

1 ANEXOS

1.1 PROTOCOLO BLAST LOCAL DE PARA LA EXTRACCIÓN DE SECUENCIAS:

- 1.1.1 Descargar el programa BLAST de la página de la NCBI en el siguiente link:
https://blast.ncbi.nlm.nih.gov/Blast.cgi?PAGE_TYPE=BlastDocs&DOC_TYPE=Download
- 1.1.2 Guardar la “secuencia local” (su secuencia completa de la que quiere hacer la extracción) como un documento fasta en una carpeta creada previamente en el escritorio.

La secuencia local que usted observa a continuación es una secuencia FASTA descargada del NCBI que contiene por defecto 9 archivos con diferentes formatos (ej. NHR, NND, NSI, NAL, entre otros):

| | | | |
|--|-----------------------|----------------|--------------|
|  refseq_rna.00.nhr | 26/08/2016 2:03 a ... | Archivo NHR | 392,644 KB |
|  refseq_rna.00.nin | 26/08/2016 2:03 a ... | Archivo NIN | 29,350 KB |
|  refseq_rna.00.nnd | 26/08/2016 2:03 a ... | Archivo NND | 19,804 KB |
|  refseq_rna.00.nni | 26/08/2016 2:03 a ... | Archivo NNI | 78 KB |
|  refseq_rna.00.nog | 26/08/2016 2:03 a ... | Archivo NOG | 9,784 KB |
|  refseq_rna.00.nsd | 26/08/2016 2:03 a ... | Archivo NSD | 104,632 KB |
|  refseq_rna.00.nsi | 26/08/2016 2:03 a ... | Archivo NSI | 2,177 KB |
|  refseq_rna.00.nsq | 26/08/2016 2:03 a ... | Archivo NSQ | 976,562 KB |
|  refseq_rna.00.tar.gz | 26/08/2016 2:40 p.... | Archivo WinRAR | 1,005,187 KB |
|  refseq_rna.nal | 26/08/2016 2:15 a ... | Archivo NAL | 1 KB |

Se recomienda en este paso revisar los ejecutables de la página del NCBI en el siguiente link <https://www.ncbi.nlm.nih.gov/guide/howto/run-blast-local/> para que pueda seguir el mismo ejemplo.

- 1.1.3 Descargar la secuencia “consulta” (o de referencia) deseada del gen bank y guardarla en la misma carpeta del escritorio en formato .txt (bloc de notas) con códigos fasta. Debe tener en cuenta que esta secuencia es la que usted quiere alinear con su secuencia local.

| | | | |
|--|----------------------|---------------------|------|
|  test_query.txt | 27/08/2016 11:20 ... | Documento de tex... | 3 KB |
|--|----------------------|---------------------|------|

1.1.4 Abrir el cmd del ordenador (“command prompt” o “símbolo del sistema”). Este lo encuentra en el buscador de su ordenador.

1.1.5 Ejecutar en el cmd los siguientes comandos:

- C:\Users>cd\“ubicación de la carpeta” (sin comillas)

```
C:\Users>cd\Users\David Arley\Desktop\blast-2.4.0+\db
```

En este caso, la ubicación de la carpeta es: Users\David Arley\Desktop\blast-2.4.0+\db

- >makeblastdb –in “nombre del documento fasta” –dbtype nucl –parse_seqids (sin comillas)

```
C:\Users\David Arley\Desktop\blast-2.4.0+\db>makeblastdb -in refseq_rna.00 -dbtype nucl -parse_seqids
```

Aquí el nombre del documento fasta se llama: **refseq_rna.00**

Aparecerá:

```
Building a new DB, current time: 11/15/2017 22:22:40
New DB name: C:\Users\David Arley\Desktop\blast-2.4.0+\db\refseq_rna.00
New DB title: refseq_rna.00
Sequence type: Nucleotide
Keep MBits: T
Maximum file size: 1000000000B
BLAST options error: File refseq_rna.00 does not exist
```

- >blastdbcmd –db “documento fasta” –info (sin comillas)

```
C:\Users\David Arley\Desktop\blast-2.4.0+\db>blastdbcmd -db refseq_rna.00 -info
```

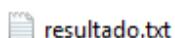
Aparecerá:

```
Database: NCBI Transcript Reference Sequences
          2,504,470 sequences; 3,991,728,284 total bases
Date: Aug 26, 2016 3:00 AM      Longest sequence: 107,499 bases
Volumes:
          C:\Users\David Arley\Desktop\blast-2.4.0+\db\refseq_rna.00
```

- >blastn –query “nombre de la secuencia consulta.txt” –db “documento fasta” –out “nombre deseado para el archivo de salida.txt”

```
C:\Users\David Arley\Desktop\blast-2.4.0+\db>blastn -query test_query.txt -db refseq_rna.00 -out resultado.txt
```

- Finalmente se obtienen los fragmentos del documento fasta que se alinean significativamente con la secuencia consulta en formato .txt en la carpeta creada con anterioridad.



resultado.txt

15/11/2017 10:55 ... Documento de tex...

