id	age	primary si	OS	ydtime	SEX	sexcode	grade	group7nan	group7nn	group4nn
1	58	C163	1	1.73	M	1	2	LC	1	1
2	85	C168	1	1.04	F	0	4	LCGCEP	7	3
3	82	C163	0	1.08	M	1	4	LCEP	5	2
4	78	C163	0	2.18	M	1	2	GC	2	1
5	68	C162	0	2.82	M	1	4		0	0
6	62	C163	0	4.38	M	1	4	LCGCEP	7	3
7	60	C163	0	1.53	F	0	4	LCGC	4	2
8	52	C162	0	10.58	F	0	4		0	0
9	81	C162	1	8.30	M	1	4	LC	1	1
10	57	C163	0	4.45	M	1	4		0	0
11	65	C163	0	0.98	F	0	4	LCGCEP	7	3
12	85	C164	1	0.05	F	0	4		0	0
13	79	C163	0	4.37	M	1	4		0	0
14	62	C163	1	1.43	M	1	4	LCGCEP	7	3
15	65	C164	0	4.47	F	0	4		0	0
16	64	C162	1	1.40	M	1	4	LCGCEP	7	3
17	66	C163	0	0.53	M	1	2		0	0
18	80	C162	1	3.10	F	0	2	LC	1	1
19	60	C165	0	8.04	F	0	4		0	0
20	75	C164	1	1.21	M	1	2	LCGCEP	7	3
21	67	C163	1	3.70	F	0	4	GCEP	6	2
22	70	C162	1	0.33	M	1	2		0	0
23	79	C163	1	0.50	M	1	4	LCGCEP	7	3
24	62	C162	0	1.06	M	1	4		0	0
25	79	C163	1	0.02	M	1	2	LCGCEP	7	3
26	85	C162	1	0.09	M	1	2		0	0
27	67	C165	1	3.09	M	1	2	LC	1	1
28	61	C162	0	0.98	F	0	4	LCEP	5	2
29	83	C163	1	0.54	M	1	4	LCGCEP	7	3
30	77	C162	0	5.48	M	1	4	LC	1	1
31	82	C163	0	7.13	M	1	4		0	0
32	76	C162	1	1.61	M	1	2	LCGCEP	7	3
33	84	C163	1	0.22	F	0	4	LCGCEP	7	3
34	81	C165	0	1.75	F	0	4		0	0
35	83	C163	1	0.08	M	1	4	LCGCEP	7	3
36	71	C163	1	1.29	M	1	4	LCGCEP	7	3
37	85	C162	0	0.86	F	0	4		0	0
38	63	C162	0	6.33	F	0	4		0	0
39	82	C162	1	1.17	M	1	2	EP	3	1
40	79	C164	0	3.10	F	0	4		0	0
41	76	C163	0	6.32	M	1	4		0	0
42	69	C163	1	4.42	M	1	4	LCGCEP	7	3
43	71	C163	1	1.90	F	0	4	LC	1	1
44	73	C163	0	0.39	F	0	2		0	0
45	70	C163	1	4.69	F	0	4	LCGCEP	7	3
46	62	C163	0	8.47	M	1	4		0	0
47	81	C163	1	0.79	F	0	4	LCGC	4	2
48	75	C163	1	1.75	M	1	4	LCEP	5	2
49	79	C163	0	2.73	F	0	4		0	0
50	70	C163	1	1.39	F	0	4	LCGC	4	2

51	47	C163	0	0.16	F	0	4	0	0
52	70	C163	0	2.80	F	0	4	0	0
53	55	C163	0	5.95	F	0	4	0	0
54	76	C163	0	8.27	F	0	2	0	0
55	69	C162	0	0.11	F	0	4	0	0
56	79	C163	1	2.88	M	1	4 EP	3	1
57	72	C163	0	3.22	M	1	2 GCEP	6	2
58	86	C163	1	0.13	M	1	4 GC	2	1
59	77	C164	1	6.39	F	0	4 GC	2	1
60	58	C163	0	2.83	M	1	4	0	0
61	80	C163	1	1.65	F	0	2 LCGCEP	7	3
62	67	C163	0	4.71	M	1	4 LCGCEP	7	3
63	75	C162	1	0.05	M	1	4	0	0
64	55	C163	1	0.41	F	0	4 GCEP	6	2
