Fig. S3. Two different conserved N-terminal domains of *Bd* CRN proteins. Profile A was generated from an alignment of the first 55 amino acids of 42 DHA subfamily proteins, 5 DAB subfamily proteins, and 3 DXV subfamily proteins. Profile B was generated from an alignment of the first 52 amino acids of 3 DDC subfamily proteins, 10 DFB subfamily proteins, 6 DX8 subfamily proteins, and 2 DXV subfamily proteins. The two conserved motifs in profile B are “FYIQYL(D/H/E/T)NQPV” (6-11aa) and “LVAA”(40-43aa). These logos were generated using WebLogo ([http://weblogo.berkeley.edu](http://weblogo.berkeley.edu)).