

	#winners <sup>a</sup>	total <sup>b</sup>	%	Fisher's exact test	% difference	Fisher's exact test	% difference
<b>exons vs. pseudo exons</b>							
ESE	478	979	48.8%	-	-	-	-
ESS	194	496	39.1%	ESE vs. ESS	P<0.001	9.7%	
Neutral	1118	2621	42.7%	ESE vs. neutral	P=0.001	6.1%	ESS vs. neutral <i>P=0.15</i> -3.6%
<b>exons vs. decoy regions</b>							
ESE	513	979	52.4%	-	-	-	-
ESS	193	496	38.9%	ESE vs. ESS	P<0.001	13.5%	
Neutral	1198	2621	45.7%	ESE vs. neutral	P<0.001	6.7%	ESS vs. neutral <i>P=0.006</i> -6.8%
<b>exons vs. intron flanks</b>							
ESE	489	979	49.9%	-	-	-	-
ESS	202	496	40.7%	ESE vs. ESS	P<0.001	9.2%	
Neutral	1119	2621	42.7%	ESE vs. neutral	P<0.001	7.2%	ESS vs. neutral <i>P=0.43</i> -2.0%

Table S5: Comparison of PU values of splicing motifs between data sets.