**S6 Table. The COI pairwise genetic distances calculated in RaxML for four *Calyptorhynchus* sp. sequences.** (Downloaded from GenBank: JF414241.1, JN801424.1, JF414242.1, JF414279.1, JF414274.1).

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | *C. lathami* | *C. banksii* | *C. baudinii* | *C. funereus* | *C. latirostris* |
| *C. lathami* | \* |  |  |  |  |
| *C. banksii* | 0.151 | \* |  |  |  |
| *C. baudinii* | 0.632 | 0.499 | \* |  |  |
| *C. funereus* | 0.686 | 0.627 | 0.036 | \* |  |
| *C. latirostris* | 0.594 | 0.497 | 0.009 | 0.03 | \* |