-	-	-	-
80	BRCA1	17	41,229,780	0	-	-	-	-	1	G	GT	0/1	PASS
81	BRCA1	17	41,229,812	1	A	G	0 1	PASS	1	A	G	0/1	PASS
82	BRCA1	17	41,229,857	1	G	A	0 1	PASS	1	G	A	0/1	PASS
83	BRCA1	17	41,229,908	1	T	A	0 1	PASS	0	-	-	-	-
84	BRCA1	17	41,230,105	1	CT	C	0 1	PASS	0	-	-	-	-
85	BRCA1	17	41,230,119	0	-	-	-	-	1	TG	G	0/1	PASS
86	BRCA1	17	41,230,228	1	G	A	0 1	PASS	1	G	A	0/1	PASS
87	BRCA1	17	41,230,336	1	A	G	0 1	PASS	1	A	G	0/1	PASS
88	BRCA1	17	41,230,376	1	A	G	0 1	PASS	1	A	G	0/1	PASS
89	BRCA1	17	41,230,524	1	T	G	0 1	PASS	1	T	G	0/1	PASS
90	BRCA1	17	41,230,537	1	A	T	0 1	PASS	1	A	T	0/1	PASS
91	BRCA1	17	41,230,990	1	A	G	0 1	PASS	1	A	G	0/1	PASS
92	BRCA1	17	41,231,221	1	A	C	0 1	PASS	1	A	C	0/1	PASS
93	BRCA1	17	41,231,516	1	C	T	0 1	PASS	1	C	T	0/1	PASS
94	BRCA1	17	41,231,902	1	G	A	0 1	PASS	1	G	A	0/1	PASS
95	BRCA1	17	41,232,344	1	G	C	1 1	PASS	1	G	C	1/1	PASS
96	BRCA1	17	41,232,698	1	C	T	0 1	PASS	1	C	T	0/1	PASS
97	BRCA1	17	41,234,470	1	A	G	0 1	PASS	1	A	G	0/1	PASS
98	BRCA1	17	41,235,799	1	G	A	0 1	PASS	1	G	A	0/1	PASS
99	BRCA1	17	41,237,953	1	G	A	0 1	PASS	1	G	A	0/1	PASS
100	BRCA1	17	41,238,126	1	T	TACACAC	0 1	PASS	1	T	TACACAC	0/1	PASS
101	BRCA1	17	41,239,472	1	G	A	0 1	PASS	1	G	A	0/1	PASS
102	BRCA1	17	41,239,491	1	T	C	0 1	PASS	1	T	C	0/1	PASS
103	BRCA1	17	41,239,628	1	A	G	0 1	PASS	1	A	G	0/1	PASS
104	BRCA1	17	41,239,915	1	AT	A	0 1	PASS	0	-	-	-	-
105	BRCA1	17	41,239,928	0	-	-	-	-	1	TC	C	0/1	PASS
106	BRCA1	17	41,240,277	1	T	C	0 1	PASS	1	T	C	0/1	PASS
107	BRCA1	17	41,240,365	0	-	-	-	-	1	C	A	0/1	PASS
108	BRCA1	17	41,241,390	1	C	A	0 1	PASS	1	C	A	0/1	PASS
109	BRCA1	17	41,241,503	1	T	C	0 1	PASS	1	T	C	0/1	PASS
110	BRCA1	17	41,241,568	1	T	TC	0 1	PASS	1	T	TT	0/1	PASS
111	BRCA1	17	41,241,569	0	-	-	-	-	1	T	C	0/1	PASS
112	BRCA1	17	41,242,066	1	G	GA	0 1	PASS	0	-	-	-	-
113	BRCA1	17	41,242,074	0	AAAAG	A	0 1	OverlapConflict	0	-	-	-	-
114	BRCA1	17	41,242,075	0	AAAG	A	0 1	OverlapConflict	0	-	-	-	-

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idx	gene	chr	position	varcall	ref	alt	var	filter	varcall	ref	alt	var	filter		
115	BRCA1	17	41,242,083	0	-	-	-	-	1	AGAG	G	0/1	PASS		
116	BRCA1	17	41,242,285	1	T	G	0 1	PASS	1	T	G	0/1	PASS		
117	BRCA1	17	41,243,190	1	T	G	0 1	PASS	1	T	G	0/1	PASS		
118	BRCA1	17	41,244,000	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
119	BRCA1	17	41,244,435	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
120	BRCA1	17	41,244,936	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
121	BRCA1	17	41,245,237	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
122	BRCA1	17	41,245,466	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
123	BRCA1	17	41,247,122	1	A	ACCT	0 1	PASS	1	A	ACCT	0/1	PASS		
124	BRCA1	17	41,247,604	1	A	C	0 1	PASS	1	A	C	0/1	PASS		
125	BRCA1	17	41,248,164	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
126	BRCA1	17	41,248,484	1	G	C	0 1	PASS	1	G	C	0/1	PASS		
127	BRCA1	17	41,248,588	0	CA	C	0/1	Silver	0	-	-	-	-		
128	BRCA1	17	41,249,094	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