462 KB

Al abrir este archivo .txt encontrará el blast local como se muestra a continuación:

Score = 4916 bits (2662), Expect = 0.0
 Identities = 2662/2662 (100%), Gaps = 0/2662 (0%)
 Strand=Plus/Plus

| | | |
|-----------|--|-----|
| Query 1 | GAAGAGACCCAGCAACCCACAGAGTTGAGAAATTGACTGGCATTCAAGCTGTCCAATCA | 60 |
| | | |
| Sbjct 1 | GAAGAGACCCAGCAACCCACAGAGTTGAGAAATTGACTGGCATTCAAGCTGTCCAATCA | 60 |
| Query 61 | ATAGCTGCCGCTGAAGGGTGGGCTGGATGGCGTAAGCTACAGCTGAAGGAAGAACGTGA | 120 |
| | | |
| Sbjct 61 | ATAGCTGCCGCTGAAGGGTGGGCTGGATGGCGTAAGCTACAGCTGAAGGAAGAACGTGA | 120 |
| Query 121 | GCACGAGGCACTGAGGTGATTGGCTGAAGGCACTTCCGTTGAGCATCTAGACGTTCCCTT | 180 |
| | | |
| Sbjct 121 | GCACGAGGCACTGAGGTGATTGGCTGAAGGCACTTCCGTTGAGCATCTAGACGTTCCCTT | 180 |
| Query 181 | GGCTCTTCTGGCGCCAAAATGTCGTCGTTGGCAGGGTTATTGGCGCTGGACGAGACA | 240 |
| | | |
| Sbjct 181 | GGCTCTTCTGGCGCCAAAATGTCGTCGTTGGCAGGGTTATTGGCGCTGGACGAGACA | 240 |

1.2 COMPILACIÓN DE TODOS LOS SITIOS HETEROPLÁSMICOS PARA CADA INDIVIDUO EVALUADO:

1.1. Heteroplasmia de Ei1:

| Heteroplasmia | Cobertura | Nucleótidos | Región | Longitud |
|---------------|--|-----------------------|--------------------------------------|----------|
| 1 | secuencia referencia TRINITY_DN32984_c0_g2_i1 TRINITY_DN34938_c0_g1_i1 TRINITY_DN34938_c0_g2_i1 | A A A C | (484) - rRNA-12S ribosomal RNA | 1 |
| 2 | secuencia referencia TRINITY_DN32984_c0_g2_i1 TRINITY_DN34938_c0_g1_i1 TRINITY_DN34938_c0_g2_i1 TRINITY_DN32984_c0_g1_i1 | G G G G A | (484) - rRNA-12S ribosomal RNA | 1 |
| 3 | secuencia referencia TRINITY_DN45273_c0_g1_i1 TRINITY_DN50508_c0_g1_i3 | TA TA CT | (1526-1527) - rRNA-16S ribosomal RNA | 2 |
| 4 | secuencia referencia TRINITY_DN45273_c0_g1_i1 TRINITY_DN50508_c0_g1_i3 | C C A | (2705) - rRNA-16S ribosomal RNA | 10 |
| 5 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | AC AC AC TT | (5371-5372) - tRNA-Tyr | 2 |
| 6 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | C C C A | (5374) - tRNA-Tyr | 1 |

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|----|--|---|---|----|
| 7 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | AA AA AA CT | (5376-5377) - tRNA-Tyr | 2 |
| 8 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | ACACT ACACT ACACT TGGGG | (5391-5395) - tRNA-Tyr | 5 |
| 9 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | GCCA GCCA GCCA TGAG | (5386-5389) - tRNA-Tyr | 4 |
| 10 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | CCTA CCTA CCTA AGGC | (5391-5394) - tRNA-Tyr | 4 |
| 11 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 | G G A | (7012) - tRNA-Asp | 1 |
| 12 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 | CGT CGT AAA | (7014-7016) - tRNA-Asp | 3 |
| 13 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 | AATTATAGGTG AATTATAGGTG CCGATCTCTAT | (7051-7060) - tRNA-Asp | 11 |
| 14 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 | A A T | (7064) - tRNA-Asp | 1 |
| 15 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 | CCCTT CCCTT GGTGA | (7066-7070) - tRNA-Asp | 5 |
| 16 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 | CGA CGA TTG | (7073-7075) - tRNA-Asp | 3 |
| 17 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 | T T A | (7077) - tRNA-Asp | 1 |
| 18 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 | A A C | (7079) - tRNA-Asp | 1 |
| 19 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g1_i2 | A A A T | (7730) - cytochrome c oxidase subunit II | 1 |

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|----|--|---|---|---|
| 20 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g1_i2 | A A A T | (7730) - cytochrome c oxidase subunit II | 1 |
| 21 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g1_i2 | AC AC AC GA | (7732-7733) - cytochrome c oxidase subunit II | 2 |
| 22 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g1_i2 | TTGA TTGA TTGA AGCC | (7735-7738) - cytochrome c oxidase subunit II | 4 |
| 23 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | G A A C | (7740) - cytochrome c oxidase subunit II | 1 |
| 24 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | CTGA CTGA CTGA GGCT | (7742-7745) - cytochrome c oxidase subunit II | 4 |
| 25 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | CTTC CTTC CTTC TGCA | (7747-7750) - cytochrome c oxidase subunit II | 4 |
| 26 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | TTAA CTAA CTAA GAGT | (7752-7755) - cytochrome c oxidase subunit II | 4 |
| 27 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | T T T C | (7759) - cytochrome c oxidase subunit II | 1 |
| 28 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | TCCTAACAC TCTAACAC TCTAACAC ATAGTTGT | (7761-7769) - cytochrome c oxidase subunit II; tRNA-Lys | 9 |
| 29 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | ATAGAA ATAGAG ATAGAA TCTGG | (7771-7776) - tRNA-Lys | 6 |
| 30 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | C C C A | (7778) - tRNA-Lys | 1 |

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|----|--|----------------------------------|------------------------|---|
| 31 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | AA AA AA TG | (7780-7781) - tRNA-Lys | 2 |
| 32 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | G G G T | (7785) - tRNA-Lys | 1 |
| 33 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | A A A C | (7787) - tRNA-Lys | 1 |
| 34 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | AG AG AG GC | (7789-7790) - tRNA-Lys | 2 |
| 35 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | GCTA GCTA GCTA AGAG | (7792-7795) - tRNA-Lys | 4 |
| 36 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | CCTT CCTT CCTT GAAAG | (7797-7801) - tRNA-Lys | 5 |
| 37 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | A A A C | (7803) - tRNA-Lys | 1 |
| 38 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | GCTAG GCTAG GCTAG AAATT | (7805-7809) - tRNA-Lys | 5 |
| 39 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | AGAG AGAG AGAG CCTT | (7814-7817) - tRNA-Lys | 4 |
| 40 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | AC AC AC TT | (7819-7820) - tRNA-Lys | 2 |
| 41 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | CC CC CC AA | (7822-7824) - tRNA-Lys | 2 |

