

Supplemental Table 2. Percentage of PolyA and PolyG sequences that reside in NFRs for all species.

Seq.	Scer	Spar	Smik	Sbay	Cgla	Scas	Klac	Kwal	Sklu	Dhan	Calb	Ylip	Spom
G5	56.14	40.18	45.48	40.35	66.84	42.68	42.94	46.09	61.51	43.81	44.52	29.00	47.71
G6	55.79	42.99	45.05	38.07	71.50	34.45	48.61	48.79	61.36	50.57	43.43	32.39	53.18
G7	69.09	54.93	55.40	37.81	73.35	44.41	52.11	51.92	62.33	56.44	47.47	33.72	55.69
G8	63.99	54.21	56.06	32.60	76.59	33.11	44.72	50.12	65.11	45.30	45.59	43.92	53.95
A5	58.23	55.80	61.19	63.67	69.09	55.94	60.47	59.27	63.79	54.17	59.57	53.47	46.77
A6	59.99	58.71	63.06	67.51	71.04	60.26	68.24	68.89	68.35	59.78	64.55	55.86	49.52
A7	63.76	63.83	69.10	74.01	75.44	61.72	73.16	71.59	73.05	65.88	68.19	62.74	53.55
A8	69.94	70.09	74.26	75.21	80.93	67.38	76.35	74.56	75.93	71.76	72.42	62.53	61.37