

Macrostomum lignano

Name	Start	Stop	Length	Start codon	Stop codon	Strand
cox1	1	1548	1548	att	taa	+
intergenic spacer	1549	1559	11			
cob	1560	2660	1101	att	tag	+
trnH(gtg)	2661	2720	60			+
trnQ(ttg)	2721	2775	55			+
trnW(tca)	2776	2837	62			+
trnT(tgt)	2838	2896	59			+
trnP(tgg)	2897	2959	63			+
intergenic spacer	2960	2968	9			
rrnL	2969	3835	867			+
intergenic spacer	3836	3845	10			
trnV(tac)	3846	3905	60			+
trnE(ttc)	3906	3965	60			+
trnS1(gct)	3963	4015	53			+
nad1	4015	4887	873	ttg	taa	+
intergenic spacer	4888	4912	25			
atp6	4913	5590	678	ttg	taa	+
intergenic spacer	5591	5600	10			
nad3	5601	5953	353	att	ta	+
intergenic spacer	5954	5985	32			
cox3	5986	6774	789	gtt	taa	+
intergenic spacer	6775	6776	2			
nad6	6777	7214	438	atg	taa	+
trnM(cat)	7213	7272	60			+
trnA(tgc)	7273	7334	62			+
trnG(tcc)	7335	7395	61			+
cox2	7396	8070	675	ata	tag	+
trnK(ctt)	8069	8121	53			+
trnF(gaa)	8122	8183	62			+
nad2	8181	9071	891	gtg	tag	+
trnI(gat)	9072	9133	62			+
nad4l	9133	9390	258	atg	taa	+
trnS2(tga)	9390	9443	54			+
intergenic spacer	9444	10165	722			
nad4	10166	11392	1227	ttg	taa	+
trnD(gtc)	11391	11448	58			+
trnR(tcg)	11449	11508	60			+
rrnS	11450	12188	739			+
intergenic spacer	12189	12204	16			
nad5	12205	13761	1557	atg	taa	+
trnN(gtt)	13760	13819	60			+
trnY(gta)	13819	13880	62			+
trnL2(taa)	13881	13940	60			+
trnC(gca)	13941	13999	59			+
intergenic spacer	14000	14012	13			
atp8	14013	14180	168	ata	taa	+
intergenic spacer	14181	14193	13			

Stenostomum sthenum

Name	Start	Stop	Length	Start codon	Stop codon	Strand
cox1	1	1557	1557	atc	tag	+
intergenic spacer	1558	1566	9			
trnA(tgc)	1567	1623	57			+
trnK(ttt)	1624	1687	64			+
intergenic spacer	1688	1695	8			
trnE(ttc)	1696	1759	64			+
intergenic spacer	1760	1760	1			
trnL2(taa)	1761	1825	65			+
intergenic spacer	1826	1828	3			
trnH(gtg)	1829	1889	61			-
intergenic spacer	1890	1939	50			
rrnS	1940	2618	679			-
intergenic spacer	2619	2638	20			
trnV(tac)	2639	2698	60			-
nad3-1	2699	3062	364	atg	t	-
intergenic spacer	3063	3114	52			
trnL1(tag)	3115	3174	60			-
intergenic spacer	3175	3179	5			
trnM(cat)	3180	3245	66			-
intergenic spacer	3246	3247	2			
trnI(gat)	3248	3306	59			+
intergenic spacer	3307	4146	840			
atp8	4147	4276	130	atc	t	-
intergenic spacer	4277	4424	148			
trnN(gtt)	4425	4484	60			+
cob	4485	5579	1095	atc	tag	+
intergenic spacer	5580	5585	6			
trnP(tgg)	5586	5649	64			+
nad6	5650	6153	504	att	tag	+
trnS1(gct)	6119	6173	55			+
cox3	6174	6953	780	atc	taa	+
intergenic spacer	6954	6963	10			
nad1	6964	7860	897	atg	taa	+
intergenic spacer	7861	7862	2			
trnY(gta)	7863	7924	62			+
intergenic spacer	7925	7926	2			
trnC(gca)	7927	7989	63			+
intergenic spacer	7990	8015	26			
rrnL	8016	8965	950			+
intergenic spacer	8966	9030	65			
trnI(gat)	9031	9089	59			-
intergenic spacer	9090	9091	2			
trnM(cat)	9092	9157	66			+
intergenic spacer	9158	9162	5			
trnL1(tag)	9163	9222	60			+
intergenic spacer	9223	9274	52			
nad3-0	9275	9638	364	atg	t	+
trnV(tac)	9639	9698	60			+
intergenic spacer	9699	9718	20			
rrnS	9719	10397	679			+

intergenic spacer	10398	10447	50		
trnH(gtg)	10448	10508	61		+
intergenic spacer	10509	10511	3		
trnL2(taa)	10512	10576	65		-
intergenic spacer	10577	10577	1		
trnE(ttc)	10578	10641	64		-
intergenic spacer	10642	10649	8		
trnK(ttt)	10650	10713	64		-
trnA(tgc)	10714	10770	57		-
intergenic spacer	10771	10801	31		
cox2	10802	11479	678 ata	taa	-
intergenic spacer	11480	11540	61		
trnM(cat)	11541	11606	66		-
nad4l	11602	11859	258 ata	taa	-
intergenic spacer	11860	11866	7		
trnW(tca)	11867	11930	64		+
intergenic spacer	11931	11931	1		
trnQ(ttg)	11932	11994	63		-
intergenic spacer	11995	11996	2		
trnT(tgt)	11997	12059	63		-
intergenic spacer	12060	12061	2		
nad4	12062	13294	1233 ata	taa	+
trnS2(tga)	13293	13354	62		-
intergenic spacer	13355	13357	3		
trnD(gtc)	13358	13420	63		+
atp6	13421	14086	666 ata	taa	+
intergenic spacer	14087	14089	3		
nad5	14090	15721	1632 att	tag	+
trnF(gaa)	15696	15755	60		+
intergenic spacer	15756	15787	32		
nad2	15788	16750	963 att	taa	+
trnG(tcc)	16749	16809	61		+
trnR(tcg)	16808	16866	59		-
intergenic spacer	16867	16944	78		