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|----|--|--|---|---|
| 42 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | A A A T | (7826) - tRNA-Lys | 1 |
| 43 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | TCTCC CCTCC CCTCC TGATA | (7828-7832) - tRNA-Lys | 5 |
| 44 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | G G G C | (7836) - tRNA-Lys | 1 |
| 45 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | GA GA GA TC | (7838-7839) - tRNA-Lys | 2 |
| 46 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | T C C A | (7846) - ATP synthase F0 subunit 8 | 1 |
| 47 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 | C C C A | (7855) - ATP synthase F0 subunit 8 | 1 |
| 48 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | CTT CCT CCT CCT TAA | (7876-7878) - ATP synthase F0 subunit 8 | 3 |
| 49 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | CTCTCC CTCTCC CTCTCC CTCTCC AAGGAA | (7880-7885) - ATP synthase F0 subunit 8 | 6 |
| 50 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | CATG CATG CATG CATG GGAA | (7887-7890) - ATP synthase F0 subunit 8 | 4 |
| 51 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | CTAATTAA CTAATTAA CTAATTAA CTAATTAA GATTACAT | (7892-7899) - ATP synthase F0 subunit 8 | 8 |

| | | | | |
|----|--|---|---|----|
| 52 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | CC CC CC CC AA | (7902-7903) - ATP synthase F0 subunit 8 | 2 |
| 53 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | TTATTT TCATCTT TCATCTT TCATCTT AGGAAAC | (7905-7911) - ATP synthase F0 subunit 8 | 7 |
| 54 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | C C C C A | (7913) - ATP synthase F0 subunit 8 | 1 |
| 55 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | CCA CCA CCA CCA AAC | (7916-7918) - ATP synthase F0 subunit 8 | 3 |
| 56 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN45232_c0_g1_i1 TRINITY_DN50508_c0_g1_i1 TRINITY_DN50508_c0_g4_i1 | ATCTCATCTTA ATTACATCTTA ATTACATCTTA ATTACATCTTA CAGTATGGAGC | (7922-7932) - ATP synthase F0 subunit 8 | 11 |
| 57 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | TGGCCC TGGCCC AAAAAA AAAAAA | (8680-8685) - cytochrome c oxidase subunit III | 6 |
| 58 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | C C A A | (8687) - cytochrome c oxidase subunit III | 1 |
| 59 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | T C A A | (8692) - cytochrome c oxidase subunit III | 1 |
| 60 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | CGCCT CGCCT AAAAAA AAAAAA | (8696-8700) - cytochrome c oxidase subunit III | 1 |
| 61 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | CATAGT ATATAG CACAAAC CACAAAC | (8705-8710) - cytochrome c oxidase subunit III | 6 |

| | | | | |
|----|--|---|--|---|
| 62 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | GACC TGAT TACA TACA | (8712-8715) - cytochrome c oxidase subunit III | 4 |
| 63 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | AG AA TG TG | (8718-8719) - cytochrome c oxidase subunit III | 2 |
| 64 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | TGA CAT TGA TGA | (8724-8726) - cytochrome c oxidase subunit III | 3 |
| 65 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | CACTAAC ACCATTAA TTTGGGGG TTTGGGGG | (8728-8735) - cytochrome c oxidase subunit III | 8 |
| 66 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | G G C C | (8738) - cytochrome c oxidase subunit III | 1 |
| 67 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | AGC GCA AAG AAG | (8741-8743) - cytochrome c oxidase subunit III | 3 |
| 68 | secuencia referencia TRINITY_DN50508_c0_g1_i3 TRINITY_DN50508_c0_g5_i1 TRINITY_DN50508_c0_g4_i1 | GCA CAG AGT AGT | (8745-8747) - cytochrome c oxidase subunit III | 3 |
| 69 | secuencia referencia TRINITY_DN4102_c0_g2_i1 TRINITY_DN4102_c0_g1_i1 TRINITY_DN30895_c0_g1_i1 | A G A A | (9646) - NADH dehydrogenase subunit 3 | 1 |
| 70 | secuencia referencia TRINITY_DN4102_c0_g2_i1 TRINITY_DN4102_c0_g1_i1 TRINITY_DN30895_c0_g1_i1 | A T A A | (9665) - NADH dehydrogenase subunit 3 | 1 |
| 71 | secuencia referencia TRINITY_DN4102_c0_g2_i1 TRINITY_DN4102_c0_g1_i1 TRINITY_DN30895_c0_g1_i1 TRINITY_DN37761_c0_g1_i1 | GGC AAA AAA GGT GGT | (9882-9884) - tRNA-Arg | 3 |

| | | | | |
|----|--|--------------------------------------|---|---|
| 72 | secuencia referencia TRINITY_DN4102_c0_g2_i1 TRINITY_DN4102_c0_g1_i1 TRINITY_DN30895_c0_g1_i1 TRINITY_DN37761_c0_g1_i1 | CT AA AA CT CT | (9887-9888) - tRNA-Arg | 2 |
| 73 | secuencia referencia TRINITY_DN30895_c0_g1_i1 TRINITY_DN37761_c0_g1_i1 | GTT AAA GTT | (11622-11624) - tRNA-His | 3 |
| 74 | secuencia referencia TRINITY_DN30895_c0_g1_i1 TRINITY_DN37761_c0_g1_i1 | T A T | (11627) - tRNA-His | 1 |
| 75 | secuencia referencia TRINITY_DN30895_c0_g1_i1 TRINITY_DN37761_c0_g1_i1 | T A T | (11629) - tRNA-His | 1 |
| 76 | secuencia referencia TRINITY_DN34579_c0_g1_i1 TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | A A A A G | (14076) - NADH dehydrogenase subunit 6 | 1 |
| 77 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | TG AA TG TG | (14156-14157) - tRNA-Glu | 2 |
| 78 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | TC AC TT TT | (14161-14162) -tRNA-Glu | 2 |
| 79 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | AGACC TCCAA AGACC AGACC | (14169-14173) -tRNA-Glu | 5 |
| 80 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | TGATTG AACAAA TGATTG TGATTG | (14176-14186) -tRNA-Glu | 8 |
| 81 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | ACCG CAAA ACCG ACCG | (14190-14193) -tRNA-Glu | 4 |
| 82 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | TGT AAA TGT TGT | (14195-14197) -tRNA-Glu | 3 |

