

Supporting Information

Factors influencing the detergent-free membrane protein isolation using synthetic nanodisc-forming polymers

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^aequal contribution

Amino acid sequences

Rat flavin mononucleotide binding domain (FBD) of cytochrome P450 reductase

MGDSHEDTSATMPEAVAEEVSLFSTTDMVLFSLIVGVLTYWFIFRKKKEEIPFESKIQTAPPV
KESFVEKMKKTGRNIIVFYGSQTGTAEFFANRLSKDAHRYGMRGMSADPEEYDLADLSSLPEI
DKSLVVFCMATYGECDPTDNAQDFYDWLQETDVDLTGVKFAVFGLGNKTYEHFNAMGKYVDQRL
EQLGAQRIFELGLGDDDDGNLEEDFITWREQFWPAVCEFFGVEATGEEHHHHHH

Rabbit cytochrome-b5

GHHHHHAAQSDKDVKYITLEEIKKHNSKSTWLIILHHKVYDLTKFLEEHPGGEEVLREQAGD
ATENFEDVGHSTDARELSKTFIIGELHPDDRSKLSKPMETLITTVDSSNSSWWTNWVIPAISALI
VALMYRLYMADD

Transmembrane domains are underlined.

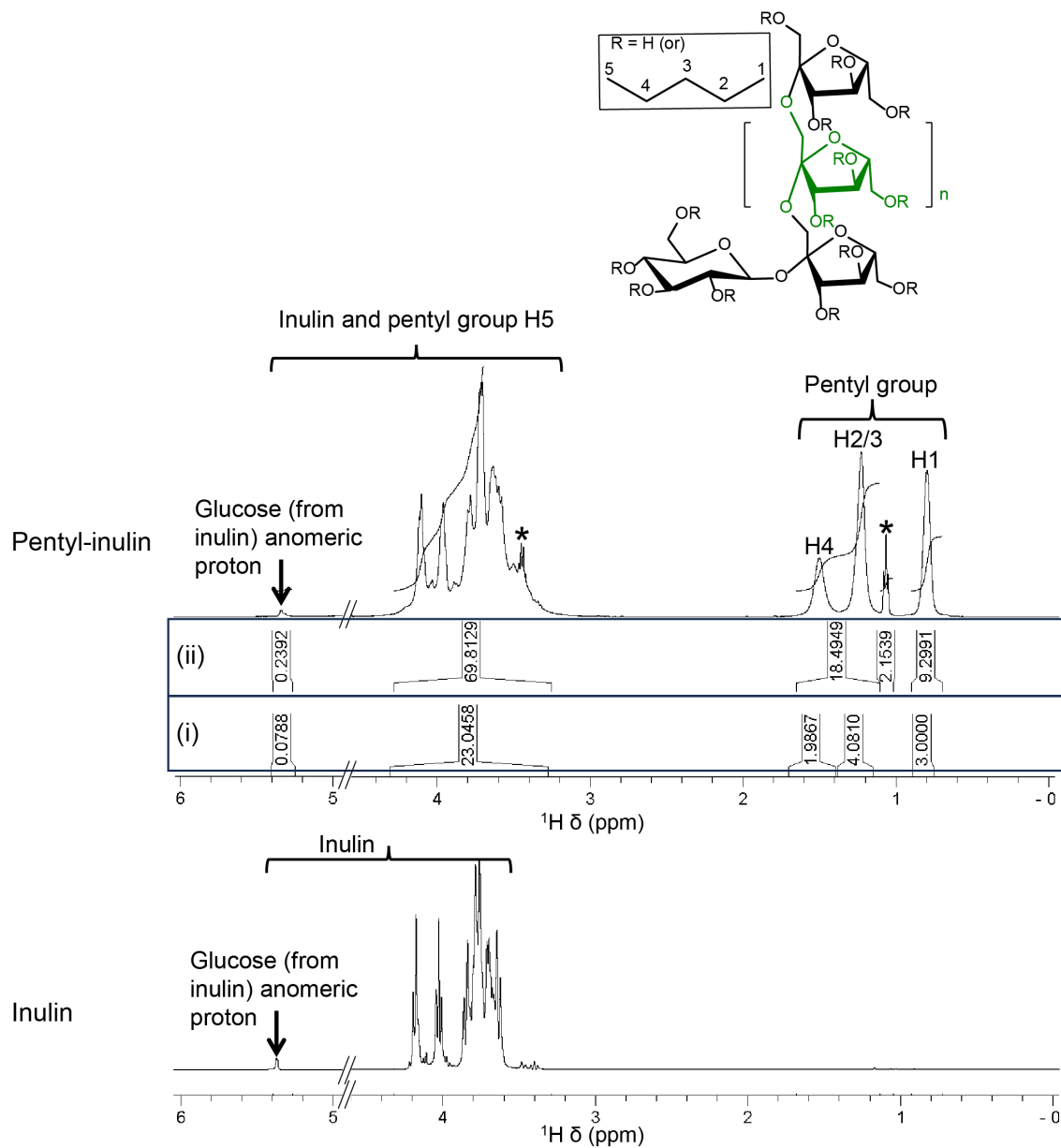


Figure S1A. ^1H NMR analysis of inulin and pentyl-inulin. The peaks are labelled with assignments. The degree of functionalization and the percentage of impurities (indicated with ‘*’) are measured using peak intensities in NMR spectra. (box-i) integral values to calculate the degree of functionalization (~ 0.33 ; see methods of the equation). (box-ii) integral values to estimate the percentage of impurities (possibly diethyl ether) in the pentyl-inulin sample. The chemical structure of pentyl-inulin is shown at the top. The water proton peak is omitted from the NMR spectra for clarity.

