**Supplementary file 2. Annotation of proteins used in this study**

| **FungiDB ID** | **Systematic/Protein name** | **Trafficking process** |
| --- | --- | --- |
| AN0411 | SarASar1 | ER exit sites (ERes) |
| AN3720 | Sec24 | ER exit sites (ERes) |
| AN4317 | Sec13 | ER exit sites (ERes) |
| AN6257 | Sec31 | ER exit sites (ERes) |
| AN11127 | Sec12 | ER exit sites (ERes) |
| AN3026 | CopACopI | COPI coat |
| AN1126 | ArfAArf1/2 | Early/late Golgi sorting |
| AN0112 | GeaAGea1/2 | Early Golgi sorting |
| AN6709 | HypBSec7 | Late Golgi sorting |
| AN4281 | RabOYpt1/Rab1 | Early Golgi sorting |
| AN0347 | RabEYpt31/32/Rab11 | Post-Golgi |
| AN7682 | Ap1σAps1 | Late Golgi sorting |
| AN6974 | RabDSec4 | exocyst |
| AN8488 | Ykt6 | R-SNARE |
| AN8769 | SynASnc1/2 | R-SNARE |
| AN0571 | NyvANyv1 | R-SNARE |
| ChrI\_A\_nidulans\_FGSC\_A4 2,596,682-2,597,694 | Sec22 | R-SNARE |
| AN3416 | SsoASso1/2 | Qa-SNARE |
| AN9526 | SedVSed5 | Qa-SNARE |
| AN2419 | Sec9 | Qb-SNARE |
| AN11900 | Bos1 | Qb-SNARE |
| AN10508 | Sft1 | Qc-SNARE |
| AN2523 | ChsBChs3 | apical cargo |
| AN6932 | UapA | non-polar cargo |