**Table 5:** $RF$ and $RF'$ distance between each pair of trees for nucleotide sequence data partitions. Unshaded cells show $RF$ distances, and shaded cells show $RF'$ distances.

|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | **A2M** | **ACE** | **APBB2** | **APP** | **BLMH** | **MPO** | **NOS3** | **PLAU** | **SORL1** |
| **A2M** | -- | 0.379 | 0.405 | 0.417 | 0.417 | 0.444 | 0.472 | 0.430 | 0.404 |
| **ACE** | 0.969 | -- | 0.380 | 0.430 | 0.379 | 0.379 | 0.379 | 0.391 | 0.391 |
| **APBB2** | 0.905 | 0.968 | -- | 0.380 | 0.392 | 0.405 | 0.380 | 0.392 | 0.405 |
| **APP** | 0.875 | 0.844 | 0.968 | -- | 0.391 | 0.379 | 0.417 | 0.379 | 0.430 |
| **BLMH** | 0.875 | 0.969 | 0.937 | 0.938 | -- | 0.391 | 0.430 | 0.404 | 0.391 |
| **MPO** | 0.812 | 0.969 | 0.905 | 0.969 | 0.938 | -- | 0.404 | 0.444 | 0.404 |
| **NOS3** | 0.750 | 0.969 | 0.968 | 0.875 | 0.844 | 0.906 | -- | 0.417 | 0.417 |
| **PLAU** | 0.844 | 0.938 | 0.937 | 0.969 | 0.906 | 0.812 | 0.875 | -- | 0.404 |
| **SORL1** | 0.906 | 0.938 | 0.905 | 0.844 | 0.938 | 0.906 | 0.875 | 0.906 | -- |