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|----|--|--|------------------------------|---|
| 83 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | T A T T | (14199) -tRNA-Glu | 1 |
| 84 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | CA GC CA CA | (14201-14202) -tRNA-Glu | 2 |
| 85 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | CT AA CT CT | (14204-14205) -tRNA-Glu | 2 |
| 86 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | C T C C | (14207) -tRNA-Glu | 1 |
| 87 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | A C A A | (14209) -tRNA-Glu | 1 |
| 88 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | A T A A | (14211) -tRNA-Glu | 1 |
| 89 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | C A T T | (14214) - Zona intergénica | 1 |
| 90 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | GGCCA TTTCG GGCTA GGCTA | (14219-14223) - cytochrome b | 5 |
| 91 | secuencia referencia TRINITY_DN40849_c0_g1_i1 TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | A C A A | (14225) - cytochrome b | 1 |
| 92 | secuencia referencia TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 | G G T | (14791) - cytochrome b | 5 |
| 93 | secuencia referencia TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 TRINITY_DN28150_c0_g2_i1 TRINITY_DN28150_c0_g3_i1 TRINITY_DN28150_c0_g1_i1 | TTCT AAAA AAAA TTCT TTCT TTCT | (15362-15365) - tRNA-Thr | 4 |

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|-------|--|----------------------------------|-----------------------------|-----|
| 94 | secuencia referencia TRINITY_DN40849_c0_g1_i2 TRINITY_DN38087_c0_g1_i1 TRINITY_DN28150_c0_g2_i1 TRINITY_DN28150_c0_g3_i1 TRINITY_DN28150_c0_g1_i1 | GT A- AA GT GT GT | (15367-15368) - tRNA-Thr | 2 |
| 95 | secuencia referencia TRINITY_DN38087_c0_g1_i1 TRINITY_DN28150_c0_g2_i1 TRINITY_DN28150_c0_g3_i1 TRINITY_DN28150_c0_g1_i1 | GCT AAA GCT GCT GCT | (15370-15372) - tRNA-Thr | 3 |
| TOTAL | | | | 286 |

1.2. Heteroplasmia de Cc1

| Heteroplasmia | Cobertura | nucleótidos | Región | Longitud |
|---------------|--|--|--|----------|
| 1 | secuencia referencia TRINITY_DN17717_c0_g1_i1 TRINITY_DN25688_c0_g1_i1 | C G C | (1123) - rRNA-16S ribosomal RNA | 1 |
| 2 | secuencia referencia TRINITY_DN17717_c0_g1_i1 TRINITY_DN25688_c0_g1_i1 | A T A | (1125) - rRNA-16S ribosomal RNA | 1 |
| 3 | secuencia referencia TRINITY_DN17717_c0_g1_i1 TRINITY_DN25688_c0_g1_i1 | C T C | (1127) - rRNA-16S ribosomal RNA | 1 |
| 4 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | ACCTCAACAA ACCTCAACAA GAGATTGT ACCTCAACAA | (4604-4613) - NADH dehydrogenase subunit 2 | 10 |
| 5 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | AT AT TG AT | (4615-4616) - NADH dehydrogenase subunit 2 | 2 |
| 6 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | A A T A | (4621) - NADH dehydrogenase subunit 2 | 1 |
| 7 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | T T G T | (4623) - NADH dehydrogenase subunit 2 | 1 |

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|----|--|--|---|----|
| 8 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | TAAAAC TAAAAC GTTTTA TAAAAC | (4626-4631) - NADH dehydrogenase subunit 2 | 6 |
| 9 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | C C A C | (4634) - NADH dehydrogenase subunit 2 | 1 |
| 10 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | A A T A | (4636) - NADH dehydrogenase subunit 2 | 1 |
| 11 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | CA CA AT CA | (4641-4642) - NADH dehydrogenase subunit 2 | 2 |
| 12 | secuencia referencia TRINITY_DN68855_c0_g1_i1 TRINITY_DN13494_c0_g1_i1 TRINITY_DN13494_c0_g2_i1 | ACAAAATC ACAAAATC TTGTTGAG ACAAAATC | (4644-4651) - NADH dehydrogenase subunit 2 | 8 |
| 13 | secuencia referencia TRINITY_DN13494_c0_g2_i1 TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 | A A G G | (5074) - tRNA-Trp | 1 |
| 14 | secuencia referencia TRINITY_DN13494_c0_g2_i1 TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | A A A A C | (5370) - tRNA-Tyr | 1 |
| 15 | secuencia referencia TRINITY_DN13494_c0_g2_i1 TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | A C A A A | (5389) - tRNA-Tyr | 1 |
| 16 | secuencia referencia TRINITY_DN13494_c0_g2_i1 TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | CCTGTGTTTT AAAAAA CCTGTGTTTT CCTGTGTTTT CCTGTGTTTT | (5395-5405) - tRNA-Tyr; cytochrome c oxidase subunit I | 11 |