65	36	C163	0	9.25	M	1	4	0	0
66	60	C163	0	7.52	F	0	4	0	0
67	78	C162	0	4.18	F	0	4	0	0
68	71	C163	0	10.10	F	0	4	0	0
69	68	C163	1	7.92	M	1	4 GC	2	1
70	68	C163	1	0.48	F	0	4 LCGCEP	7	3
71	63	C163	1	1.50	M	1	2 LCGCEP	7	3
72	76	C163	1	2.88	M	1	2 GC	2	1
73	54	C168	1	0.06	F	0	4 LCGCEP	7	3
74	53	C162	0	8.81	M	1	2 LC	1	1
75	66	C161	1	1.37	F	0	4 LCGCEP	7	3
76	53	C163	1	1.26	F	0	4 LCEP	5	2
77	58	C163	0	1.87	M	1	2	0	0
78	61	C163	0	7.58	F	0	2	0	0
79	67	C163	0	2.25	M	1	2 LC	1	1
80	80	C162	0	6.77	M	1	4	0	0
81	84	C163	1	10.23	F	0	4	0	0
82	76	C163	0	1.22	F	0	2	0	0
83	67	C162	0	10.47	F	0	4	0	0
84	79	C163	0	8.24	M	1	4 GC	2	1
85	74	C168	1	2.09	M	1	2	0	0
86	69	C162	1	1.73	M	1	4 LCEP	5	2
87	73	C163	1	1.91	M	1	4 GC	2	1
88	69	C163	0	7.66	M	1	4 GC	2	1
89	55	C163	0	5.92	M	1	2	0	0
90	58	C163	0	5.48	M	1	4	0	0
91	60	C163	0	2.44	M	1	4	0	0
92	81	C164	1	0.20	M	1	2 GCEP	6	2
93	60	C163	1	0.93	M	1	2 LCGC	4	2
94	59	C162	1	1.16	F	0	4 GCEP	6	2
95	75	C163	1	2.90	M	1	2 LC	1	1
96	96	C162	1	0.17	M	1	2 LCGC	4	2
97	49	C162	0	7.83	F	0	4	0	0
98	61	C162	0	2.62	M	1	4 LCEP	5	2
99	75	C162	0	0.23	F	0	4	0	0
100	76	C163	0	2.17	F	0	2	0	0
101	45	C163	0	1.96	F	0	4 EP	3	1

102	66 C163	1	1.25 F	0	4 LCGCEP	7	3
103	66 C168	1	0.23 M	1	4 LCGCEP	7	3
104	46 C163	0	9.68 F	0	4	0	0
105	74 C163	0	4.07 M	1	4 LC	1	1
106	53 C163	1	1.14 F	0	4 GC	2	1
107	67 C162	1	1.72 M	1	2 LCGC	4	2
108	67 C168	1	2.25 M	1	4 GC	2	1
109	78 C164	1	3.53 M	1	2 GCEP	6	2
110	85 C163	0	0.16 M	1	4 LCEP	5	2
111	73 C168	1	1.61 F	0	4 LCGCEP	7	3
112	52 C162	1	2.53 F	0	4 LCEP	5	2
113	63 C162	1	3.45 M	1	4	0	0
114	64 C162	0	3.45 M	1	4	0	0
115	71 C163	0	0.16 M	1	4 GCEP	6	2
116	83 C163	1	2.61 M	1	4	0	0
117	76 C164	0	4.93 M	1	2	0	0
118	44 C168	0	8.79 F	0	2	0	0
119	87 C163	1	4.29 M	1	4	0	0
120	68 C164	1	6.36 M	1	2 LCGC	4	2
121	71 C162	0	5.88 M	1	4	0	0
122	57 C162	1	1.20 M	1	4 GCEP	6	2
123	55 C163	0	5.98 M	1	4 LCGC	4	2
124	74 C163	1	0.15 M	1	2	0	0
125	75 C163	0	3.18 M	1	4 LCGCEP	7	3
126	75 C162	1	1.22 M	1	4 LCGCEP	7	3
127	49 C163	0	0.15 F	0	4	0	0
128	56 C163	0	6.80 F	0	4 LC	1	1
129	67 C162	0	1.48 M	1	4 LC	1	1
130	78 C163	0	2.93 M	1	2 GC	2	1
131	75 C163	1	0.18 F	0	4 GC	2	1
