16-1 TA	32.8				11.47	94.09	885.6	2134.36	2.46	109.49	18.07	30.64	3.31	46.03
16-1 TB	13.86				11.3	249.72	5474.7	4033.26	4.16	483.71	79.97	269.44	3.82	184.89
31-1 DA	25.76				11.08	167.18	515.2	1236.48	11.08	279	19.58	277.43	34	641.94
31-1 DB	18.12				14.24	222.08	308.04	724.8	8.52	258.57	16.49	31.53	13.41	93.32
33-2 BA	2.24		4.95	5.5	11.1	70.79	241.42	1543.36	1.66	86.11	40.63	39	3.25	37.59
33-2 BB	10.41		1.8	6.33	11.22	26.69	3664.32	1894.62	3.12	81.2	2.91	109.93	6.97	221.42
33-2 BC	5.8		1.95	6.1	11.04	113.83	1363	1705.2	9.34	117.86	9.69	10.9	4.18	50.52
33-2 LA	7.64		1.92	6.31	11.46	236.1	7922.68	2926.12	9.7	188.55	55.08	152.65	7.41	74.03
33-2 TA	8.25				10.88	194.54	1237.5	1625.25	1.16	48.92	58.82	2.71	3.47	38.03
33-2 TB	4.29				10.91	65.34	1698.84	1613.04	2.23	111.03	34.32	35.22	4.41	50.58
33-2 TC	8.45				11.01	113.99	3058.9	1664.65	4.14	122.1	32.87	96.84	7.01	121.93
33-2 TD	11.68				11.18	219.79	4765.44	1495.04	11.68	124.16	25.93	128.36	8.88	118.32
33-2 TE	5.62				11.03	152.25	1871.46	1562.36	13.04	180.29	12.76	137.41	5.73	80.25
5-12 DA	6.24		4.38	4.69	10.93	105.33	1291.68	1597.44	2.31	141.52	44.99	49.92	5.18	64.33
5-12 LA	24.83		5.59	6.39	11.06	122.35	7225.53	844.22	26.07	413.92	3.73	395.29	19.37	152.95
7-1 DA	1.99				11.07	139.25	300	1197.98	7.32	117.17	2.57	25.41	2.13	12.77
7-1 LA	1.93				11.9	43.6	191.07	1186.95	0.81	40.14	1.88	22.66	3.05	12.7
7-1 LB	2.07				11.44	50.74	204.93	1324.8	1.3	38.67	5.38	10.56	0.97	7.06
7-1 TA	7.6				13.1	143.42	319.2	1444	1.82	168.95	26.45	96.82	6.38	79.8
7-1 TB	6.05				11.34	95.95	502.15	1542.75	0.54	69.88	15.12	87.3	3.27	54.87
7-1 TC	8.94				11.1	95.93	616.86	1993.62	0	93.42	14.3	75.99	6.62	33.08
E-7 BA	3.6		2.84	5.39	10.91	117.69	1026	1760.4	2.81	102.38	33.05	65.52	4.21	57.85
E-7 BB	5.57		1.71	5.47	10.92	87.02	1002.6	1665.43	81.45	122.6	36.82	85.95	4.96	52.64
E-7 BC	5		2.47	6.07	11.05	214.75	1140	1530	1.6	129.25	33.05	36.5	6	70.3
E-7 LA	5.87		2.92	6.6	11.14	216.54	5911.09	2477.14	12.44	184.96	32.11	39.27	1.82	20.19
16-1 A1	3.69	CORR			10.85	80.04	6077.43	2210.31	1.33	54.8	12.95	18.78	1.66	18.41
16-1 A2	14.02				10.96	22.98	16165.06	2397.42	1.68	80.76	31.97	71.5	12.48	61.27
16-1 A3	9.94				11.09	271.46	9969.82	2246.44	3.28	62.72	19.48	27.73	3.18	14.61
31-1 A1	4.45				11.41	122.82	8895.55	1962.45	4.72	38.72	12.33	36.67	2.54	119.97
31-1 A2	3.04				11.23	82.9	4350.24	2760.32	3.53	30.4	10.76	17.94	1.7	24.35
31-1 C1	7.61				10.97	171.23	13911.08	4253.99	1.52	52.97	22.3	19.02	2.66	29.45
31-1 C2	3.19				11.13	78.95	4568.08	258.38	1.31	10.91	17.51	12.41	1.82	27.24
32-3 A1	3.39				11.32	108.17	1342.44	3681.54	2.34	35.6	35.22	13.46	2.17	18.41
32-3 A2	4.01				11.28	110.6	1652.12	2165.4	2.05	21.65	8.78	21.61	1.6	10.71