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|----|--|--------------------------------------|--|---|
| 17 | secuencia referencia TRINITY_DN13494_c0_g2_i1 TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | CCG AAA CCG CCG CCG | (5409-5411) - cytochrome c oxidase subunit I | 3 |
| 18 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | G T G G | (6535) - cytochrome c oxidase subunit I | 1 |
| 19 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | G T G G | (6548) - cytochrome c oxidase subunit I | 1 |
| 19 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | GCCGT AAAAG GCCGT GCCGT | (6550-6554) - cytochrome c oxidase subunit I | 5 |
| 20 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | TTTGCT AAGATA TTTGCT TTTGCT | (6556-6561) - cytochrome c oxidase subunit I | 6 |
| 21 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | T A T T | (6563) - cytochrome c oxidase subunit I | 1 |
| 22 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | A C A A | (6565) - cytochrome c oxidase subunit I | 1 |
| 23 | secuencia referencia TRINITY_DN25186_c0_g1_i2 TRINITY_DN25186_c0_g1_i1 TRINITY_DN22337_c0_g1_i1 | CAGGAT GGCTCA CAGGAT CAGGAT | (6569-6574) - cytochrome c oxidase subunit I | 6 |
| 24 | secuencia referencia TRINITY_DN26422_c0_g1_i1 TRINITY_DN22108_c0_g1_i1 | TGAGG AAGAT TGAGG | (9453-9457) - cytochrome c oxidase subunit III | 5 |
| 25 | secuencia referencia TRINITY_DN26422_c0_g1_i1 TRINITY_DN22108_c0_g1_i1 | TC GG TC | (9459-9460) - cytochrome c oxidase subunit III | 2 |

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| 26 | secuencia referencia TRINITY_DN26422_c0_g1_i1 TRINITY_DN22108_c0_g1_i1 TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 | GCTCCCCTAG GA----- AAAAAAAAGA - CTCCCCTAG - CTCCCCTAG | (9463-9472) - tRNA-Gly | 10 |
| 27 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | TTAC TTAC TTAC TTAC AGTA | (9817-9820) - NADH dehydrogenase subunit 3 | 4 |
| 28 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | CTCC CTCC CTCC CTCC GGGG | (9823-9826) - NADH dehydrogenase subunit 3 | 4 |
| 29 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | CAC CAC CAC CAC TTA | (9830-9832) - NADH dehydrogenase subunit 3 | 3 |
| 30 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | G G G G A | (9835) - NADH dehydrogenase subunit 3 | 1 |
| 31 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | CCT CCT CCT CCT TGA | (9837-9839) - NADH dehydrogenase subunit 3 | 3 |
| 32 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | ATTTA ATCTA ATCTA ATCTA GGGT | (9841-9845) - NADH dehydrogenase subunit 3 | 5 |
| 33 | secuencia referencia TRINITY_DN13439_c0_g1_i1 TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | GAATGAATCCAAGGAGGCTTAGAAT GAATGAGGCCACCCAACTCCCATCTC GAATGAGTCCAAGGAGGCTTAGAAT GAATGAGTCCAAGGAGGCTTAGAAT TGTAATTGGGTGGTCATATTGCTA | (9847-9871) - NADH dehydrogenase subunit 3 | 25 |
| 34 | secuencia referencia TRINITY_DN13439_c0_g2_i1 TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | GGC AAA GGC GGG | (9882-9884) - tRNA-Arg | 3 |

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| 35 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | AG GG AT | (9889-9890) - tRNA-Arg | 2 |
| 36 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | CTAAC CTAAC TGCTT | (9892-9897) - tRNA-Arg | 6 |
| 37 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | A A T | (9899) - tRNA-Arg | 1 |
| 38 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | ACTA ACTA TAGT | (9905-9908) - tRNA-Arg | 4 |
| 39 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 | TTTCGACTTAGTTAAC TTTCGACTTAGTTAAC GAGTTGAGATTAAAGTGGG | (9910-9927) - tRNA-Arg | 18 |
| 40 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | GATTAAACT GATTAAACT TCGATGTAA ----GATGTAA | (9929-9937) - tRNA-Arg | 9 |
| 41 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | C C G G | (9939) - tRNA-Arg | 1 |
| 42 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | TG TG GA GA | (9941-9942) - tRNA-Arg | 2 |
| 42 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | TTTCCC TTTCCC GCCTAT GCCTAT | (9949-9954) - tRNA-Arg; NADH dehydrogenase subunit 4L | 6 |
| 43 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | AC AC CT AC | (9954-9955) - NADH dehydrogenase subunit 4L | 2 |
| 44 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | CC CC AT CC | (9957-9958) - NADH dehydrogenase subunit 4L | 2 |
| 45 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | T T A T | (9960) - NADH dehydrogenase subunit 4L | 1 |

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| 46 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | C C A C | (9963) - NADH dehydrogenase subunit 4L | 1 |
| 47 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | CTTCAGC CTTCAGC GGCGGAG CTTCAGC | (9965-9971) - NADH dehydrogenase subunit 4L | 7 |
| 48 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | ATC ACC GGT ACC | (9973-9975) - NADH dehydrogenase subunit 4L | 3 |
| 49 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | CTCCGCC CTCCGCC GCTGAAG CTCCGCC | (9977-9983) - NADH dehydrogenase subunit 4L | 7 |
| 50 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | T T G T | (9985) - NADH dehydrogenase subunit 4L | 1 |
| 51 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | T T A T | (9988) - NADH dehydrogenase subunit 4L | 1 |
| 52 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | AT AT GG AT | (9990-9991) - NADH dehydrogenase subunit 4L | 1 |
| 53 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | AG AG GT AG | (9993-9994) - NADH dehydrogenase subunit 4L | 2 |
| 54 | secuencia referencia TRINITY_DN21587_c0_g1_i1 TRINITY_DN24818_c0_g3_i1 | TGGCCAC AAAAACA TGGCCAC | (14218-14224) - cytochrome b | 7 |
| 55 | secuencia referencia TRINITY_DN21587_c0_g1_i1 TRINITY_DN24818_c0_g3_i1 | CT AA CT | (14229-14230) - cytochrome b | 2 |
| 56 | secuencia referencia TRINITY_DN21587_c0_g1_i1 TRINITY_DN24818_c0_g3_i1 TRINITY_DN78593_c0_g1_i1 | AT AT GG AT | (14229-14230) - cytochrome b | 1 |