132	66 C162	0	0.28 M	1	4	0	0
133	63 C163	0	2.11 F	0	4	0	0
134	43 C163	0	1.10 F	0	4 GCEP	6	2
135	43 C163	0	0.87 F	0	4	0	0
136	79 C162	0	2.42 M	1	4	0	0
137	89 C163	1	0.65 M	1	2 GCEP	6	2
138	81 C162	1	2.39 M	1	4 LC	1	1
139	81 C163	1	2.39 F	0	4 LCEP	5	2
140	73 C168	1	0.59 F	0	4 LCGCEP	7	3
141	60 C162	0	8.11 M	1	2 LCGCEP	7	3
142	71 C163	0	8.51 F	0	2	0	0
143	49 C163	0	7.15 F	0	4 LCGCEP	7	3
144	59 C162	0	9.70 M	1	2 LC	1	1
145	73 C168	1	1.10 F	0	4 LCGCEP	7	3
146	80 C163	0	4.36 M	1	4 LCGCEP	7	3
147	70 C163	0	3.25 M	1	4 LCGCEP	7	3
148	79 C165	0	8.91 M	1	2 LC	1	1
149	63 C162	1	4.54 F	0	4 GCEP	6	2
150	52 C163	0	6.27 F	0	4 EP	3	1
151	73 C163	0	2.41 M	1	4	0	0
152	71 C161	1	1.46 M	1	4 GCEP	6	2

153	77 C163	0	5.50 M	1	4 LCGC	4	2
154	68 C162	0	0.39 M	1	2	0	0
155	54 C163	0	5.07 M	1	2	0	0
156	57 C163	0	8.05 M	1	4	0	0
157	57 C164	1	0.91 M	1	4 LCEP	5	2
158	64 C163	0	0.09 M	1	4	0	0
159	54 C162	0	8.77 F	0	4 LCGC	4	2
160	71 C168	1	0.95 M	1	4 LCGCEP	7	3
161	75 C164	0	0.29 M	1	2 LC	1	1
162	69 C162	1	1.40 M	1	2	0	0
163	71 C163	0	9.34 M	1	2	0	0
164	37 C163	0	7.54 F	0	4	0	0
165	83 C163	1	4.36 M	1	4 GCEP	6	2
166	68 C163	0	8.35 F	0	4 LCEP	5	2
167	55 C163	0	3.52 M	1	2 LC	1	1
168	55 C168	1	1.28 M	1	4 LCGCEP	7	3
169	69 C163	0	5.78 M	1	4 LC	1	1
170	77 C163	1	4.82 F	0	4 LCEP	5	2
171	79 C162	1	0.37 M	1	2 GC	2	1
172	88 C168	1	0.13 F	0	2	0	0
173	53 C163	0	4.32 M	1	2	0	0
174	67 C163	0	3.12 M	1	2 LC	1	1
175	59 C165	1	0.42 F	0	4 LCEP	5	2
176	72 C162	1	0.59 M	1	4 LCGCEP	7	3
177	72 C163	1	0.68 F	0	4 LCGCEP	7	3
178	31 C163	1	3.76 M	1	4 LC	1	1
179	73 C163	1	0.32 M	1	4 GC	2	1
180	85 C163	1	2.54 M	1	2 GC	2	1
181	69 C163	0	3.53 M	1	4 EP	3	1
182	73 C168	1	2.16 M	1	4 LCGCEP	7	3
183	56 C164	1	1.17 M	1	2 GCEP	6	2
184	65 C163	0	9.60 F	0	4	0	0
185	55 C163	1	2.65 M	1	4 LCGCEP	7	3
186	79 C168	1	2.70 M	1	4 LCGCEP	7	3
187	57 C162	0	3.80 M	1	4	0	0
188	60 C162	0	3.05 F	0	4 GC	2	1
189	69 C163	0	5.72 F	0	4 GC	2	1
190	53 C164	1	3.04 F	0	2 GCEP	6	2
191	79 C163	1	1.36 M	1	2 LCGCEP	7	3
192	67 C162	1	4.03 M	1	4	0	0
193	75 C163	1	0.11 M	1	4 GC	2	1
194	75 C162	0	0.54 F	0	4 LCGC	4	2
195	79 C168	1	0.30 F	0	4 LCGCEP	7	3
