

B

5SrRNA gene	TriTrypDB ID	Lenght	Strand	Position	Termination Type	Lenght of 1st T Cluster	Lenght of 2nd T Cluster	Distance 1st & 2nd T cluster	
5	LmjF.05.5SrRNA.01	119	+	360083..360201	D	8	-	-	
9-1	LmjF.09.5SrRNA.03	119	-	277005..276887	C	6	-	-	
9-3	LmjF.09.5SrRNA.02	118	+	405671..405788	D	27	-	-	
9-4	LmjF.09.5SrRNA.01	119	-	418738..418620	D	14	-	-	
11-1	LmjF.11.5SrRNA.01	119	-	156905..156787	D	11	-	-	
Lm 11-2	LmjF.11.5SrRNA.02	119	+	163008..163126	D	11	-	-	
11-3	LmjF.11.5SrRNA.03	118	+	381623..381740	D	16	-	-	
15	LmjF.15.5SrRNA.01	119	+	323210..323328	D	14	-	-	
21-1	LmjF.21.5SrRNA.01	119	-	166286..166168	D	10	-	-	
21-2	LmjF.21.5SrRNA.02	119	+	448851..448969	B	2	Avg: 9	4	
23	LmjF.23.5SrRNA.01	119	-	225680..225562	D	11	11.8	-	
5	LmxM.05:rRNA:rfamscan:361991-362109	120	+	361991..362110	D	10	-	-	
9-1	LmxM.09:rRNA:rfamscan:251967-252085	119	-	252085..251967	C	7	-	-	
9-3	LmxM.09:rRNA:rfamscan:380921-381039	119	+	380921..381039	D	12	-	-	
9-4	LmxM.09:rRNA:rfamscan:391554-391672	119	-	391672..391554	D	6	-	-	
11-1	LmxM.11:rRNA:rfamscan:157979-158097	119	-	158097..157979	B	5	4	12	
Lx 11-2	LmxM.11:rRNA:rfamscan:163755-163875	119	+	163755..163873	B	5	4	12	
11-3	LmxM.11:rRNA:rfamscan:361629-361747	119	+	361629..361747	D	12	-	-	
15	LmxM.15:rRNA:rfamscan:288997-289115	119	+	288997..289115	D	16	-	-	
21-1	LmxM.21:rRNA:rfamscan:156060-156178	119	-	156178..156060	C	4	-	-	
21-2	LmxM.21:rRNA:rfamscan:430811-430929	119	+	430811..430929	D	7	Avg: -	-	
23	LmxM.23:rRNA:rfamscan:217667-217785	120	-	217785..217666	B	5	8.09	5	8
9-3	LbrM.09.rRNA1	119	+	395413..395531	D	14	-	-	
9-4*	*	123	-	403745..403623	C	5	-	-	
11-1*	*	119	-	63370..63252	D	12	-	-	
11-2	LbrM.11.rRNA1	119	-	68610..68492	D	12	-	-	
Lb 21-1*	*	123	-	159391..159269	C	5	-	-	
21-2	LbrM.21.rRNA1	119	+	430680..430798	D	7	-	-	
23	lbrM.23.rRNA1	119	-	215667..215549	D	11	-	-	
31*	*	119	-	965247..965129	D	16	Avg: -	-	
NA*	*	119	-	607851..607733	D	7	9.88	-	-
9-1	LinJ.09.rRNA1	118	-	266769..266652	C	7	-	-	
9-3	LinJ.09.rRNA2	119	+	400163..400281	B	5	4	18	
9-4	LinJ.09.rRNA3	119	-	413091..412973	D	12	-	-	
11-1*	*	119	-	162781..162663	B	7	4	11	
Li 11-2*	*	119	-	212062..211944	B	7	4	11	
11-3	LinJ.11.rRNA1	119	+	371388..371506	D	10	-	-	
21-1*	*	125	-	163366..163242	D	8	-	-	
21-2	LinJ.21.rRNA1	118	+	445667..445784	D	15	Avg: -	-	
NA* 23	LinJ.23.rRNA1	119	-	228303..228185	B	6	8.55	4	9
5	*	119	+	359812..359930	D	10	-	-	
9-1	*	119	-	249842..249724	D	6	-	-	
9-2	*	118	-	257062..256945	B	7	5	26	
9-3	*	118	+	377890..378007	D	29	-	-	
Lt 9-4	*	118	-	390508..390391	D	21	-	-	
11-3	*	121	+	390508..390391	D	11	-	-	
21-1	*	121	-	167666..167546	C	5	-	-	
21-2	*	119	+	442984..443102	D	14	-	-	
23	*	122	-	213231..213110	D	27	Avg: -	-	
33	*	119	+	649716..649834	D	11	14.1	-	-