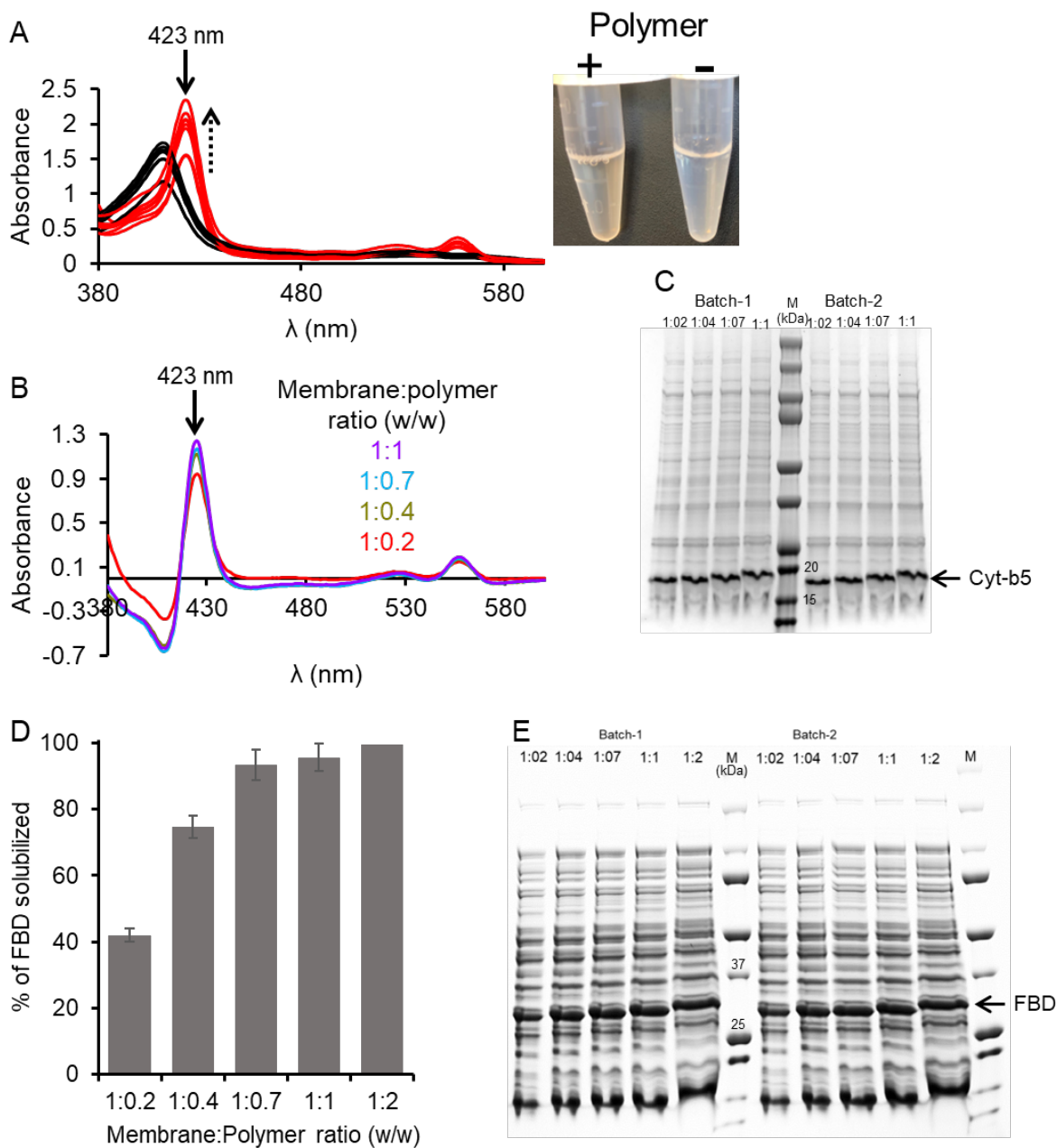


Figure S1B. (A) Absorbance spectra of oxidized (black) and sodium dithionite reduced Cyt-b5 in *E. coli* lipid-pentyl-inulin nanodiscs. The samples were made using 4 different membrane:polymer (w/w) ratios as indicated. The absorbance peak intensity change at 423 nm is indicated with an upside dotted arrow. Picture: The light-red colored (left-side) and colorless (right-side) solutions in tubes are supernatants after centrifugation of Cyt-b5-enriched cell membranes with and without polymer, respectively. (B) Difference absorbance spectra (reduced minus oxidized) of pentyl-

inulin-solubilized *E. coli* membranes enriched with a ~15.7-kDa rabbit cytochrome-b5 showing the maximal absorbance differences at 409, 423, 526 and 556 nm. The data were collected on the solubilization samples prepared using 4 different polymer concentrations as indicated. (C) SDS-PAGE analysis of pentyl-inulin-solubilized Cyt-b5-rich *E. coli* cell membranes. (D) Bar plot depicting the percentage of 27.8-kDa FBD protein band intensities at 5 different membrane:polymer ratios (w/w) as indicated. (E) SDS-PAGE analysis of pentyl-inulin-solubilized FBD-rich *E. coli* cell membranes. Both Cyt-b5 and FBD are anchored to the lipid membrane via a transmembrane helical domain. The solubilization experiments were performed in technical replicates (Batch 1 and 2). Cyt-b5 and FBD proteins have been directly isolated using SMA-EA and pentyl-inulin polymers, respectively [1-4].

