**Table 6:** $RF$ and $RF'$ distance between each pair of trees for protein sequence data partitions. Unshaded cells show $RF$ distances, and shaded cells show $RF'$ distances.

|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | **A2M** | **ACE** | **APBB2** | **APP** | **BLMH** | **MPO** | **NOS3** | **PLAU** | **SORL1** |
| **A2M** | -- | 0.492 | 0.438 | 0.421 | 0.472 | 0.634 | 0.553 | 0.745 | 0.590 |
| **ACE** | 0.709 | -- | 0.401 | 0.402 | 0.423 | 0.456 | 0.449 | 0.463 | 0.424 |
| **APBB2** | 0.826 | 0.915 | -- | 0.435 | 0.400 | 0.418 | 0.426 | 0.443 | 0.436 |
| **APP** | 0.864 | 0.911 | 0.833 | -- | 0.384 | 0.402 | 0.408 | 0.386 | 0.402 |
| **BLMH** | 0.750 | 0.860 | 0.917 | 0.957 | -- | 0.454 | 0.465 | 0.461 | 0.454 |
| **MPO** | 0.455 | 0.786 | 0.872 | 0.911 | 0.789 | -- | 0.468 | 0.562 | 0.526 |
| **NOS3** | 0.592 | 0.800 | 0.854 | 0.897 | 0.765 | 0.760 | -- | 0.499 | 0.487 |
| **PLAU** | 0.294 | 0.769 | 0.814 | 0.951 | 0.774 | 0.577 | 0.696 | -- | 0.584 |
| **SORL1** | 0.527 | 0.857 | 0.830 | 0.911 | 0.789 | 0.643 | 0.720 | 0.538 | -- |