129	BRCA1	17	41,249,363	1	TA	T	0 1	PASS	1	TA	T	0/1	PASS		
130	BRCA1	17	41,250,047	1	C	CT	0 1	PASS	0	-	-	-	-		
131	BRCA1	17	41,250,678	1	C	CT	0 1	PASS	1	C	CT	0/1	PASS		
132	BRCA1	17	41,250,923	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
133	BRCA1	17	41,251,495	1	C	G	0 1	PASS	1	C	G	0/1	PASS		
134	BRCA1	17	41,251,646	1	T	A	0 1	PASS	1	T	A	0/1	PASS		
135	BRCA1	17	41,252,575	1	G	A	0 1	PASS	1	GTATATATATA	A	0/1	PASS		
136	BRCA1	17	41,252,589	0	-	-	-	-	1	AA	A	0/1	PASS		
137	BRCA1	17	41,252,591	0	-	-	-	-	1	C	T	0/1	PASS		
138	BRCA1	17	41,252,613	0	-	-	-	-	1	T	A	0/1	PASS		
139	BRCA1	17	41,252,615	0	-	-	-	-	1	T	C	0/1	PASS		
140	BRCA1	17	41,252,627	0	A	ATAT	0/1	Silver	0	-	-	-	-		
141	BRCA1	17	41,254,174	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
142	BRCA1	17	41,254,405	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
143	BRCA1	17	41,254,486	1	T	G	0 1	PASS	1	T	G	0/1	PASS		
144	BRCA1	17	41,254,664	0	-	-	-	-	1	A	C	0/1	PASS		
145	BRCA1	17	41,254,965	1	C	CT	0/1	Silver	1	C	CT	0/1	PASS		
146	BRCA1	17	41,255,102	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
147	BRCA1	17	41,255,111	1	A	T	0 1	PASS	1	A	T	0/1	PASS		
148	BRCA1	17	41,256,074	0	CA	C	0 1	OverlapConflict	0	-	-	-	-		
149	BRCA1	17	41,256,075	0	AAAAAAAAAAAAAG	A	0 1	OverlapConflict	0	-	-	-	-		
150	BRCA1	17	41,256,086	0	-	-	-	-	1	AGAAAAAAAAAAAAAG	GAA	0/1	PASS		
151	BRCA1	17	41,256,089	0	AAAAAAAAAGAAAAAG	A	0 1	OverlapConflict	0	-	-	-	-		
152	BRCA1	17	41,256,100	0	AAAG	A	0 1	OverlapConflict	0	-	-	-	-		

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153	BRCA1	17	41,257,134	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
154	BRCA1	17	41,257,458	1	A	C	0 1	PASS	1	A	C	0/1	PASS		
155	BRCA1	17	41,258,043	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
156	BRCA1	17	41,258,135	1	T	TA	0 1	PASS	0	-	-	-	-		
157	BRCA1	17	41,258,153	0	-	-	-	-	1	G	AG	0/1	PASS		
158	BRCA1	17	41,258,682	0	-	-	-	-	1	T	G	0/1	PASS		
159	BRCA1	17	41,259,049	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
160	BRCA1	17	41,259,079	0	A	ATT	0/1	Silver	0	-	-	-	-		
161	BRCA1	17	41,259,113	1	G	A	0 1	PASS	0	-	-	-	-		
162	BRCA1	17	41,259,390	0	-	-	-	-	1	A	T	0/1	PASS		
163	BRCA1	17	41,260,352	1	C	CA	0 1	PASS	1	C	CA	0/1	PASS		
164	BRCA1	17	41,260,808	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
165	BRCA1	17	41,261,058	1	T	TCTATCTATCTACCTAC	0 1	PASS	1	T	TCTATCTATCTACCTAC	0/1	PASS		
166	BRCA1	17	41,261,233	1	C	T	0 1	PASS	1	C	T	0/1	PASS		
167	BRCA1	17	41,261,915	0	-	-	-	-	1	C	CT	0/1	PASS		
168	BRCA1	17	41,263,044	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
169	BRCA1	17	41,263,117	1	C	CA	0 1	PASS	1	C	CA	0/1	PASS		
170	BRCA1	17	41,263,566	1	T	C	0 1	PASS	1	T	C	0/1	PASS		
171	BRCA1	17	41,264,146	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
172	BRCA1	17	41,264,364	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
173	BRCA1	17	41,264,739	1	C	T	0 1	PASS	1	CTGACTATCACAAGCTTGAAC	TTCATCAGTAGGGGTTT	0/1	PASS		