Figure S1

C

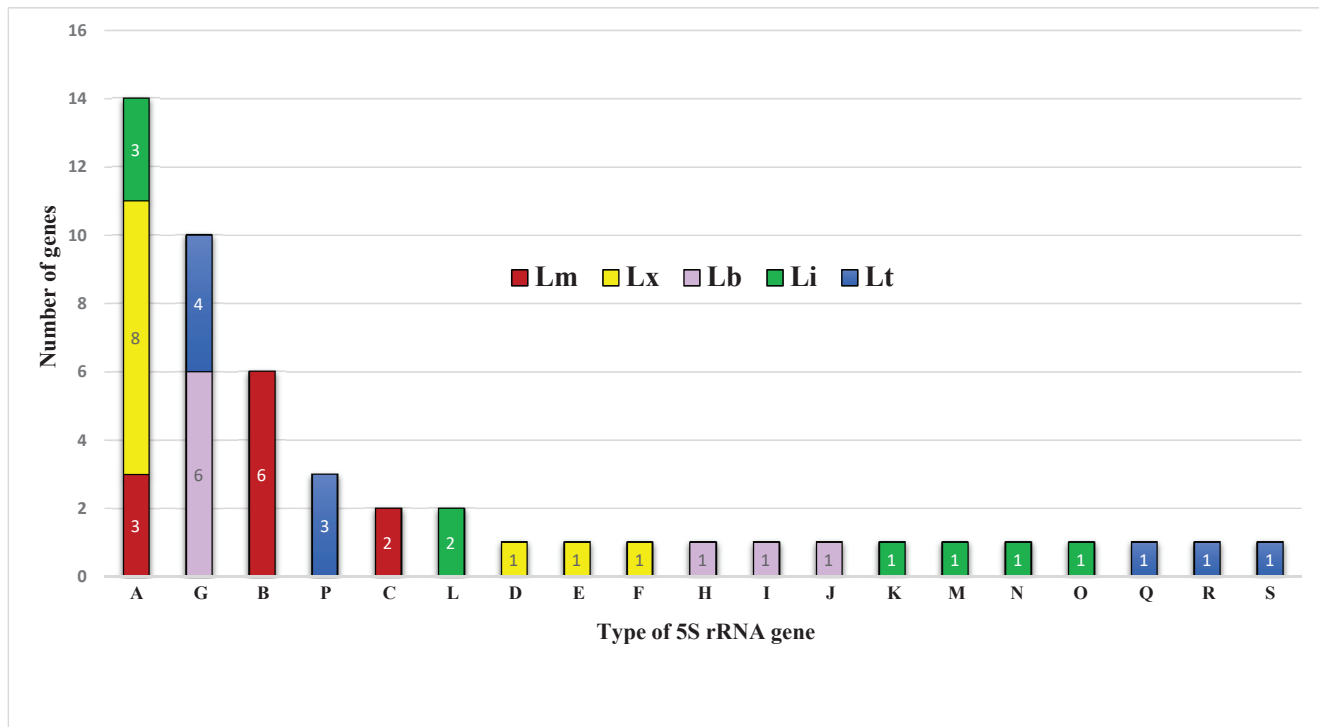


Figure S1