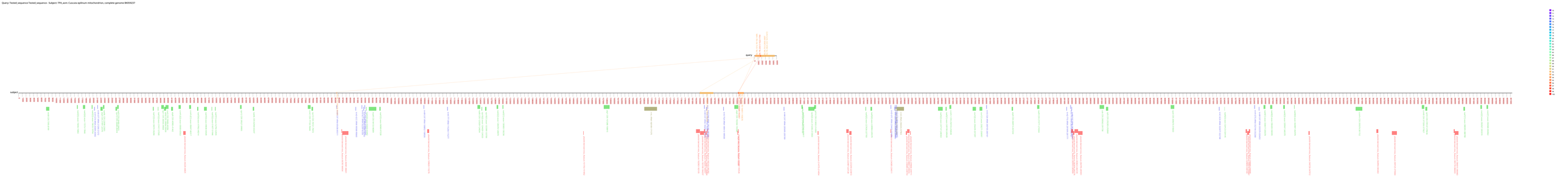

Query: Tested_sequence Tested_sequence - Subject: TPA_asm: Cuscuta epilinum mitochondrion, complete genome BK059237

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mttB CDS 7395-8199

nad2[5] exon 15694-15882

nad2[4] exon 17273-17846

nad2[3] exon 19633-19794

nad2[2] exon 128478-128870

nad2[1] exon 130083-130236

trnY-GTA tRNA 20309-20392

trnN-GTT tRNA 21200-21272

atp4 CDS 21991-22570

nad4L CDS 22698-23001

rps12 CDS 26040-26418

nad3 CDS 26466-26823

nad1[3] exon 36074-36266

nad1[2] exon 37417-37500

nad1[4] exon 93433-93492

nad1[5] exon 96969-97228

nad1[1] exon 388395-388780

rps1 CDS 38350-39004

sdh4 CDS 39094-39559

cox3 CDS 39486-40284

atp8 CDS 40985-41462

ccmFc2[2] exon 43000-43562

ccmFc2[1] exon 45847-46317

nad7[4] exon 47952-48214

nad7[3] exon 49838-50549

nad7[2] exon 51877-51946

nad7[1] exon 52792-52935

rps12 CDS 59527-59905

nad4L CDS 62944-63247

ccmC CDS 77720-78446

rps13 CDS 78721-79072

trnM-CAT tRNA 85438-85511

trnG-GCC tRNA 90588-90660

trnS-GCT tRNA 92238-92326

trnF-GAA tRNA 92681-92755 trnP-TGG tRNA 92990-93065

matR CDS 94133-96089

trnM-CAT tRNA 108852-108926

trnK-TTT tRNA 115204-115277

nad6 CDS 123329-124034

rps10[2] exon 124420-124533

rps10[1] exon 125399-125700

atp1 CDS 157288-158815

rrnL rRNA 168189-171575

trnM-CAT tRNA 184297-184370

trnL-TAA tRNA 184803-184917 trnF-GAA tRNA 185186-185259

trnS-TGA tRNA 189413-189500

atp6 CDS 192366-193380

trnM-CAT tRNA 205695-205769

rps19 CDS 210445-210730 rps3[1] exon 210743-210817

rps3[2] exon 212282-213903

rpl16 CDS 213832-214309

nad5[5] exon 227636-227783

nad5[4] exon 228984-229379

nad5[2] exon 247147-248363

nad5[1] exon 249198-249428

nad5[3] exon 324123-324145

trnD-GTC tRNA 234422-234496

trnF-AAA tRNA 235428-235493 trnW-CCA tRNA 235589-235663 rrn5 rRNA 235720-235839 trnQ-CTG tRNA 235932-235993

rrnS rRNA 236028-237963

ccmFc1 CDS 250155-250638

cox1[2] exon 256439-257297

cox1[1] exon 258271-258997

trnS-CGA tRNA 260103-260161

atp9 CDS 266913-267234

nad9 CDS 273771-274344

trnQ-TTG tRNA 281699-281771

trnE-TTC tRNA 282720-282792 trnM-CAT tRNA 283016-283087

cob CDS 290562-291738

rpl5 CDS 292283-292868

rps4 CDS 309670-310603

trnC-GCA tRNA 322677-322748

trnH-GTG tRNA 332194-332268

trnM-CAT tRNA 333379-333453

nad4[1] exon 334617-335078

nad4[2] exon 336420-336935

nad4[3] exon 339947-340370

nad4[4] exon 342887-342976

ccmFn CDS 359376-361152

ccmB CDS 377200-377827

rpl10 CDS 378147-378636

cox2[2] exon 392927-393319

cox2[1] exon 394580-394962

plastid-derived misc_feature 44220-44825

plastid-derived misc_feature 86787-86934

plastid-derived misc_feature 86981-88601

plastid-derived misc_feature 109820-110276

plastid-derived misc_feature 151753-151852

plastid-derived misc_feature 182083-183105

plastid-derived misc_feature 183190-184237

plastid-derived misc_feature 184245-184457 plastid-derived misc_feature 184637-184890 plastid-derived misc_feature 185143-185282

atp6 misc_feature 193241-193367 plastid-derived misc_feature 193387-193526

plastid-derived misc_feature 214775-214954

plastid-derived misc_feature 222487-223128

plastid-derived misc_feature 223259-223813

plastid-derived misc_feature 234385-234511

plastid-derived misc_feature 238489-238753

plastid-derived misc_feature 238807-239455

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plastid-derived misc_feature 282933-283409 plastid-derived misc_feature 283434-283776

plastid-derived misc_feature 283823-284729

plastid-derived misc_feature 284795-285951

plastid-derived misc_feature 329850-330165 plastid-derived misc_feature 330210-330687 plastid-derived misc_feature 330689-330999

plastid-derived misc_feature 346726-347012

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plastid-derived misc_feature 385778-386003

plastid-derived misc_feature 386015-386983

92.1% 3824 233 1673-5472

92.1% 3824 233 183094-186708

92.7% 1312 78 192-1495

92.7% 1312 78 194911-193670

96.1% 233 0 1522-1754

96.1% 233 0 193589-193357

93.3% 75 0 2933-3007

93.3% 75 0 85435-85509

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