

Supplementary information, Fig. S2 Mass spectrometry analysis of lipids isolated from MIaFEDCB in DDM and in nanodiscs. a, Mass spectra of lipids purified from buffer alone. b, Mass spectra of lipids isolated from purified MIaFEDCB in DDM, the ions at m/z 733.5, 747.5, 761.5, 773.5, 787.5 were identified as PG. The ions at m/z 702.5, 716.5 and 742.5 were identified as PE. c, Mass spectra of lipids purified from MIaFEDCB in POPG nanodisc. d, Mass spectra of lipids purified from MIaFEDCB in POPG nanodisc at zoomed range from m/z 680 to 800, e. Tandem MS spectra of the fragments at m/z 733.5 from purified MIaFEDCB in DDM. The fragment at m/z 733.5 is consistent with PG with two acyl chain of C16:0 and C17:1. f Tandem MS spectra of the fragments at m/z 747.5 from purified MIaFEDCB in DDM. The fragment at m/z 747.5 is consistent with PG with two acyl chain of C16:0 and C18:1. g, Tandem MS spectra of the fragments at m/z 773.5 from purified MIaFEDCB in DDM. The fragment at m/z 773.5 is consistent with PG with two acyl chain of C18:1 and C18:1. h. Tandem MS spectra of the fragments at m/z 702.5 from purified MIaFEDCB in DDM. The fragment at m/z 702.5 is consistent with PE with two acyl chain of C16:0 and C17:1. i, Tandem MS spectra of the fragments of YAGALVLGQYYK from MIaC in MIaFEDCB in nanodisc. **j**, Tandem MS spectra of the fragments of NSQTGNWQAYDMIAEGVSMITTK from MlaC in MIaFEDCB in nanodisc. k, Tandem MS spectra of the fragments of LMMVALLVIAPLSAATAADQTNPYK from MIaC in MIaFEDCB in nanodisc. I, Tandem MS spectra of the fragments of TIVDQELLPYVQVK from MIaC in MIaFEDCB in nanodisc.