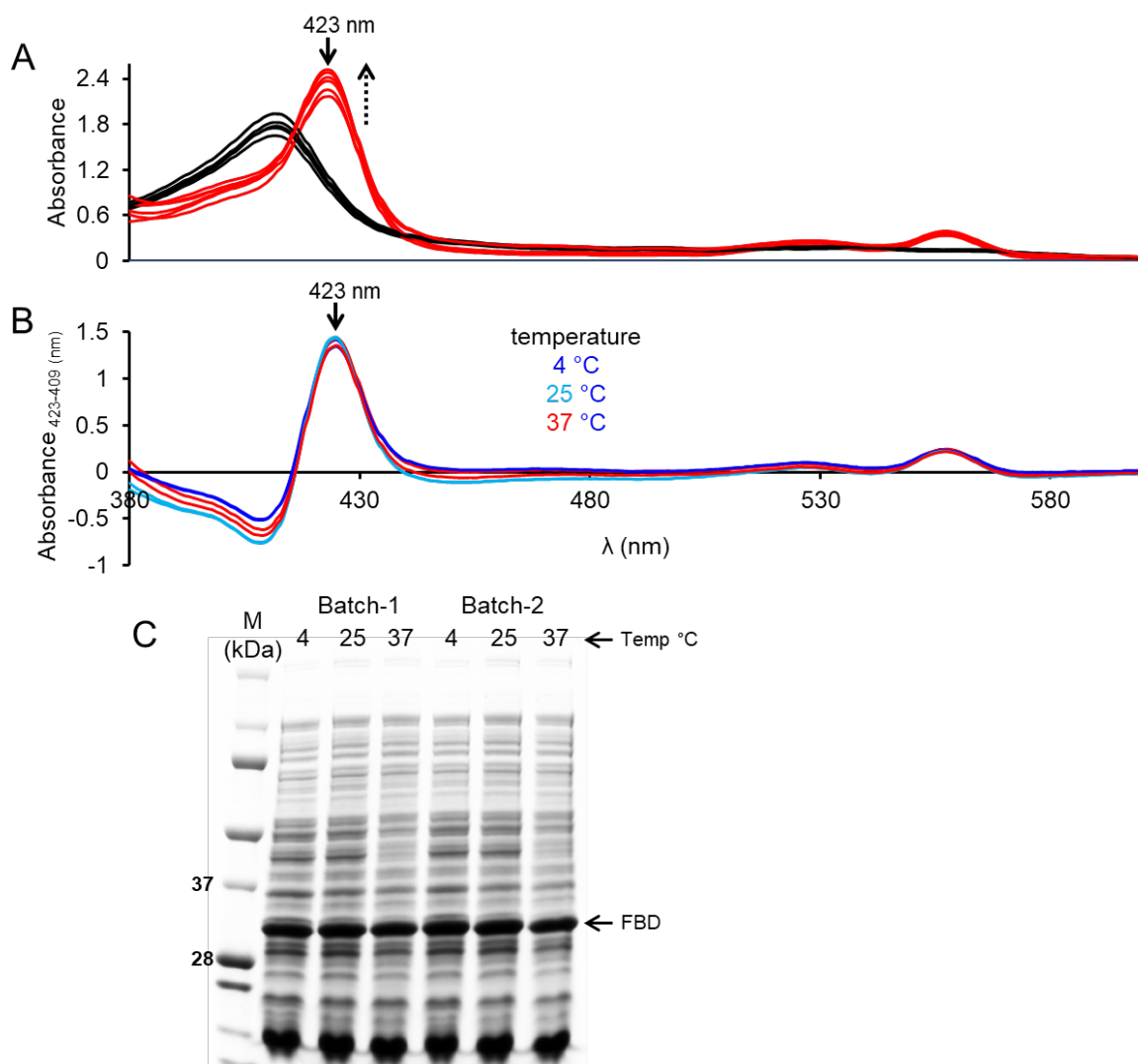


Figure S2. (A) Absorbance spectra of oxidized (black) and sodium dithionite reduced Cyt-b5 in *E. coli* lipid-pentyl-inulin nanodiscs. The samples were made using 1:1 membrane:polymer (w/w) ratio and solubilization was performed at 3 different temperatures as indicated. The absorbance peak intensity change at 423 nm is indicated with upside dotted arrow. (B) Difference absorbance spectra (reduced minus oxidized) of pentyl-inulin-solubilized *E. coli* membranes enriched with a ~15.7-kDa rabbit cytochrome-b5 showing the maximal absorbance differences at 409 and 423 nm. The data were collected on the solubilization samples prepared at 3 different temperature conditions as indicated. (C) SDS-PAGE analysis of pentyl-inulin-solubilized 27.8-kDa FBD-rich *E. coli* cell membranes obtained at different temperature conditions. M denotes the protein marker. The solubilization experiments were performed in technical replicates (Batch 1 and 2).