174	BRCA1	17	41,264,740	1	TGA	T	0 1	PASS	0	-	-	-	-		
175	BRCA1	17	41,264,743	1	CT	C	0 1	PASS	0	-	-	-	-		
176	BRCA1	17	41,264,749	1	C	G	0 1	PASS	0	-	-	-	-		
177	BRCA1	17	41,264,750	1	A	T	0 1	PASS	0	-	-	-	-		
178	BRCA1	17	41,264,751	0	A	AGGG	0 1	OverlapConflict	0	-	-	-	-		
179	BRCA1	17	41,264,753	0	C	T	0 1	OverlapConflict	0	-	-	-	-		
180	BRCA1	17	41,264,755	0	TGAAAC	T	0 1	OverlapConflict	0	-	-	-	-		
181	BRCA1	17	41,264,760	0	C	T	0 1	OverlapConflict	0	-	-	-	-		
182	BRCA1	17	41,265,776	1	A	G	0 1	PASS	1	A	G	0/1	PASS		
183	BRCA1	17	41,266,137	0	-	-	-	-	1	T	G	0/1	PASS		
184	BRCA1	17	41,266,407	1	C	CT	0 1	PASS	1	C	CT	0/1	PASS		
185	BRCA1	17	41,267,050	1	G	A	0 1	PASS	1	G	A	0/1	PASS		
186	BRCA1	17	41,267,518	1	CA	C	0 1	PASS	1	CA	C	0/1	PASS		
187	BRCA1	17	41,268,206	1	A	C	0 1	PASS	1	AAC	CACT	0/1	PASS		
188	BRCA1	17	41,268,208	1	C	CT	0 1	PASS	0	-	-	-	-		
189	BRCA1	17	41,270,229	1	T	G	0 1	PASS	1	T	G	0/1	PASS		
190	BRCA1	17	41,270,277	1	C	T	0 1	PASS	1	C	T	0/1	PASS		

<b>Color Code Description</b>	White = Identical Calls (based on the corresponding reported data)
	Green = Consistent Calls (based on Crystal's visual analysis of the reads supporting the variants)
	Red = Inconsistent Calls/NoCalls (based on Crystal's visual analysis of the reads supporting the variants)
	Yellow = Equivalent and Consistent Calls for Adjacent Rows (based on Crystal's visual analysis of the reads supporting the variants)
	Grey = Hard-to-assess Calls (insufficient data in Crystal's visual analysis of the reads supporting the variants)

Features & Coordinates				Illumina's Platinum Genomes Variants <a href="http://www.illumina.com/platinumgenomes">www.illumina.com/platinumgenomes</a>						Crystal Genetics' Variants <a href="http://www.crystalgenetics.com">www.crystalgenetics.com</a>					
idx	gene	chr	position	varcall	ref	alt	var	filter	varcall	ref	alt	var	filter		
191	BRCA1	17	41,270,463	1	G	A	0 1	PASS	1	G	A	0 1	PASS		
192	BRCA1	17	41,270,666	1	C	A	0 1	PASS	1	C	A	0 1	PASS		
193	BRCA1	17	41,270,778	1	C	CT	0 1	PASS	1	C	CT	0 1	PASS		
194	BRCA1	17	41,271,293	1	GA	G	0 1	PASS	0	-	-	-	-		
195	BRCA1	17	41,271,305	0	-	-	-	-	1	AG	G	0 1	PASS		
196	BRCA1	17	41,273,095	1	G	A	0 1	PASS	1	G	A	0 1	PASS		
197	BRCA1	17	41,273,348	1	T	C	0 1	PASS	1	T	C	0 1	PASS		
198	BRCA1	17	41,273,379	1	G	C	0 1	PASS	1	G	C	0 1	PASS		
199	BRCA1	17	41,273,537	1	A	C	0 1	PASS	1	A	C	0 1	PASS		
200	BRCA1	17	41,274,778	1	G	A	0 1	PASS	1	G	A	0 1	PASS		
201	BRCA1	17	41,274,906	1	G	A	0 1	PASS	1	G	aA	0 1	PASS		
202	BRCA1	17	41,275,081	1	G	GA	0 1	PASS	1	G	GA	0 1	PASS		
203	BRCA1	17	41,275,151	1	G	C	0 1	PASS	1	G	C	0 1	PASS		
204	BRCA1	17	41,275,366	0	GTTTTTT	G	0 1	Silver	0	-	-	-	-		
205	BRCA1	17	41,275,645	1	A	G	0 1	PASS	1	A	G	0 1	PASS		
206	BRCA1	17	41,276,247	1	A	G	0 1	PASS	1	A	G	0 1	PASS		
207	BRCA1	17	41,276,348	1	T	C	0 1	PASS	1	T	C	0 1	PASS		
208	BRCA1	17	41,277,187	1	G	C	0 1	PASS	1	G	C	0 1	PASS		