

Table S4. Pairwise comparison of DIA1 and DIA1R proteins.

Protein		DIA1										DIA1R									
	Species ^a	Hsapi ^b	Mmula	Btaur	MmusC	Rnorv	Mdome	Ggall	Dr(a)	Dr(b)	Hsapi ^b	Mmula	Btaur	MmusC	Rnorv	Mdome	Ggall	Dreri			
Percentage amino acid similarity ^c	DIA1																				
	Hsapi	-	100	99	98	98	97	90	78	82	28	29	28	26	26	29	28	27			
	Mmula	100	-	99	98	98	97	90	78	82	28	29	28	26	26	29	28	27			
	Btaur	100	100	-	98	98	97	90	78	82	28	29	28	26	26	29	28	27			
	MmusC	99	99	99	-	100	96	89	78	82	28	29	27	27	27	29	28	27			
	Rnorv	99	99	99	100	-	96	89	78	82	28	29	27	27	27	29	28	27			
	Mdome	99	99	100	99	99	-	88	77	82	28	28	28	26	27	27	29	27			
	Ggall	96	96	96	99	96	96	-	78	82	29	29	28	27	27	28	28	29			
	Dreri(a)	96	96	96	97	95	96	95	-	88	28	28	28	26	27	27	26	25			
	Dreri(b)	97	97	97	97	97	97	97	98	-	28	28	28	25	26	28	29	26			
Percentage amino acid identity ^c	DIA1R																				
	Hsapi	62	62	62	63	63	62	62	62	62	-	97	83	81	80	74	65	45			
	Mmula	63	63	63	64	64	63	63	63	62	99	-	83	81	83	73	64	44			
	Btaur	61	61	61	64	64	62	62	63	62	96	94	-	79	78	70	62	46			
	MmusC	63	63	63	64	64	62	64	63	62	94	95	93	-	91	71	63	45			
	Rnorv	60	60	60	60	60	62	60	62	62	95	94	94	97	-	70	61	45			
	Mdome	63	63	63	63	63	63	63	65	62	91	91	90	90	89	-	66	48			
	Ggall	61	61	61	61	61	62	62	63	63	87	87	86	86	86	90	-	45			
	Dreri	59	59	59	59	59	60	62	60	60	78	78	77	76	76	78	76	-			

^aSpecies with both a DIA1 and DIA1R orthologue (on the current databases) are presented.

^bSpecies abbreviations are the first letter of the genus name, followed by the first four letters of the species name e.g. Homo sapiens = Hsapi. Full details of species names and protein accession numbers can be found in from Tables S1 and S5.

^cThe percent identity (above diagonal) and similarity (below diagonal) between proteins aligned pair-wise using CLUSTALW [47] are displayed.

Percentage amino acid identity^c