

| SARS-CoV-2 viral proteins significant between variants at 24 hpi | | | | | | | |
|--|-------------|--------------|----------------|------------------|--------------|----------------|------------------|
| Protein id | Gene symbol | Significance | | | FDR q-value | | |
| | | Delta vs B.1 | Omicron vs B.1 | Omicron vs Delta | Delta vs B.1 | Omicron vs B.1 | Omicron vs Delta |
| P0DTC7 | 7a | ns | ns | ns | 0,43059 | 0,465696 | 0,0727006 |
| P0DTD2 | 9b | + | ns | + | 0 | 0,253294 | 0,0136 |
| P0DTC2 | S | ns | ns | + | 0,0561951 | 0,146333 | 0,0118261 |
| P0DTC3 | 3a | ns | ns | + | 0,881793 | 0,167333 | 0,02125 |
| P0DTC6 | NSP6 | ns | ns | + | 0,747202 | 0,159636 | 0,0126512 |
| P0DTC5 | M | ns | ns | + | 0,803253 | 0,143429 | 0,0437667 |
| P0DTC9 | N | ns | ns | + | 0,0729362 | 0,112 | 0,0114805 |
| P0DTC8 | NSP8 | + | ns | + | 0,0472727 | 0,125778 | 0,0188889 |
| P0DTD1 | REP | ns | ns | + | 0,440881 | 0,219429 | 0,0100741 |

Viral proteins detected in whole cell proteome of cells infected with B.1, Delta (B.1.167.2) or Omicron BA.1 at 24 hpi

No significance was observed at 6 and 12 hpi and significance determined as described in Proteomics methods.

Significant values were FDR corrected at q-value < 0.05 was considered significant (+). ns: not significant.

N=3 biologically independent replicates