Table S6: Corrected number of proteins annotated with different targeting peptides for the A. thaliana genome using the confusion matrix from Table [S3](supplementary/index.html#table:cm_plant).

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| |  |  |  |  |  | | --- | --- | --- | --- | --- | |  | Class | Original | Corrected | Shift | |  | SP | 4115 | 4109.3 | -0.1% | |  | mTP | 1095 | 1106.5 | +1.1% | |  | cTP | 1448 | 1410.6 | -2.6% | |  | luTP | 127 | 158.6 | +24.9% | |