Code	E1	E2	E3	E4	Pscie	Psalt	Porga	Pmicr	Pcells	Pfrac	Pmen	Pdial	Penzl	Pmito	Pmeic	Pprot	Q1sal	Q2sci	Q3sal	Q4orc	Q5mi	Q6cel	Q7fra	Qmer	Qenz	Qmitc	Crick	Dialy	Enzyr	Extra	Total	%	Grade
0	100	100	100	200	5	5	5	10	5	5	5	5	5	5	5	5	20	20	20	20	20	20	20	20	20		20	50	70	0	885	100.0	
2	49	94.5	74	170	5	5	4.5	10	5	5	5	5			4.5	5	17	18	10.5	19	14	15	14	14		20	18	34	64	15	714	80.7	B
7	94	79.5	72	144	5	5	5	9.5	5	5	4.5	5	5	5	4.5	5	20	19.5	18.5	17		12	18	17	12	13	20	38	66	33	757	85.5	B
8	40	82	68	131	4.5	5	4.5	10	5	5	5	5	5	5	5	5	19	17	15	16.5		18	18	18	15	20	19	43.5	67	18	689	77.9	C
10	54	76	79	123	5	5	4.5	5	5	4.5	5	5	4.5				18		16.5	15	13.5	19	19	15	20	16	19	37.5	48		632	71.4	C
11	87	95.5	84	138	5	5	5	9.5	5	4.5	5	5	5	5	5	4.5	17		17.5	17.5	15	17	19	20	19	18	19	46	66	24	783	88.5	B
12	62	87.5	63	143	5	5			4.5	5	5	5	5	4.5	5	4	20	12		8	16	12	18	13	13	20	19	40.5	60	3	658	74.4	C
13	76	83.5	50	128	5	5	4.5	9.5	5	5	5	5	5		5	5	13	16	16	19	15.5	19	19	15	20		18.5	40.5	60		668	75.5	C
14	56	96	83.5	157	5	5	5	10	5	4.5	5	5	5		4.5	5	19.5	20	19.5	19	15	20	18		20	18	19	47.5	70	15	772	87.2	B
15	99	99.5	88	183	5	5	5	9.5	5	5		5	4.5	5	5	5	19	17	20	19	18	20	18		20	20	20	25	66	30	841	95.0	A
16	53	66	73	128	5	5	5	10	5	5	5	5	5	5	5	5	18		18	16	11	17	19	20	12	12	18	39	60	9	654	73.9	C
17	84	95.5	87	176	4	5	5	8.5	5	5	4.5	5	5	5	5	5	20	18		15	14	18	18	20	16	18	19.5	30	55	18	784	88.6	B
18	71	71	57	117	5	5	5	10	5	4.5	5	5	5	5	5	4.5	20	17	18.5	17.5	16.5	18	19	18	14		20	40.5	64	15	678	76.6	C
19	45	80	26	108	4.5	4.5	4.5	9	5	4.5	5	5		5			19	11	17	19	10	12.5	18	12	9		20	39	53		546	61.6	D
20	53	66	52		4.5	5	5	9	5	4.5	5		5	5	5		19	14	17	16	13.5	12	15	10	8		20	39	60	9	477	53.8	F
21	37	79.5	67	109	5	5	4	9	5	4.5	4.5	5	4	5	4.5	5	13	13	15	16.5	13		18	10	6	12	17	32	52	13	584	65.9	D
22	52	70.5	82	128	5	5	5	9	5	4.5	5		5	5	5	4.5	17	18	19	19	15		14	20	14	16	20	44.5	57	9	673	76.0	C
23	57	63	68	135	4.5	5	4.5	9.5	5	5	5	5	4.5	5		5	19	17	18	19	13	14	14	15	16		18.5	40	65	6	656	74.1	C
29	63	57	61	105	5	5	5	10		3.5	3.5	5	4.5		5		20	17	17	19		19	12	13.5	14	11	18	42	51		586	66.2	D
30	19	24	30	100	4	4.5	2	9	5	3	4	2	4.5	3.5	4.5	5	18	12	10.5	10	11	9	3		7	4	17	15	23		364	41.1	F
31	61	51	42	93	5	5	4.5	9	5	4.5	4.5	5		3			18	16	16	17.5	17	12	11	13.5	7		19	33	46		519	58.6	F
33	70	44	44	152	5	5	4.5	9	5	4.5	5	5	5	5			19	18	18	19	18	19	19	17	17		19	41	29	6	622	70.3	C
34	38	64	63	119		5	5	9	5	5		5					14	12	17.5	17		7	10	14	5	18	19	31.5	46		529	59.8	D
35	44	70	65		5	5	4	10	5	4.5	5	5	5	5	5		19	17	18	18	17	20	20		19	16	19	34	61		516	58.2	F
36	96	99	95	171	5	5	5	10	5	4.5	5	5	4.5	5	5	5	20	19	18.5	19	20	20	20	18	17		20	48.5	68	15	848	95.8	A
37	84	79	80	113	5	5	5	10	5	4.5	5	5	5		4		18	17	17.5	18	16	17	16	16	17		17.5	32	50		662	74.7	C
38	84	90	71	151	5	5	5	9.5	5	4.5	5	5	5	5	5	5	20	19.5	17.5		18	20	20	17	18	20	19.5	45	70	12	777	87.7	B
39	83	80	79	122	5	5	5	9.5	5	4.5	4.5	5	4.5	5	4	5	15		17	16.5	12.5	13	18	18	10	15.5	17	42	61	24	706	79.7	B
40	61	60	54	85	5	5		9.5	4.5	5	4.5	3.5	5	5	5	5	13	13	18	13		14	17	14	20	10	20	34	68	3	574	64.9	D
41	92	91	82	172	5	5	5	10	5	4.5	4.5	5	5	5	5	4.5	20	18.5	18.5	19		13	19	20	17	20	19.5	49	70	33	837	94.6	A
42	42	68	51.5	99	5	5	5	10	5	5	4	4	4.5	5	5	5	19	17	17	17.5		11	17	14	14	13	17.5	31	64	3	578	65.3	D
42	90	101	83	173	5	5	5	10	5	4.5	5	5	4.5	5	5	4.5	20	20	18.5	18		16	19	18	19	16	20	44	64	35	838	94.7	A
43	69	68	57	118	5	5	5	9	5	4.5	4.5	5	5	5	5		14	17	16.5	16.5		12	13	12	15	11	20	42	63	21	643	72.7	C
44	83	84	68.5	139	5	4.5	5	9.5	5	5	5	5	5	5	5	5	13	16	17	17.5	11	14	17		12	18	54	42	50	24	744	84.1	B
45	67	83	64	124	5	5	5	9	5	4.5	5	5	5	5	5	4.5	20	15	18	16	19	19	17	19	15		20	43.5	52		675	76.2	C
51	50	95	79.5	159	5	4.5	5	9	5	4	4.5	5	5	5	5	5	16	16.5	17.5	18		19	19	16	16	20	20	36	56	18	734	82.9	B
53	27	24	40	104	4.5	4.5	5	10	4.5	4.5	4.5	5	5		3.5	5	20	14	19	16	11.5	12	15	14		9	19	30			431	48.6	F