A

B

No.	sequence	M.W.	Predicted peptide
1,	40-KVIDPQHSDK	54.6	40-509
2,	40-KVIDPQHSDK	46.4	40-460
3,	127-SINPDRNTVT	44.4	127-509
4,	158-AEPAEIKYDY	42.0	158-509
5,	158-AEPAEIKYDY	39.6	158-495
6,	241-GELQDYVHQD	29.0	241-495

C

Supplemental Fig. 3. Proteinase K-digested peptides of Ndi1. *A*, The numbers indicate the digested peptides, and correspond to the numbers in panels *B* and *C*. *B*, N-terminal amino acid sequence of each peptide. Molecular weight of each peptide was calculated from the mobility on SDS-PAGE. *C*, Visualized models of digested peptides. Peptides number 1 to 5 contain the NADH binding motif (GXGXXG).