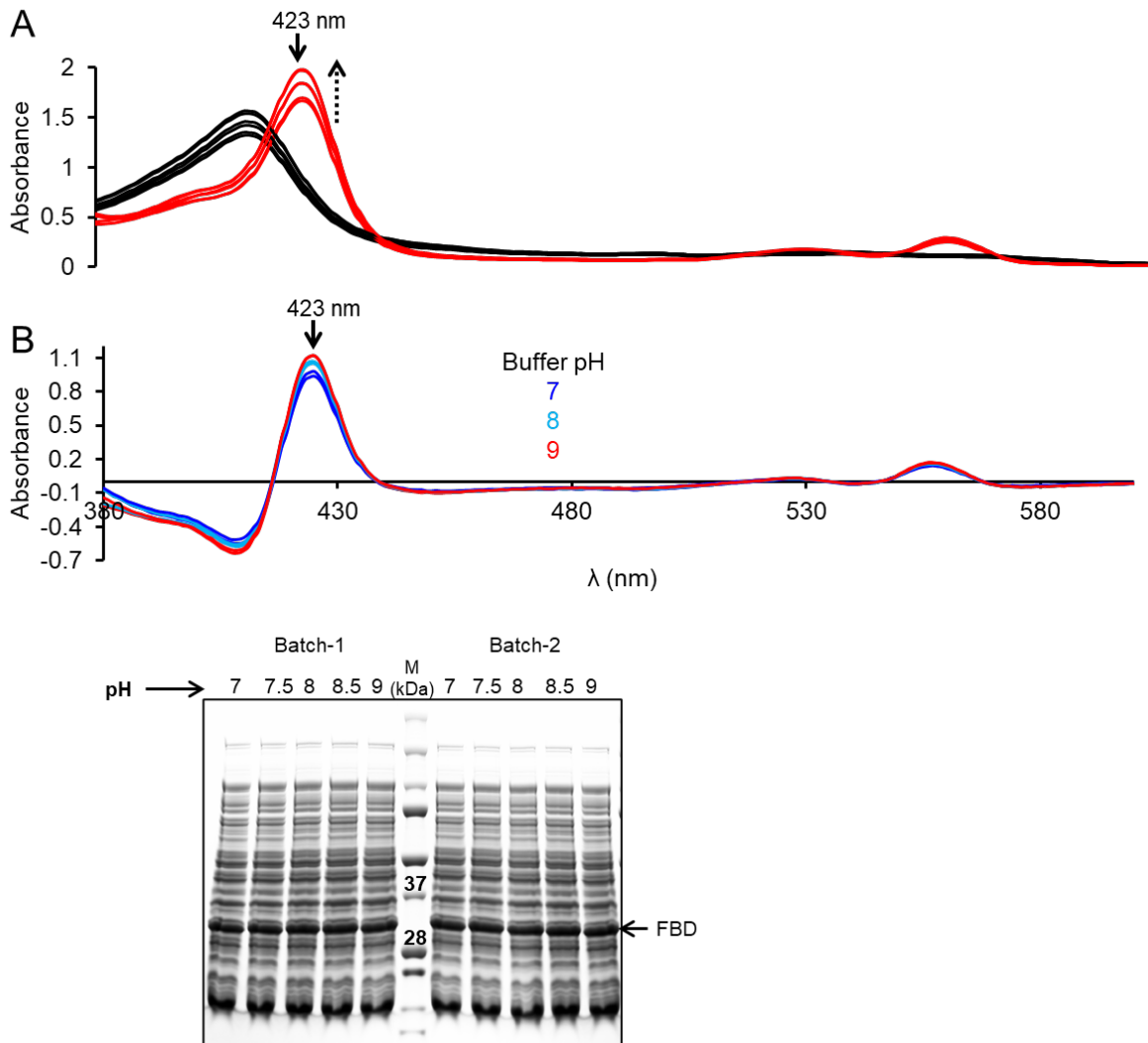


Figure S3. (A) Absorbance spectra of oxidized (black) and sodium dithionite reduced Cyt-b5 in *E. coli* lipid-pentyl-inulin nanodiscs. The samples were made using 1:1 membrane:polymer (w/w) ratio and solubilization was performed at 3 different pH conditions as indicated. The absorbance peak intensity change at 423 nm is indicated with an upside dotted arrow. (B) Difference absorbance spectra (reduced minus oxidized) of pentyl-inulin-solubilized *E. coli* membranes enriched with a ~15.7-kDa rabbit cytochrome-b5 showing the maximal absorbance differences at 409 and 423 nm. The data were collected on the solubilization samples prepared at 3 different pH conditions as indicated. (C) SDS-PAGE analysis of pentyl-inulin-solubilized 27.8-kDa FBD-rich *E. coli* cell membranes at different pH conditions. M denoted the protein marker. The solubilization experiments were performed in technical replicates.

