Additional file 2

Supplemental Figure S2. Sequences of EF-hand motifs in Nox5, Duox, plant Nox, and NoxC The original alignment data are shown in Additional file 8. EF (I) to EF (IV) represent four EF-hands of Nox5. Each EF-hand domain contains a twelve residue loop that is involved in calcium-binding. The letters X, Y, Z, -Y, -X, and -Z indicate the six residues that directly participate in calcium-binding, and these six residues correspond to positions 1, 3, 5, 7, 9 and 12. The consensus sequences of these six positions are: D at position 1, D/N/S at position 3, D/E/N/S/T/G at position 5, G/P at position 7, D/E/N/Q/S/T/A/G/C at position 9, and D/E at position 12 (PROSITE documentation number PDOC00018), and the most conserved positions are 1, 3, and 12. Asterisks indicate atypical EF-hand motif that is not consistent with the consensus sequences at positions 1, 3, or 12. Predicted T. rubripes (Tr)-Nox5 sequence was not a complete sequence; therefore the 1st EF-hand motif sequence was not shown. The arrows indicate motifs that have 17 to 23 additional amino acids inserted in the indicated position. Abbreviations of species are as in Figure 2.

	EF-I	EF-II	EF-III	EF-IV
	X Y Z-Y-X-Z	X Y Z-Y-X - Z	X Y Z-Y-X - Z	X Y Z-Y-X - Z
human-Hs-Nox5α	AG-EDGEISLQE*	DSDRSGTITLQE	DIDGSGSIDPDE	DADGNGAITFEE
			DVDGSGS1DADE DVDGSGS1DADE	DKDCNGAITFDE
dog-Cf-Nox5	AE-KDREINLQQ*		DVDGSGSIDADE	
cow-Bt-Nox5	AG-EDGEINLQD*	DSDGSGTITLQE		DODGNOSTITEE
chicken-Gg-Nox5	AG-HDEEIGLEE*		DVDGSGSIDAAE	DQDGNGSITFQE
frog-Xt-Nox5	AG-DDKEIDLEE*		DVDGSGSIDPSE	DKDHSGSITFQE
opposum-Md-Nox5	AG-EDREIDLQE*		DVDGNGSIDPDE	DKDHSGSITFEE
fugu-Tr-Nox5	AC DEVELVE OF	DSDGSSSISLDE	DVDGSGSIDPDE	DKDHSGSITFEE
medaka-Ol-Nox5	AG-DDKEINLCE*		DVQGSGSIDPDE*	DTDNSGSITFEE
sea urchin-Sp-Nox5A	AG-EDRQIDEDE*	•	DVDGNGAIDHEE	DTDGSGAISFEE
sea urchin-Sp-Nox5B	AG-DDNLIDLDE*	•	DVDGSGFIDFDE	DVDGDGEVSFEE
fruit fly-Dm-Nox5	VG-NEQEIRREE*	•	DIDGDGLIQHKE	DPHNSGEITYEA*
mosquito-Ag-Nox5	VG-NEKEIRREE*	•	DLDGDGLIQHRE	DKYNRGAITYEA*
honeybee-Am-Nox5	VG-NEKEIRREE*	DKDNSGTISLQE	DIDGDGLIQLRE	DQSNRGAITFEA*
fungus-Mg-NoxC			DHDGDGCIDYSE	
fungus-Fg-NoxC			DHDNDGHINYEE	
At-rbohF			DKNEDGRITEEE	DPERLGYIELWQ*
At-rbohl			CYQLSSNLVKHI *	APDGLYYIELKD*
At-rbohC			DKDADGRLTEDE	DPDNIGYIMLES*
At-rbohG			DKDSDGRLTEDE	DPDHMGYIMMES*
At-rbohA			DKDSDGRLNEAE	DPYHYGYIMIEN*
At-rbohD			DKDEDGRVTEEE	DPDNAGFIMIEN*
At-rbohB			DKNLDGRITGDE	DRDNLGYIELHN*
At-rbohE			DSNEDGKITREE	DPENFGYIELWQ*
At-rbohH			DKNGDGKLTEEE	DPDHKGYIEMWQ*
At-rbohJ			DKDGDGKLTEEE	DPNEQGYIEMWQ*
amoeba-Dd-NoxC			DIYDKGFISRDD*	DKNMDGYIDFEE
mouse-Mm-Duox1		DKDGNGYLSFRE	DFDGNGLISKDE	
mouse-Mm-Duox2		DKDGNGYISFRE	DLDGNGFLSKDE	
dog-Cf-Duox1		DKDGNGYLSFRE	DFDGNGLISKDE	
dog-Cf-Duox1		DKDGNGYLSFRE	DLDANGFLSKDE	
human-Hs-Duox1		DKDGNGYLSFRE	DFDGNGLISKDE	
human-Hs-Duox2		DKDGNGYLSFRE	DLDENGFLSKDE	
rat-Rn-Duox1		DKDGNGYLSFRE	DFDGNGLISKDE	
rat-Rn-Duox2		DKDGNGYISFRE	DLDGNGFLSKEE	
chicken-Gg-Duox		DKDGNGYISFRE	DIDENGFLSKEE	
frog-Xt-Duox1		DKDHNGYLSFEE	DVNGNGILPKEE	
frog-Xt-Duox2		DEDGNGYLSFRE	DVDGNGFLSKEE	
fugu-Tr-Duox		DKDGNGSLSFQE	DIGGTGSLSKGE*	
tetraodon-Tn-Duox		DKDGNGYLSFQE		
zebrafish-Dr-Duox		DKDGNGYLSFQE	DIKGDGFLSKEE*	
medaka-OI-Duox		DTDHSGYLSFQE	DVGGNGYLSKEE	
ascidian-Ci-Duox-B		DTDHSGYLSFRE	DVDHSGEINREE	
ascidian-Ci-Duox-A		DSDGSGAISFRE	DLDKSGELSKKE	
ascidian-Ci-Duox-C		DSDEDGTISFRE	DLDKSGGLSKEE	
ascidian-Ci-Duox-D		DEDQDGFISFHD	DLNQNGSLTKQQ *	
sea urchin-Sp-Duox		DQDNSGS1SFRE	DIDRSGHLSREE	
fuit fly-Dm-Duox		DKDQDGRISFQE	DNDRNGVIDKGE	
mosquito-Ag-Duox		DKDKDGRISFQE	DNDRNGV I DKGE	
honeybee-Am-Duox		DKDRDGRISFQE	DKDCNGVIDKEE	
nematode-Ce-Duox1		AKHNEDSLSFNE*	DLEGKNKVLRKD*	
nematode-Ce-Duox2		AKHNEDSLSFNE *		
		THE TOPOLOGICAL	PPPOINTILLFIRID	