

Supplementary Table 4 — Gene Length, Structure and Functional Type Statistics on Final Annotation Gene Sets

	Intersection Set	Aragorn-Only	Union Set
<b>#tRNA genes</b>	3579	36	3616
<b>Avg. gene length</b>	74	98	75
<b>Min. gene length</b>	68	71	68
<b>Max. gene length</b>	89	206	206
<b>%intron</b>	2	28	3
<b>%GC</b>	58	59	58
<b>#Ala tRNA genes</b>	210	2	212
<b>#Cys tRNA genes</b>	64	1	65
<b>#Asp tRNA genes</b>	105	1	106
<b>#Glu tRNA genes</b>	160	1	161
<b>#Phe tRNA genes</b>	104	2	106
<b>#Gly tRNA genes</b>	228	3	231
<b>#His tRNA genes</b>	80	4	84
<b>#Ile tRNA genes</b>	171	1	172
<b>#Lys tRNA genes</b>	183	1	184
<b>#Leu tRNA genes</b>	335	6	341
<b>#Met tRNA genes</b>	97	0	97
<b>#Asn tRNA genes</b>	125	0	125
<b>#Pro tRNA genes</b>	200	0	200
<b>#Gln tRNA genes</b>	161	0	161
<b>#Arg tRNA genes</b>	348	2	350
<b>#Ser tRNA genes</b>	228	7	235
<b>#Thr tRNA genes</b>	218	4	222
<b>#Val tRNA genes</b>	236	0	236
<b>#Trp tRNA genes</b>	52	1	53
<b>#iMet tRNA genes</b>	76	0	76
<b>#Tyr tRNA genes</b>	88	0	88
<b>#SeC tRNA genes</b>	76	0	76
<b>#Ambiguous tRNA genes</b>	34	0	35