196	81 C163	0	2.35 M	1	2	0	0
197	46 C168	0	8.69 F	0	4	0	0
198	57 C168	1	0.45 F	0	4 LCGCEP	7	3
199	81 C165	0	0.72 M	1	2 LC	1	1
200	71 C166	1	1.22 M	1	4 GC	2	1
201	45 C165	0	7.45 M	1	4 LC	1	1
202	79 C163	1	6.24 M	1	4	0	0
203	72 C162	0	4.38 F	0	4	0	0

204	50 C162	1	1.40 F	0	4 LCEP	5	2
205	56 C163	1	5.37 F	0	4 LC	1	1
206	72 C163	1	0.77 M	1	4 LCGCEP	7	3
207	72 C163	1	1.28 M	1	4 LCGCEP	7	3
208	72 C163	1	0.11 M	1	2	0	0
209	33 C163	0	0.62 M	1	2 LCGC	4	2
210	56 C163	0	4.17 M	1	4	0	0
211	48 C163	0	3.79 M	1	4 LCGC	4	2
212	56 C163	0	3.63 M	1	4 LCGCEP	7	3
213	63 C163	1	0.98 M	1	4	0	0
214	82 C163	1	0.77 F	0	4 LCGCEP	7	3
215	40 C164	0	0.82 M	1	4	0	0
216	76 C162	0	0.53 M	1	4 GCEP	6	2
217	34 C163	0	0.37 F	0	4	0	0
218	54 C163	0	0.09 M	1	4	0	0
219	64 C163	1	7.72 M	1	4 LC	1	1
220	81 C163	1	3.94 F	0	4 LCGCEP	7	3
221	57 C163	0	3.45 M	1	4 LCGCEP	7	3
222	62 C163	0	3.18 F	0	4 GC	2	1
223	82 C163	1	0.81 F	0	4 LCGCEP	7	3
224	52 C162	0	5.05 F	0	4 EP	3	1
225	62 C162	0	7.64 M	1	4 GCEP	6	2
226	80 C164	0	7.55 F	0	2	0	0
227	72 C163	1	2.00 F	0	4 LCGC	4	2
228	66 C168	1	0.95 F	0	4 LCGCEP	7	3
229	63 C163	1	4.95 M	1	2 LCGCEP	7	3
230	64 C168	0	3.94 M	1	4 LCGCEP	7	3
231	82 C163	0	0.92 F	0	4 LCGCEP	7	3
232	56 C163	0	0.16 F	0	4 LCGC	4	2
233	82 C163	0	0.13 M	1	4 LCGCEP	7	3
234	54 C161	1	4.01 M	1	2 LC	1	1
235	71 C163	1	0.58 M	1	2 GCEP	6	2
236	68 C163	0	2.32 F	0	4 LCGCEP	7	3
237	58 C163	1	0.66 F	0	2 LC	1	1
238	60 C164	0	0.94 M	1	4 GC	2	1
239	56 C163	0	7.58 M	1	4 LCGC	4	2
240	77 C163	1	2.24 F	0	4 LCGCEP	7	3
241	59 C163	1	0.37 F	0	4 LCGCEP	7	3
242	61 C162	0	0.48 M	1	4 LCEP	5	2
243	71 C163	1	2.92 M	1	4 LC	1	1
244	45 C162	0	3.97 F	0	4	0	0
245	76 C162	1	4.12 M	1	4	0	0
246	58 C165	0	7.80 M	1	2 LCGCEP	7	3
247	83 C163	1	2.07 F	0	2 LCGC	4	2
248	43 C162	0	0.58 F	0	4 LCGCEP	7	3
249	68 C163	0	7.27 F	0	2	0	0
250	58 C164	0	5.31 M	1	2 GCEP	6	2
251	73 C164	1	6.25 F	0	2	0	0
252	56 C163	0	4.75 M	1	4 LCGCEP	7	3
253	48 C163	0	1.97 M	1	4 LCGCEP	7	3
254	74 C163	0	2.28 M	1	4	0	0

255	71 C163	0	3.34 M	1	4	0	0
256	79 C163	0	2.36 F	0	2 LCGC	4	2
257	59 C162	1	1.01 F	0	4 LCGCEP	7	3
258	77 C164	1	0.31 M	1	2 EP	3	1
259	70 C163	0	0.18 F	0	2 LCGCEP	7	3