32-3 A3	4.45				11.26	102.26	1539.7	2420.8	1.56	15.4	10.9	16.15	2.58	19.34
32-3 A4	3.8				11.25	103.97	6220.6	2861.4	1.9	36.82	14.9	8.74	1.6	11.02
37-4 D1	7.16				10.77	140.91	1303.12	1890.24	1.72	75.61	8.52	22.84	3.87	35.44
37-4 D2	23.19				11.02	368.95	3733.59	8162.88	64.24	640.74	187.84	90.21	9.97	153.98
37-4 D3	2.36				11.01	83.56	3091.6	2267.96	4.3	56.76	6.75	15.86	2.41	19.33
37-4 D4	22.28				10.8	1032.45	8845.16	3007.8	2.67	58.59	95.58	42.11	10.47	122.09
COW2 C1	9.28				11.05	220.86	900.16	2403.52	1.21	121.2	10.91	22.55	2.6	24.59
COW4 A1	5.21				11	89.87	6048.81	2203.83	1.04	41.26	12.92	28.24	2.34	36.42
COW4 A2	3.66				11.1	156.57	5948.58	3272.04	2.49	62.77	12.26	15.41	2.16	17.31
COW4 A3	2.49				10.95	86.03	3234.51	2828.64	1.52	25.85	6.6	11.85	1.37	12.08
COW4 A4	4.33				11.1	121.15	5490.44	2589.34	1.56	68.54	10.39	25.72	3.29	13.37
COW4 D1	11.84				10.99	69.38	1160.32	3031.04	1.18	120.77	36.7	98.75	7.22	29.36
COW4 D2	6.89				11.2	103.49	1378	5863.39	2.41	110.03	32.8	105.55	1.93	23.15
COW4 D3	15.18				11.21	215.25	1776.06	2419.2	1.82	219.5	57.84	115.98	11.08	69.07

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COW4 D4	7.78						10.13	152.95	972.5	2715.22	0.62	120.51	24.12	48.55	3.42	17.27	
GF116 A1	7.71						11.23	137.78	3068.58	2760.18	3.93	13.49	10.33	27.83	2.85	14.73	
GF116 A2	2.99						11.2	128.69	3286.01	2305.29	6.99	39.34	14.26	8.41	1.41	12.85	
GF116 C1	13.61						11.33	115.14	830.21	2340.92	21.64	265.56	96.09	61.52	3.27	63.01	
GF116 C2	13.48						11.24	221.21	1334.52	2655.56	1.89	266.1	23.59	44.62	7.95	58.23	
GF118 C1	6.11						11.15	79.55	6916.52	2413.45	2.99	13.67	17.41	22.91	3.48	32.38	
PDB A1	2.54						10.85	87.88	4460.24	2547.62	2.79	44.5	7.26	7.21	1.37	9.02	
PDB A2	3.41						11.25	79.66	4238.63	2537.04	1.26	30.55	8.38	8.56	0.99	10.74	
PDB C1	2.82						11.33	166.75	3307.86	4230	2.76	10.97	119.91	10.24	2.62	20.13	
PDB D1	6.58						10.85	273.44	552.72	3303.16	1.71	91.4	4.97	58.36	3.55	24.81	
PDB D2	8.29						11.05	126.17	8.29	9442.31	0.83	96.66	17.74	72.37	4.15	11.11	
PDB D3	4.11						11	106.98	793.23	3497.61	2.14	76.4	10.77	31.93	2.59	17.26	
PDB D4	3.1						11.05	53.72	700.6	3658	0.34	46.03	7.38	5.18	1.36	69.07	
PHIL1 A1	4.58						11.22	68.79	6311.24	2748	1.01	34.35	11.63	21.89	2.2	64.17	
PHIL1 A2	11.82						11.27	147.63	16370.7	2174.88	0.24	141.37	32.51	84.04	6.5	147.51	
PHIL1 A3	3.51						11.29	179.96	4780.62	2446.47	1.23	34.71	8.25	10.32	1.68	88	
PHIL1 A4	4.32						11.3	55.38	4497.12	2191.56	0.65	42.12	4.45	18.71	1.73	116.73	
33-2 01	10.03	CONT	0	2.06	7.05	10.78	263.59	18966.73	2377.11	3.11	77.33	13.24	70.11	3.71	48.95	0	
33-2 02	5.3		-0.2	2.29	6.93	11.12	300.77	3874.3	3084.6	6.51	43.51	9.59	18.6	2.33	17.17	-0.2	
33-2 03	5.98		-0.27	2.72	6.29	11.65	423.03	16761.94	4287.66	18.06	146.69	53.1	34.98	3.11	13.04	-0.27	
