| GO | P-Value | Description |
| --- | --- | --- |
| "GO:0070069" | 5.66E-004 | "cytochrome complex"  |
| "GO:0005746" | 2.12E-003 | "mitochondrial respiratory chain"  |
| "GO:0005750" | 6.88E-003 | "mitochondrial respiratory chain complex III"  |
| "GO:0045275" | 6.88E-003 | "respiratory chain complex III"  |
| "GO:1990204" | 9.58E-003 | "oxidoreductase complex"  |
| "GO:0016021" | 1.02E-002 | "integral component of membrane"  |
| "GO:0031224" | 1.02E-002 | "intrinsic component of membrane"  |
| "GO:1902495" | 1.95E-002 | "transmembrane transporter complex"  |
| "GO:0005874" | 2.08E-002 | "microtubule"  |
| "GO:0044425" | 2.45E-002 | "membrane part"  |
| "GO:0015630" | 2.96E-002 | "microtubule cytoskeleton"  |
| "GO:0044455" | 3.09E-002 | "mitochondrial membrane part"  |
| "GO:0009277" | 3.69E-002 | "fungal-type cell wall"  |
| "GO:0031300" | 4.66E-002 | "intrinsic component of organelle membrane"  |
| "GO:0031301" | 4.66E-002 | "integral component of organelle membrane"  |
| "GO:0016020" | 4.81E-002 | "membrane"  |

Supplementary Table 12: List of overrepresented GO terms in the Cellular Components category for the genes upregulated at 1C1W