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|----|--|--|---------------------------------|---|
| 57 | secuencia referencia TRINITY_DN21587_c0_g1_i1 TRINITY_DN24818_c0_g3_i1 TRINITY_DN78593_c0_g1_i1 | CGAA ACCC CGAA -GAA | (14232-14235) - cytochrome b | 4 |
| 58 | secuencia referencia TRINITY_DN24442_c0_g1_i1 TRINITY_DN24442_c0_g1_i4 TRINITY_DN24442_c0_g1_i3 | C T C C | (14239) - cytochrome b | 1 |
| 59 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | GCTGACC GCTGATC TTCTCCA GCTGATC | (15207-15213) - cytochrome b | 7 |
| 60 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | A A C A | (15215) - cytochrome b | 1 |
| 61 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | C C T C | (15222) - cytochrome b | 1 |
| 62 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | AC AC TT AC | (15225-15226) - cytochrome b | 2 |
| 63 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | TG TG CT TG | (15228-15229) - cytochrome b | 2 |
| 64 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | TGGAGG TGGAGG ACCCAC TGGAGG | (15233-15238) - cytochrome b | 6 |
| 65 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | CA CA TT CA | (15240-15241) - cytochrome b | 2 |
| 66 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | CAGTCGA CAGTCGA ACACATC CAGTCGA | (15244-15250) - cytochrome b | 7 |
| 67 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | G G A G | (15252) - cytochrome b | 1 |

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|----|--|----------------------------------|---------------------------------|---|
| 68 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | T T A T | (15254) - cytochrome b | 1 |
| 69 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | C C A C | (15256) - cytochrome b | 1 |
| 70 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | TTC TTC CGA TTC | (15258-15260) - cytochrome b | 3 |
| 71 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | TTA TTA CAG TTA | (15262-15264) - cytochrome b | 3 |
| 72 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | ATGGT ATGGT TCATT ATGGT | (15267-15272) - cytochrome b | 6 |
| 73 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | A A G A | (15274) - cytochrome b | 1 |
| 74 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | AT AT CC AT | (15276-15277) - cytochrome b | 2 |
| 75 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | GC GC CT GC | (15279-15280) - cytochrome b | 2 |
| 76 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | T T A T | (15282) - cytochrome b | 1 |
| 77 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | TAGCC TAGCC CCAAA TAGCC | (15284-15288) - cytochrome b | 5 |
| 78 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | T T C T | (15290) - cytochrome b | 1 |

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| 79 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | AC AC TA AC | (15292-15293) - cytochrome b | 2 |
| 80 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | T T G T | (15298) - cytochrome b | 1 |
| 81 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | AT AT TC AT | (15300-15301) - cytochrome b | 2 |
| 82 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | T T C T | (15303) - cytochrome b | 1 |
| 83 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | CTA GAT CTA | (15413-15415) - tRNA-Thr | 3 |
| 84 | secuencia referencia TRINITY_DN78593_c0_g1_i1 TRINITY_DN24818_c0_g2_i1 | AACT GGAA AACT | (15417-15420) - tRNA-Thr | 4 |
| 85 | secuencia referencia TRINITY_DN24818_c0_g2_i1 TRINITY_DN40817_c0_g1_i1 | G G T | (15445) - tRNA-Pro | 1 |
| TOTAL | | | | 299 |

1.3. Heteroplasmia Cc2:

| Heteroplasmia | Cobertura | Nucleótidos | Región | Longitud |
|---------------|--|----------------------|---------------------------------------|----------|
| 1 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | C C C T | (3551) - NADH dehydrogenase subunit 1 | 1 |
| 2 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | GG AA GG AA | (3764-3765) - tRNA-Ile | 2 |

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| 3 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | C A C A | (3767) - tRNA-Ile | 1 |
| 4 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | CGTGC AAAAAA CGTGC AAAAAA | (3769-3774) - tRNA-Ile | 6 |
| 5 | secuencia referencia TRINITY_DN28027_c0_g2_i1 TRINITY_DN26863_c0_g1_i2 TRINITY_DN27634_c0_g1_i1 | T T T A | (3934) - tRNA-Met | 1 |
| 6 | secuencia referencia TRINITY_DN28027_c0_g2_i1 TRINITY_DN26863_c0_g1_i2 TRINITY_DN27634_c0_g1_i1 | C C C T | (3936) - tRNA-Met | 1 |
| 7 | secuencia referencia TRINITY_DN28027_c0_g2_i1 TRINITY_DN26863_c0_g1_i2 TRINITY_DN27634_c0_g1_i1 | CCGAA CCGAA CCGAA AAACC | (3938-3942) - tRNA-Met | 5 |
| 8 | secuencia referencia TRINITY_DN28027_c0_g2_i1 TRINITY_DN26863_c0_g1_i2 TRINITY_DN27634_c0_g1_i1 | TGT TGT TGT ACG | (3945-3947) - tRNA-Met | 3 |
| 9 | secuencia referencia TRINITY_DN28027_c0_g2_i1 TRINITY_DN26863_c0_g1_i2 TRINITY_DN27634_c0_g1_i1 | GGTT GGTT GGTT CACCA | (3949-3952) - tRNA-Met | 4 |
| 10 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | A A A C | (3954) - tRNA-Met | 1 |
| 11 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | ATCCCTCCT ATCCCTCCT ATCCCTCCT CGATACAAA | (3956-3964) - tRNA-Met | 9 |
| 12 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | TACT TACT TACT GGAG | (3966-3969) - tRNA-Met | 4 |
| 13 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | AT AT AT CA | (3971-3972) - NADH dehydrogenas e subunit 2 | 2 |