260	59 C165	0	0.17 F	0	4	0	0
261	55 C161	1	1.18 M	1	4 EP	3	1
262	71 C163	0	2.84 F	0	4	0	0
263	58 C162	0	5.57 F	0	4	0	0
264	56 C162	0	0.57 F	0	4	0	0
265	73 C165	1	0.23 M	1	4 LCGCEP	7	3
266	69 C163	1	3.48 F	0	4 LCGCEP	7	3
267	66 C163	1	6.69 M	1	2 LCGC	4	2
268	81 C164	1	2.55 M	1	4 EP	3	1
269	71 C163	0	7.34 M	1	4 LCGC	4	2
270	60 C163	1	1.16 M	1	4 LCGC	4	2
271	65 C163	0	5.58 M	1	2	0	0
272	63 C164	0	1.01 M	1	2 LCGCEP	7	3
273	73 C164	1	3.65 F	0	4 LCGCEP	7	3
274	58 C163	0	4.06 M	1	4	0	0
275	75 C162	1	0.51 F	0	4 LCGCEP	7	3
276	64 C163	1	0.86 F	0	4 LCGCEP	7	3
277	65 C166	1	0.18 M	1	4 LCGCEP	7	3
278	59 C161	0	1.05 M	1	2	0	0
279	63 C163	0	2.45 M	1	4	0	0
280	44 C163	0	7.91 M	1	4	0	0
281	73 C163	1	8.87 M	1	4 LCGCEP	7	3
282	86 C168	1	1.54 M	1	2 LCGC	4	2
283	42 C168	0	9.98 M	1	4	0	0
284	73 C163	0	3.24 F	0	4	0	0
285	58 C163	1	3.93 M	1	4	0	0
286	35 C163	0	6.33 F	0	4 GCEP	6	2
287	58 C163	0	8.92 M	1	4	0	0
288	57 C165	0	8.19 M	1	2	0	0
289	71 C162	0	7.78 F	0	4	0	0
290	80 C162	0	6.08 F	0	4	0	0
291	65 C162	0	7.58 M	1	4	0	0
292	50 C162	0	7.71 F	0	4 LCGCEP	7	3
293	62 C164	0	5.53 M	1	2	0	0
294	54 C163	1	0.85 F	0	4 LCGCEP	7	3
295	66 C163	1	0.88 M	1	4 LCEP	5	2
296	73 C163	0	3.05 M	1	4 LCGCEP	7	3
297	82 C168	1	0.84 F	0	4 LCGCEP	7	3
298	86 C162	0	0.65 M	1	4 LCEP	5	2
299	68 C164	0	0.87 M	1	4 LC	1	1
300	61 C165	0	0.19 M	1	2 LCGCEP	7	3
301	72 C163	0	4.61 M	1	2	0	0
302	83 C163	1	1.35 M	1	2	0	0
303	74 C163	0	3.98 F	0	4 LCEP	5	2
304	86 C162	1	1.49 M	1	2	0	0
305	51 C162	0	5.79 M	1	4 LCGCEP	7	3

306	67 C163	0	1.40 M	1	2	0	0
307	76 C165	1	7.70 M	1	2 LC	1	1
308	72 C163	1	8.76 M	1	4 LCGCEP	7	3
309	53 C163	0	7.96 M	1	2 GC	2	1
310	75 C163	1	1.01 M	1	2 LCGCEP	7	3
311	44 C168	1	0.26 M	1	4 LCGCEP	7	3
312	52 C162	1	2.65 M	1	4 LCGCEP	7	3
313	74 C163	1	1.98 M	1	4 LCGCEP	7	3
314	51 C162	0	4.27 F	0	4	0	0
315	70 C163	0	4.36 M	1	4	0	0
316	73 C163	1	1.73 M	1	4 LCGCEP	7	3
317	56 C162	0	3.95 F	0	4 LCGCEP	7	3
318	80 C162	1	0.32 M	1	4	0	0
319	84 C163	1	2.83 M	1	2	0	0
320	50 C163	0	3.10 F	0	2 LCGC	4	2
321	55 C164	0	2.06 M	1	4	0	0
322	71 C163	0	0.43 M	1	4 LCGCEP	7	3
323	64 C163	0	1.00 M	1	4 LCGCEP	7	3
