**Supplementary Figure Legends**

**Fig. S1**: Best tree for the family *Peltulaceae* based on the nucSSU data set calculated with raxml, using a GTR+I+G model (likelihood value = -6,176.14). See caption of Figure 4 for details. Long branches are drawn by their half-length for saving space (//).

**Fig. S2**: Best tree for the family *Peltulaceae* based on the nucSSU data set calculated with raxml, using a GTR+I+G model (likelihood value = -7,912.07). See caption of Figure 4 for details.

**Fig. S3**: Best tree for the family *Peltulaceae* based on the mtSSU data set calculated with raxml, using a GTR+I+G model (likelihood value = -3,057.50). See caption of Figure 4 for details. Long branches are drawn by their half-length for saving space (//).

**Fig. S4**: Best tree for the family *Peltulaceae* based on the ITS data set calculated with raxml, using a GTR+I+G model (likelihood value = -2,486.78). See caption of Figure 4 for details.

**Fig. S5**: Best tree for the family *Peltulaceae* based on the *RPB2* data set calculated with raxml, using a GTR+I+G model (likelihood value = -10,049.06). See caption of Figure 4 for details.

**Fig. S6**: Best tree for the family *Peltulaceae* based on the β-tub data set calculated with raxml, using a GTR+I+G model (likelihood value = -6,822.62). See caption of Figure 4 for details.