Table S3: Confusion matrix for Viridiplantae representing the number of proteins for each targeting peptides predicted by TargetP 2.0 (rows) versus observed in the test set (columns).

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| |  |  |  |  |  |  |  | | --- | --- | --- | --- | --- | --- | --- | |  | Class | SP | mTP | cTP | luTP | noTP | |  | SP | 272 | 0 | 2 | 0 | 3 | |  | mTP | 0 | 117 | 2 | 0 | 11 | |  | cTP | 0 | 2 | 197 | 9 | 9 | |  | luTP | 0 | 0 | 11 | 34 | 0 | |  | noTP | 10 | 6 | 15 | 2 | 1779 | |