324	68 C163	0	1.07 F	0	4 GC	2	1
325	36 C163	0	0.10 M	1	2	0	0
326	65 C162	1	0.14 F	0	4 LCGCEP	7	3
327	57 C163	0	0.62 M	1	2 GCEP	6	2
328	76 C163	0	5.76 M	1	2 GC	2	1
329	52 C163	0	2.21 M	1	4 GC	2	1
330	53 C165	0	8.51 M	1	4	0	0
331	69 C165	1	0.86 F	0	2 LCEP	5	2
332	76 C165	1	8.41 M	1	4	0	0
333	24 C164	1	1.67 M	1	4	0	0
334	78 C163	1	0.15 M	1	2	0	0
335	50 C163	1	2.83 M	1	4 GC	2	1
336	63 C163	0	3.27 M	1	2	0	0
337	49 C162	1	2.12 M	1	4 LCGCEP	7	3
338	77 C162	1	0.50 M	1	4 GCEP	6	2
339	78 C164	0	6.22 F	0	2 LCGCEP	7	3
340	69 C168	1	1.76 M	1	4 LCGCEP	7	3
341	74 C162	0	5.81 M	1	2	0	0
342	72 C163	1	2.73 M	1	4	0	0
343	49 C163	0	4.96 F	0	4	0	0
344	81 C164	1	1.25 F	0	4 LCGC	4	2
345	75 C164	0	2.97 F	0	2 GC	2	1
346	42 C163	1	1.41 M	1	4 GCEP	6	2
347	58 C168	1	1.15 F	0	4 LCGC	4	2
348	61 C163	1	0.14 F	0	4 LCGCEP	7	3
349	56 C163	0	2.30 M	1	4 GCEP	6	2
350	57 C168	1	0.39 M	1	4 LCGCEP	7	3
351	47 C164	0	1.64 M	1	4 LCGCEP	7	3
352	58 C162	1	0.39 M	1	4 LCGCEP	7	3
353	53 C163	0	1.20 M	1	4 LCGCEP	7	3
354	69 C163	0	0.61 F	0	4 LCGC	4	2
355	64 C162	1	4.64 F	0	2 GC	2	1
356	71 C163	1	1.11 F	0	4 LCGCEP	7	3

357	78 C163	0	4.65 M	1	2 LC	1	1
358	27 C163	0	1.30 F	0	4	0	0
359	66 C162	0	8.55 M	1	4	0	0
360	44 C163	0	6.35 F	0	4	0	0
361	77 C162	0	3.23 F	0	4 LC	1	1
362	82 C162	1	0.26 M	1	4	0	0
363	63 C163	0	1.20 F	0	4 LCGC	4	2
364	60 C168	1	0.19 M	1	4 LCGC	4	2
365	71 C163	1	6.21 M	1	4 LCEP	5	2
366	78 C163	0	5.13 M	1	4 EP	3	1
367	73 C163	1	7.79 F	0	4	0	0
368	42 C162	0	3.69 M	1	4	0	0
369	85 C162	0	6.79 F	0	4	0	0
370	77 C163	1	3.53 M	1	4 LCGCEP	7	3
371	79 C168	1	0.25 M	1	4 LCGCEP	7	3
372	71 C162	1	3.10 M	1	2	0	0
373	37 C162	0	6.02 M	1	4	0	0
374	56 C163	0	6.50 M	1	2 LC	1	1
375	45 C162	0	5.83 M	1	4 LCGC	4	2
376	51 C163	0	1.25 M	1	4 EP	3	1
377	70 C162	0	3.67 F	0	4	0	0
378	61 C163	1	1.38 F	0	4 LCGCEP	7	3
379	59 C163	0	1.04 M	1	4 LCGCEP	7	3
380	79 C162	0	0.84 F	0	2	0	0
381	78 C163	0	4.19 M	1	4 LCGCEP	7	3
382	74 C162	0	0.57 M	1	2 LCGCEP	7	3
383	53 C162	0	0.31 M	1	2 LC	1	1
384	79 C163	1	1.52 M	1	2 LCGC	4	2
385	54 C165	0	8.81 M	1	4	0	0
386	66 C163	1	0.89 F	0	4 LC	1	1
387	68 C162	1	7.36 M	1	2 EP	3	1