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| 14 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | A A A G | (3975) - NADH dehydrogenase subunit 2 | 1 |
| 15 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | CT CC CC AA | (3979-3980) - NADH dehydrogenase subunit 2 | 2 |
| 16 | secuencia referencia TRINITY_DN26863_c0_g1_i2 TRINITY_DN27634_c0_g1_i1 TRINITY_DN26863_c0_g1_i1 | GCTAA GCTAA GCTAA CGATC | (5373-5377) - tRNA-Tyr | 5 |
| 17 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | ACACT ACACT ACACT CGGGG | (5379-5383) - tRNA-Tyr | 5 |
| 18 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | GCCA GCCA GCCA TGAG | (5386-5389) - tRNA-Tyr | 4 |
| 19 | secuencia referencia TRINITY_DN28115_c0_g1_i1 TRINITY_DN28027_c0_g2_i1 TRINITY_DN28115_c0_g2_i1 | CCTA CCTA CCTA AGGC | (5391-5394) - tRNA-Tyr | 4 |
| 20 | secuencia referencia TRINITY_DN27634_c0_g1_i1 TRINITY_DN16111_c0_g1_i1 | A A G | (7776) - tRNA-Lys | 1 |
| 21 | secuencia referencia TRINITY_DN27634_c0_g1_i1 TRINITY_DN16111_c0_g1_i1 | A A G | (7781) - tRNA-Lys | 1 |
| 22 | secuencia referencia TRINITY_DN27634_c0_g1_i1 TRINITY_DN16111_c0_g1_i1 | T T C | (8012) - ATP synthase F0 subunit 6 | 1 |
| 23 | secuencia referencia TRINITY_DN27634_c0_g1_i1 TRINITY_DN27634_c0_g2_i1 | A A C | (8780) - cytochrome c oxidase subunit III | 1 |
| 24 | secuencia referencia TRINITY_DN27634_c0_g1_i1 TRINITY_DN27634_c0_g2_i1 | C C A | (8799) - cytochrome c oxidase subunit III | 1 |

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| 25 | secuencia referencia TRINITY_DN27634_c0_g1_i1 TRINITY_DN27634_c0_g2_i1 TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 | A A A C A | (9471) - tRNA-Gly | 1 |
| 26 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AGCC AGCC AGCC TTTT | (11309- 11312) - NADH dehydrogenas e subunit 4 | 4 |
| 27 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11314) - NADH dehydrogenas e subunit 4 | 1 |
| 28 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | GA GA GA TT | (11316- 11317) - NADH dehydrogenas e subunit 4 | 2 |
| 29 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | GAC GAC GAC TTT | (11319- 11321) - NADH dehydrogenas e subunit 4 | 3 |
| 30 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AC AC AC TT | (11323- 11324) - NADH dehydrogenas e subunit 4 | 2 |
| 31 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | GC GC GC TT | (11327- 11328) - NADH dehydrogenas e subunit 4 | 2 |
| 32 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | T T T G | (11332) - NADH dehydrogenas e subunit 4 | 1 |
| 33 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11335) - NADH dehydrogenas e subunit 4 | 1 |
| 34 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AAC AAC AAC TGT | (11339- 11341) - NADH | 3 |

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| | | | dehydrogenas e subunit 4 | |
| 35 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TAGCCC TAGCCC TAGCCC GTTGAA | (11343- 11348) - NADH dehydrogenas e subunit 4 | 6 |
| 36 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | T T T A | (11350) - NADH dehydrogenas e subunit 4 | 1 |
| 37 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C A | (11352) - NADH dehydrogenas e subunit 4 | 1 |
| 38 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CAACCA CAACCA CAACCA GGTGGT | (11355- 11360) - NADH dehydrogenas e subunit 4 | 6 |
| 39 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AA AA AA TT | (11363- 11364) - NADH dehydrogenas e subunit 4 | 2 |
| 40 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CT CT CT AA | (11366- 11367) - NADH dehydrogenas e subunit 4 | 2 |
| 41 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AT AT AT TC | (11369- 11370) - NADH dehydrogenas e subunit 4 | 2 |
| 42 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | GGA GGA GGA TTG | (11372- 11374) - NADH dehydrogenas e subunit 4 | 3 |
| 43 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AA AA AA TC | (11376- 11377) - NADH | 2 |

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| | | | dehydrogenas e subunit 4 | |
| 44 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AACC AACC AACC GGTT | (11380- 11383) - NADH dehydrogenas e subunit 4 | 4 |
| 45 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TTA TTA TTA GAG | (11385- 11387) - NADH dehydrogenas e subunit 4 | 3 |
| 46 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TACTT TACTT TACTT CCAGG | (11389- 11393) - NADH dehydrogenas e subunit 4 | 5 |
| 47 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CT CT CT AG | (11396- 11397) - NADH dehydrogenas e subunit 4 | 2 |
| 48 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | T T T A | (11399) - NADH dehydrogenas e subunit 4 | 1 |
| 49 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CC CC CC GT | (11409- 11410) - NADH dehydrogenas e subunit 4 | 2 |
| 50 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AC AC AC TT | (11412- 11413) - NADH dehydrogenas e subunit 4 | 2 |
| 51 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A G | (11417) - NADH dehydrogenas e subunit 4 | 1 |
| 52 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11420) - NADH dehydrogenas e subunit 4 | 1 |

