

Supplementary Data

SUPPLEMENTARY TABLE S1. PROPOSED SYSTEMATIC NOMENCLATURE FOR THE CYB561 PROTEIN FAMILY

Species	Systematic name	Names used so far	Protein ID	References
<i>Arabidopsis thaliana</i>	At_CYB561B1	CYBASC1, TCytb, TCB, CLb561A, Acytb ₅₆₁ -A, b561A.tha5	Q8L856	(2), (4), (5), (7), (17), (24)
<i>Bos taurus</i>	At_CYB561B2	CYBASC2, Acytb ₅₆₁ -B, b561A.tha4	Q9SWS1	(5), (24)
<i>Citrus limon</i>	Bt_CYB561A1	cytochrome b ₅₆₁ , Cyt b561, b561Bovine	P10897	(9), (19), (21), (23), (24)
	Cl_CYB561B1	CLb561A	Q3LGX5	(14)
	Cl_CYB561B2	CLb561B	Q3LGX4	(14)
<i>Drosophila melanogaster</i>	Dm_CYB561A1	nemy, b561D.me1	Q95T77	(8), (24)
<i>Homo sapiens</i>	Hs_CYB561A2	Cybrd1, Dcytbhuman	Q53TN4	(10), (11), (15), (24)
	Hs_CYB561D1	CYB561D1	Q8N8Q1	(22)
	Hs_CYB561D2	101F6, CYB561D2, TSFhuman	O14569	(16), (18), (24)
<i>Mus musculus</i>	Mm_CYB561A1	CGCytb, CGCB, b561mouse	Q60720	(3), (24)
	Mm_CYB561A2	Dcytb, DCytb, DCB, Cybrd1, Dcytbmouse	Q925G2	(12), (20), (24)
	Mm_CYB561A3	LCytb, 'ubiquitous', LCB, b561mouse2	Q6P1H1	(20), (24), (25)
	Mm_CYB561D1	Cyb561d1	NP_001074789	(22)
	Mm_CYB561D2	mouse 101F6, Cyb561d2, TSP10, TSCytb, TSCB, TSFmouse	Q9WUJ3	(1), (4), (24)
<i>Schistosoma japonicum</i>	Sj_CYB561A1	SjCytb561, b561Schisto	Q5D8X4	(6), (24)
<i>Zea mays</i>	Zm_CYB561B1	Zmb561, b561Z.mays	Q61681	(13), (24)

The systematic name suggested for the CYB561 protein family members, is based on (i) the HUGO Gene Nomenclature Committee adopted name for the cytochrome *b* super family, 'CYB'; (ii) the cytochrome *b*561-specific extension: '561'; (iii) a letter (A-E) referring to the evolutionary clusters identified by Tsubaki *et al.* (24)*, and (iv) a sequential number (1, 2, ...) for each CYB561 isoform within each species. Note that by lack of information on subcellular localization and physiological function for several CYB561 isoforms, identical numbers (e.g., Mm_CYB561A1 and Sj_CYB561A1) not necessarily mean identical functions or localizations across species. In this article, we refer to the proposed systematic name of the proteins, preceded, for clarity, by the common name that refers to the protein localization (*i.e.*, 'CG' for chromaffin granule; 'D' for duodenum; 'L' for lysosome; 'T' for tonoplast) or function (*i.e.*, 'TS' for tumor suppression).

*Clusters D and E from [24] were exchanged to accommodate for already published names (CYB561D1 and CYB561D2).

Supplementary References

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