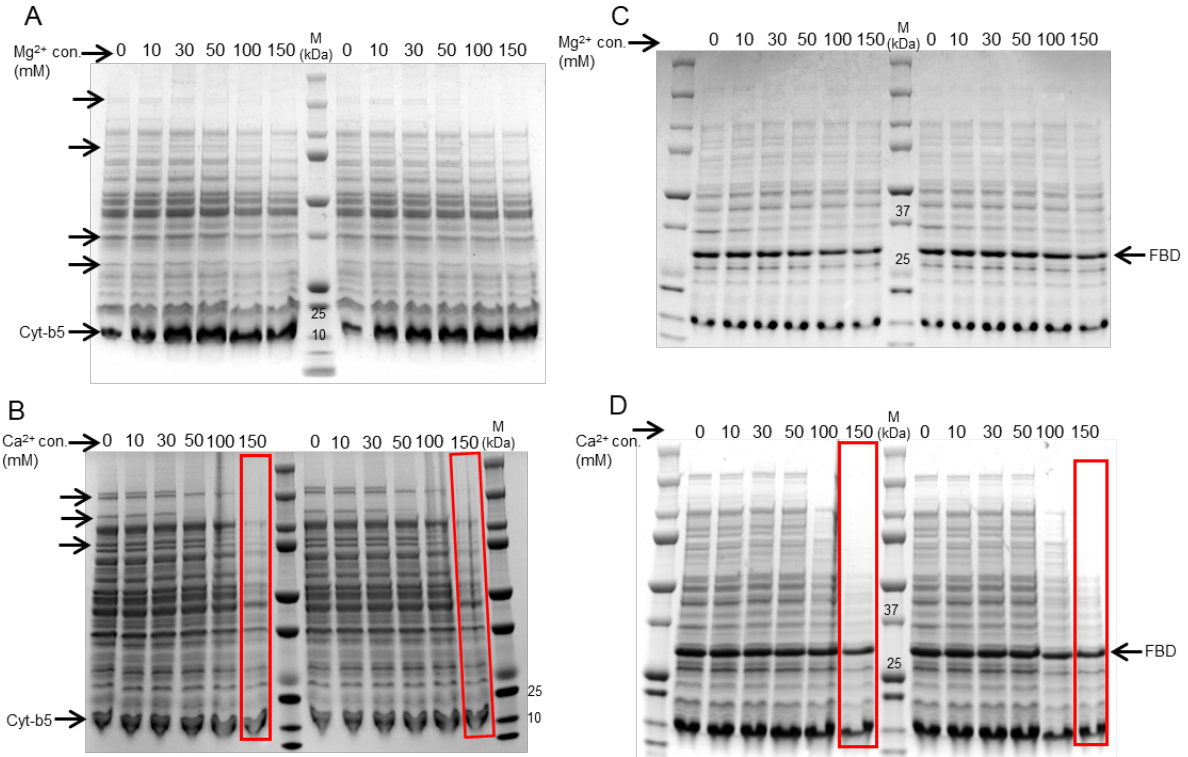


Figure S4. SDS-PAGE analysis of Cyt-b5 (A, B) and FBD-enriched (C, D) *E. coli* membranes that were solubilized using pentyl-inulin at the indicated concentrations of divalent metal ions. The protein band corresponding to Cyt-b5 and FBD are labelled, and the variations in the protein band intensity of *E. coli* membrane proteins at different metal ion concentrations are indicated with arrows. Only the well-resolved protein bands showing variation with different metal ion concentrations are indicated for clarity. The lanes highlighted in (B) and (D) with red boxes indicate a significant decrease in the intensity of protein bands at higher concentrations of divalent metal ions. The solubilization experiments were performed in technical replicates. M: protein marker

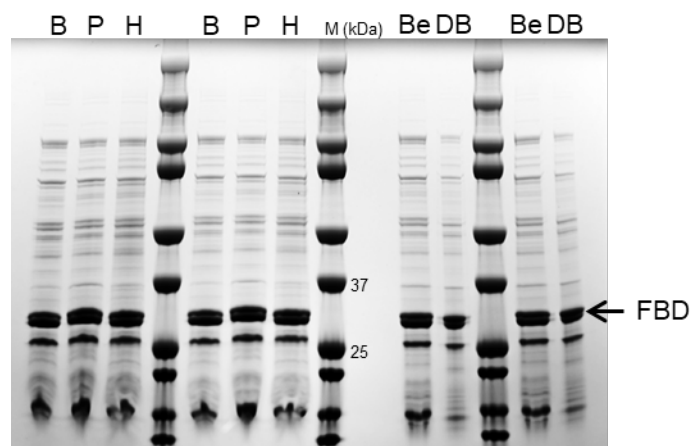


Figure S5. SDS-PAGE analysis of FBD-enriched *E. coli* membranes that were solubilized using inulin functionalized with butyl (B), pentyl (P), hexyl, (H), benzyl (Be) and di-benzyl (DB) hydrophobic groups. The solubilization experiments were performed in technical replicates.

References

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- [2] B. Krishnarjuna, T. Ravula, A. Ramamoorthy, Detergent-free extraction, reconstitution and characterization of membrane-anchored cytochrome-b5 in native lipids, *ChemComm*, 56 (2020) 6511-6514.
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- [4] T. Ravula, S.K. Ramadugu, G. Di Mauro, A. Ramamoorthy, Bioinspired, size-tunable self-assembly of polymer-lipid bilayer nanodiscs, *Angew. Chem. Int. Ed.*, 56 (2017) 11466-11470.