



Figure S1. Clustal-W multiple sequence alignment and ESPrnt (Easy Sequencing in PostScript) visualization analysis of representative members of the NapC/NirT/TorC family of cytochromes including CycB (*c_{m552}*) from *N. europaea* highlighted in blue. White letter, red box designates strict identity; Red character designates similarity in a group; Yellow box designates similarity across groups. Bold letters above the alignment are highly/strictly conserved residues of particular interest (see text). Asterisk indicate possible distal axial ligand to the Fe heme as predicted in other studies (refs 16-22).