388	70 C163	1	1.01 M	1	2	0	0
389	45 C163	1	0.22 M	1	4 LCGCEP	7	3
390	62 C163	1	0.97 F	0	4 LCGCEP	7	3
391	50 C163	1	1.05 F	0	2 LCGCEP	7	3
392	61 C161	1	4.00 M	1	4 GCEP	6	2
393	81 C163	0	4.25 F	0	4 GCEP	6	2
394	73 C162	1	0.22 F	0	2 LCGC	4	2
395	70 C164	1	4.05 M	1	4 LC	1	1
396	66 C163	0	3.61 M	1	4	0	0
397	59 C163	0	1.41 F	0	4 LCGCEP	7	3
398	91 C163	0	1.21 F	0	4 GC	2	1
399	73 C164	0	0.61 M	1	2 GC	2	1
400	61 C161	0	0.39 M	1	2 GC	2	1

nnstage7	T7ajcc	N7ajcc	stage7ajcc	N7ajcc4	nngroup2	resectionln2
2A	2	2	2B	2	1	1
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2B	3	1	2B	1	1	2
1A	1B	0	1A	0	0	2
3B	3	2	3A	2	2	2
3B	4A	2	3B	2	1	2
1A	1B	0	1A	0	0	2
2B	3	2	3A	2	1	2
1A	1B	0	1A	0	0	1
3B	3	3A	3B	3	2	1
1B	2	0	1B	0	0	2
1A	1A	0	1A	0	0	1
3C	4A	3A	3C	3	2	2
1B	2	0	1B	0	0	1
4	3	3A	4	3	0	2
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2A	2	1	2A	1	1	2
2B	4A	0	2B	0	0	2
3C	4A	3B	3C	3	2	2
3B	4A	2	3B	2	2	2
2A	3	0	2A	0	0	1
3C	4A	3B	3C	3	2	2
2B	4A	0	2B	0	0	1
4	4B	3A	4	3	2	2
1B	2	0	1B	0	0	2
3A	4A	2	3B	2	1	2
2A	1A	3A	2B	3	2	1
3C	4A	3A	3C	3	2	2
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4	4A	3B	4	3	2	2
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4	4A	3B	4	3	2	2
3B	3	3B	3B	3	2	2
2B	4A	0	2B	0	0	1
1A	1B	0	1A	0	0	2
3B	4B	1	3B	1	2	1
1B	2	0	1B	0	0	2
1A	1B	0	1A	0	0	1
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3B	3	3A	3B	3	2	1
1A	1A	0	1A	0	0	1
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3B	4A	3B	3C	3	1	2

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2A	3	0	2A	0	0	1
1A	1B	0	1A	0	0	2
1A	1A	0	1A	0	0	1
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3A	3	2	3A	2	2	1
4	3	1	4	1	1	1
2A	2	2	2B	2	1	2
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3A	2	3B	3A	3	2	2
3C	4A	3A	3C	3	2	2
2A	3	0	2A	0	0	1
3C	4B	3A	3C	3	2	1
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3B	3	3A	3B	3	2	2
3B	3	3A	3B	3	2	2
2A	2	1	2A	1	1	1
3C	4A	3B	3C	3	2	1
3A	4A	2	3B	2	1	1
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2A	2	2	2B	2	1	2
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2B	3	1	2B	1	1	2
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1B	2	0	1B	0	0	2
