Supplemental Table 1. Summary of the Overlaps in the Differentially Regulated Genes by the *cop1* Mutations and the White Light

Section A: Number and percentage of white light up-regulated genes overlap with those regulated by the three *cop1* mutations in darkness. In this study, 1065 genes were up-regulated in white light in wild type with 2-fold cut-off. Three cut-off values for the differential expression ratios were used for the *cop1* mutations.

| (i) Similarly regulated (percentage in bracket) | | | | |
|---|---------------|-------------|-------------|--------------|
| | <u>cop1-1</u> | cop1-4 | copl-6 | <u>N-282</u> |
| With 2-fold cut-off | 566 (53%) | 611 (57.4%) | 557 (52.3%) | 512 (48.1%) |
| With 1.5-fold cut-off | 807 (75.8%) | 819 (76.9%) | 773 (72.6%) | 758 (71.2%) |
| With 1.2-fold cut-off | 936 (87.9%) | 938 (88.1%) | 915 (85.9%) | 906 (85.1%) |

| (ii) Oppositely regulated (percentage in bracket) | | | | |
|---|---------------|------------|------------|------------|
| | <u>copl-l</u> | cop1-4 | copl-6 | N-282 |
| With 2-fold cut-off | 8 (0.75%) | 10 (0.94%) | 4 (0.38%) | 4 (0.38%) |
| With 1.5-fold cut-off | 19 (1.78%) | 17 (1.60%) | 16 (1.50%) | 12 (1.13%) |
| With 1.2-fold cut-off | 31 (2.9%) | 39 (3.7%) | 32 (3.00%) | 31 (2.9%) |

Section B: Number and percentage of white light down-regulated genes overlap with those regulated by the three *cop1* mutations in darkness. In this study, 774 genes were up-regulated in white light in wild type with 2-fold cut-off. Three cut-off values for the differential expression ratios were used for the *cop1* mutations.

(iii) Similarly regulated (percentage in bracket)

| | cop1-1 | cop1-4 | copl-6 | N-282 |
|-----------------------|--------------|-------------|-------------|-------------|
| With 2-fold cut-off | 432 ((55.8%) | 368 (47.5%) | 278 (40%) | 268 (34.6%) |
| With 1.5-fold cut-off | 603 (77.9%) | 633 (81.8%) | 515 (66.5%) | 505 (65.2%) |
| With 1.2-fold cut-off | 691 (89.3%) | 720 (93.0%) | 654 (84.5) | 644 (83.2%) |

(iv) Oppositely regulated (percentage in bracket)

| () | cop1-1 | cop1-4 | | <u>N-282</u> |
|-----------------------|------------|------------|------------|--------------|
| With 2-fold cut-off | 10 (1.29%) | 2 (0.26%) | 14 (1.81%) | 7 (0.90%) |
| With 1.5-fold cut-off | 18 (2.33%) | 6 (0.78%) | 25 (3.33%) | 17 (2.20%) |
| With 1.2-fold cut-off | 28 (3.61%) | 16 (1.29%) | 44 (5.68%) | 29 (3.75%) |

Section C: Number and percentage of white light total (up and down) regulated genes overlap with those regulated by the three *cop1* mutations in darkness. Together, 1839 genes were up or down-regulated in white light with 2-fold cut-off. Three cut-off values for the differential expression ratios were used for the *cop1* mutations.

| (v) Similarly regu | Similarly regulated (percentage in bracket) | | | | |
|-----------------------|---|--------------|--------------|--------------|--|
| | copl-6 | N-282 | | | |
| With 2-fold cut-off | 998 (54.3%) | 979 (53.2%) | 835 (45.4%) | 780 (42.4%) | |
| With 1.5-fold cut-off | 1410 (76.7%) | 1452 (79.0%) | 1288 (70.0%) | 1263 (68.7%) | |

With 1.2-fold cut-off 1627 (88.5%) 1658 (90.2%) 1569 (85.3%) 1550 (84.3%)

| (vi) Oppositely regulated (percentage | e in bracket) |
|---------------------------------------|---------------|
|---------------------------------------|---------------|

| | <u>cop1-1</u> | cop1-4 | <i>cop1-6</i> | N-282 |
|---------------------|---------------|------------|---------------|------------|
| With 2-fold cut-off | 18 (0.98%) | 12 (0.65%) | 18 (0.98%) | 11 (0.60%) |
| With 2-fold cut-off | 37 (2.0%) | 23 (1.25%) | 41 (2.23%) | 29 (1.58%) |
| With 2-fold cut-off | 59 (3.21%) | 55 (2.99%) | 76 (4.13%) | 60 (3.26%) |

Section D: Number and percentage of differentially regulated genes that overlap among the three cop1 mutations in darkness. The 2-fold cut-off was used to select those genes regulated by each of the three cop1 mutations.

(vii) Similarly regulated (percentage in bracket)

| | <u>cop1-1</u> | cop1-4 | copl-6 | <u>N-282</u> |
|--------|---------------|-------------|-------------|--------------|
| cop1-1 | 1321 (100%) | | | |
| cop1-4 | 1176 (70.7%) | 1176 (100%) | | |
| cop1-6 | 1162 (62.5%) | 709 (61.0%) | 1162 (100%) | |
| N282 | 673 (69.1%) | 639 (65.7%) | 666 (85.3%) | 973 (68.4%) |

(viii) Oppositely regulated (percentage in bracket)

| | cop1-1 | cop1-4 | cop1-6 | N-282 |
|--------|------------|------------|-----------|--------|
| cop1-1 | 0 (0%) | | | |
| cop1-4 | 8 (0.68%) | 0 (0%) | | |
| cop1-6 | 13 (1.1%) | 13 (1.1%) | 0 (0%) | |
| N282 | 13 (1.34%) | 18 (1.85%) | 2 (0.21%) | 0 (0%) |

Section E: Number and percentage of differentially regulated genes by the three different qualities and white light in wild type. The 2-fold cut-off was used to select those genes regulated by each of the light quality.

| (ix) | Similarly regulated (percentage in bracket) | | | | |
|------|---|-----------------|-----------------|-------------|----------------|
| | | White light | Blue light | Red light | Fare red light |
| | White light | 1839 (100%) | - | - | - |
| | Blue light | 713 (78.2%) | 911 (100%) | | |
| | Red light | 452 (67.4%) | 467 (69.6%) | 671 (100%) | |
| | Far red light | 793 (73.3%) | 498 (46.0%) | 335 (31.0%) | 1082 (100%) |
| | | | | | |
| (x) | Oppositely re | gulated (percen | tage in bracket | | |
| | | White light | Blue light | Red light | Fare red light |
| | White light | 0 (0%) | | | |
| | Blue light | 23 (2.52%) | 0 (0%) | | |
| | Red light | 26 (3.87%) | 16 (3.38%) | 0 (0%) | |
| | Far red light | 20 (1.85%) | 15 (1.39%) | 16 (1.48%) | 0 (0%) |
| | | | | | |