

# Figure 5E – source data

Table format: Grouped		Group A			Group B			Group C		
		ATP8B1/CDC50A dNC			Drs2 dNC			Na/K		
		A:Y1	A:Y2	A:Y3	B:Y1	B:Y2	B:Y3	C:Y1	C:Y2	C:Y3
1	-C-ter peptide	100	100	100	98	99	103	106.7	97.8	95.4
2	+C-ter peptide	10			100	77	83	85.9	85.5	90.1
3	Title									

# Figure 5E – statistics

ANOVA results × Multiple comparisons × Narrative results ×

2way ANOVA						
ANOVA results						
1	Table Analyzed	% of control, grouped				
2						
3	<b>Two-way ANOVA</b>	Ordinary				
4	Alpha	0.05				
5						
6	<b>Source of Variation</b>	<b>% of total variation</b>	<b>P value</b>	<b>P value summary</b>	<b>Significant?</b>	
7	Interaction	45.92	<0.0001	****	Yes	
8	C-ter peptide	65.40	<0.0001	****	Yes	
9	Protein type	45.88	<0.0001	****	Yes	
10						
11	<b>ANOVA table</b>	<b>SS (Type III)</b>	<b>DF</b>	<b>MS</b>	<b>F (DFn, DFd)</b>	<b>P value</b>
12	Interaction	3551	2	1776	F (2, 10) = 46.42	P<0.0001
13	C-ter peptide	5058	1	5058	F (1, 10) = 132.2	P<0.0001
14	Protein type	3548	2	1774	F (2, 10) = 46.38	P<0.0001
15	Residual	382.5	10	38.25		
16						
17	<b>Difference between row means</b>					
18	Predicted (LS) mean of -C-ter peptide	99.99				
19	Predicted (LS) mean of +C-ter peptide	61.28				
20	Difference between predicted means	38.71				
21	SE of difference	3.367				
22	95% CI of difference	31.21 to 46.21				
23						
24	<b>Data summary</b>					
25	Number of columns (Protein type)	3				
26	Number of rows (C-ter peptide)	2				
27	Number of values	16				
28						