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3B	4A	2	3B	2	1	2
3C	4B	3A	3C	3	2	2
3A	4A	1	3A	1	1	1
3A	3	2	3A	2	1	1
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3A	3	3A	3B	3	2	2
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1B	1A	1	1B	1	2	1

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3C	4A	3A	3C	3	2	2
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2B	2	2	2B	2	2	1
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3C	4A	3B	3C	3	2	2
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2A	2	1	2A	1	1	2
2A	2	1	2A	1	1	1
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3A	3	3A	3B	3	2	2

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3B	4A	1	3A	1	2	2
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3B	4A	1	3A	1	2	1
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3C	4B	3B	3C	3	2	2
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2B	3	1	2B	1	1	1
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3C	4B	3B	3C	3	2	2
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3C	4A	3B	3C	3	2	2
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4	4A	3B	4	3	2	2
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3C	4B	3A	3C	3	2	2
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3B	3	2	3A	2	2	1
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3C	4A	3B	3C	3	2	2

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1B	2	0	1B	0	0	2
1A	1B	0	1A	0	0	2
2B	3	1	2B	1	1	2
4	3	0	4	0	0	1
2A	1B	2	2A	2	1	2
3B	4A	3B	3C	3	1	1
2B	2	2	2B	2	2	2
3A	4A	1	3A	1	2	2
2B	4A	0	2B	0	0	1
1A	1A	0	1A	0	0	2
1A	1A	0	1A	0	0	1
3C	4A	3B	3C	3	2	2
3B	3	3B	3B	3	2	2
2A	3	0	2A	0	0	2
2A	3	0	2A	0	0	2
2A	2	1	2A	1	1	1
3A	3	2	3A	2	1	2
2B	3	1	2B	1	2	2
1A	1B	0	1A	0	0	2
4	3	3A	4	3	2	1
3C	4A	3A	3C	3	2	2
2A	3	0	2A	0	0	2
3B	3	3A	3B	3	2	2
3C	4A	3A	3C	3	2	2
2B	3	1	2B	1	1	1
3B	4A	2	3B	2	1	1
1A	1B	0	1A	0	0	1
4	4A	3A	4	3	1	1
4	3	1	4	1	2	2
1A	1A	0	1A	0	0	1
3C	4A	3A	3C	3	2	1
3C	4A	3B	3C	3	2	2
4	4A	3B	4	3	2	2
3A	3	2	3A	2	2	2
3A	3	1	2B	1	2	2
2A	1B	2	2A	2	1	2
2B	3	2	3A	2	1	2
1B	2	0	1B	0	0	1
4	4A	3B	4	3	2	1
1B	1B	1	1B	1	1	2
2B	3	1	2B	1	1	1
3A	4A	3A	3C	3	1	2