33-2 04	4.32		-0.3	0.64	4.68	11.11	81.39	280.8	1749.6	1.47	33.74	12.1	9.98	1.43	49.25	-0.3	
33-2 05	3.42		-0.6	2.02	5.93	11.5	192.96	314.64	2014.38	7.52	31.43	16.48	15.9	3.32	36.05	-0.6	
33-2 06	4.24		-0.8	1.77	5.5	11.53	155.74	864.96	4216.8	1.48	235.49	30.4	32.1	5.6	74.79	-0.8	
33-2 07	5.28		-0.9	0.93	5.35	10.93	183.22	5280	2412.96	2.9	59.93	46.99	15	3.01	23.55	-0.9	
33-2 08	6.71		-0.95	2.55	6.33	11.06	386.16	8300.27	1362.13	14.23	194.46	14.76	52.2	2.28	31.2	-0.95	
33-2 09	4.08		-1	1.97	5.91	11.37	396.53	3288.48	877.2	4	38.84	26.81	22.64	2.9	74.26	-1	
33-2 10	5.31		-1.1	2.09	6.09	11.22	265.34	5405.58	1874.43	6.32	122.71	18.05	30.85	1.81	13.43	-1.1	
33-2 11	5.47		-1.2	1.27	5.13	11.23	63.07	492.3	1564.42	0.49	48.19	28.66	22.65	1.53	14.93	-1.2	
33-2 12	4.71		-1.3	0.17	3.77	11.09	64.95	456.87	1234.02	0	55.53	0	16.53	0.75	17.76	-1.3	
33-2 12A	6.35		-1.35	1.44	3.32	11.08	370.82	6038.85	2552.7	1.97	61.4	12.06	18.35	2.35	24.64	-1.35	
33-2 13	6.94		-1.4	2.74	6.28	11.79	495.93	12068.66	7564.6	29.77	100.56	50.01	38.24	4.44	33.73	-1.4	
33-2 14	5.2		-1.45	1.95	6.51	11.04	268.58	9625.2	2043.6	3.74	48.67	15.76	104.36	2.6	17.21	-1.45	
33-2 15	7.61		-1.5	1.3	7.03	11.3	177.31	707.73	2427.59	1.98	68.33	24.12	58.75	3.96	29.91	-1.5	
33-2 16	5.94		-1.55	1.01	6.94	11.01	128.6	11975.04	2162.16	1.66	53.04	21.03	31.24	3.05	27.38	-1.55	
33-2 17	6.07		-1.6	1.45	6.84	11.48	166.01	1984.89	2725.43	2.12	267.93	12.5	46.74	3.28	28.83	-1.6	
33-2 18	6.63		-1.65	2.03	7.44	11.22	314.2	2632.11	1882.92	6.17	131.21	18.56	37.74	2.12	35.8	-1.65	
33-2 19	5.96		-1.7	0.43	6.45	10.73	371.19	3766.72	2938.28	4.05	59.84	23.84	22.83	3.69	25.27	-1.7	
33-2 20	2.98		-1.75	1.72	7.15	11.18	176.33	3045.56	4470	4.41	76.31	37.97	52.12	9.27	164.41	-1.75	
33-2 21	7.61		-1.8	1.44	6.9	11.07	206.99	3325.57	1346.97	1.29	140.79	0	15.14	0.61	13.85	-1.8	
33-2 22	5.05		-1.85	1.48	6.85	11.27	138.22	2580.55	2519.95	2.98	43.94	18.58	12.12	1.92	18.03	-1.85	
33-2 23	6.12		-1.9	2.09	7.18	11.1	220.2	1891.08	1658.52	5.08	117.93	9.79	24.3	1.71	24.79	-1.9	
33-2 24	5.68		-1.95	1.9	7.07	11.15	171.59	2766.16	1465.44	1.53	129.11	6.82	12.27	1.14	16.59	-1.95	
33-2 25	5.86		-2	1.85	7.05	11.12	316.2	4793.48	2303.98	12.36	138.3	32.7	108.35	4.45	137.65	-2	
33-2 26	5.34		-2.05	2.12	7.04	11.25	271.33	4474.92	2269.5	8.7	102.1	13.88	10.73	1.34	36.9	-2.05	
33-2 27	6.07		-2.1	1.7	6.93	11.02	199.58	3368.85	1468.94	2.61	131.48	9.71	19.85	1.58	30.11	-2.1	
33-2 28	7.71		-2.15	1.34	6.69	11.02	280.34	4602.87	2097.12	4.16	154.35	1.54	38.65	0.39	40.71	-2.15	
33-2 29	4.21		-2.2	1.07	6.56	11.7	183.72	3780.58	2888.06	14.06	147.18	19.2	30.78	3.11	16.88	-2.2	
33-2 30	8.88		-2.25	4.23	8.95	11.04	205.66	2743.92	2006.88	4.62	113.22	0	45.73	0.88	25.66	-2.25	