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| 53 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CC CC CC TT | (11422- 11423) - NADH dehydrogenas e subunit 4 | 2 |
| 54 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11426) - NADH dehydrogenas e subunit 4 | 1 |
| 55 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AACAGGA AACAGGA AACAGGA TGGGTTT | (11428- 11434) - NADH dehydrogenas e subunit 4 | 7 |
| 56 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TAGGAACCC TAGGAACCC TAGGAACCC GTTTTGTTT | (11436- 11444) - NADH dehydrogenas e subunit 4 | 9 |
| 57 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AATC AATC AATC TTGG | (11446- 11449) - NADH dehydrogenas e subunit 4 | 4 |
| 58 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CC CC CC TG | (11451- 11452) - NADH dehydrogenas e subunit 4 | 2 |
| 59 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C T | (11454) - NADH dehydrogenas e subunit 4 | 1 |
| 60 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AC AC AC GG | (11456- 11457) - NADH dehydrogenas e subunit 4 | 2 |
| 61 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TACA TACA TACA GTGT | (11459- 11462) - NADH dehydrogenas e subunit 4 | 4 |

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| 62 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CCTAT CCTAT CCTAT TTGTA | (11464- 11468) - NADH dehydrogenas e subunit 4 | 5 |
| 63 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ATA ATA ATA CCG | (11471- 11473) - NADH dehydrogenas e subunit 4 | 3 |
| 64 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TAT TAT TAT CCC | (11475- 11477) - NADH dehydrogenas e subunit 4 | 3 |
| 65 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TACAACACA TACAACACA TACAACACA ATATTATGG | (11479- 11487) - NADH dehydrogenas e subunit 4 | 9 |
| 66 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TGAGGGGAAACACC TGAGGGGAAACACC TGAGGGGAAACACC GTTTTAAGTTGGT | (11489- 11502) - NADH dehydrogenas e subunit 4 | 14 |
| 67 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TCATACATCAAAAC TCATACATCAAAAC TCATACATCAAAAC CTTGGGGCGGCTT | (11504- 11517) - NADH dehydrogenas e subunit 4 | 14 |
| 68 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ATCCCCCAACCCACAC AC ATCCCCCAACCCACAC AC ATCCCCCAACCCACAC AC GGTTGTGGTTGTTAGT TA | (11519- 11537) - NADH dehydrogenas e subunit 4 | 19 |
| 69 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AGA AGA AGA TAT | (11539- 11541) - NADH dehydrogenas e subunit 4 | 3 |

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| 70 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CATCTCTTAA CATCTCCTAA CATCTCCTAA GGGGGGTCTT | (11543- 11552) - NADH dehydrogenas e subunit 4 | 10 |
| 71 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ATCA ATCA ATCA TGTT | (11554- 11557) - NADH dehydrogenas e subunit 4 | 4 |
| 72 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TACACA TACACA TACACA CTGTAG | (11559- 11564) - NADH dehydrogenas e subunit 4 | 6 |
| 73 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CC CC CC AT | (11566- 11567) - NADH dehydrogenas e subunit 4 | 2 |
| 74 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ACCAA ACCAA ACCAA GTTTT | (11569- 11573) - NADH dehydrogenas e subunit 4 | 5 |
| 75 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C T | (11582) - NADH dehydrogenas e subunit 4 | 1 |
| 76 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11584) - NADH dehydrogenas e subunit 4 | 1 |
| 77 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TAACAA TAACAA TAACAA GGTGGC | (11586- 11591) - NADH dehydrogenas e subunit 4 | 6 |
| 78 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ACCAGAACTAA ACCAGAACTAA ACCAGAACTAA TATTAGTTATT | (11593- 11603) - NADH dehydrogenas e subunit 4 | 11 |

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| 79 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C T | (11605) - NADH dehydrogenas e subunit 4 | 1 |
| 80 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AGGC AGGC AGGC CTTA | (11608- 11611) - NADH dehydrogenas e subunit 4 | 4 |
| 81 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C T | (11613) - NADH dehydrogenas e subunit 4 | 1 |
| 82 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | T T T A | (11615) - NADH dehydrogenas e subunit 4 | 1 |
| 83 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A G | (11619) - NADH dehydrogenas e subunit 4 | 1 |
| 84 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AA AA AA GG | (11625- 11626) - tRNA-His | 2 |
| 85 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AT AT AT GG | (11628- 11629) - tRNA-His | 2 |
| 86 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | GTTT GTTT GTTT AGAG | (11631- 11634) - tRNA-His | 4 |
| 87 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AAA AAA AAA CTT | (11636- 11638) - tRNA-His | 3 |
| 88 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CAAACA CAAACA CAAACA TCCTGT | (11640- 11645) - tRNA-His | 6 |
| 89 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TAGACT TAGACT TAGACT GCTTGA | (11647- 11652) - tRNA-His | 6 |

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| 90 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TGGC TGGC TGGC GTAA | (11654- 11657) - tRNA-His | 4 |
| 91 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | T T T G | (11660) - tRNA-His | 1 |
| 92 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AAAATAGG AAAATAGG AAAATAGG GGTGGGAT | (11662- 11669) - tRNA-His | 8 |
| 93 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | G G G A | (11671) - tRNA-His | 1 |
| 94 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11674) - tRNA-His | 1 |
| 95 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AA AA AA TG | (11676- 11677) - tRNA-His | 2 |
| 96 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CTCCTTA CTCCTTA CTCCTTA GGTGAGT | (11679- 11685) - tRNA-His | 7 |
| 97 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A T | (11688) - tRNA-His | 1 |
| 98 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CCGAGAGAGG CCGAGAGAGG CCGAGAGAGG TTTAGTTCT | (11690- 11699) - tRNA-His; tRNA-Ser | 10 |
| 99 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A G | (11701) - tRNA-Ser | 1 |
| 100 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AATAC AATAC AATAC GGGGT | (11703- 11707) - tRNA-Ser | 6 |

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| 101 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ATAA ATAA ATAA TGGG | (11709- 11712) - tRNA-Ser | 4 |
| 102 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A C | (11714) - tRNA-Ser | 1 |
| 103 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C A | (11716) - tRNA-Ser | 1 |
| 104 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CTAAC CTAAC CTAAC TGTAA | (11723) - tRNA-Ser | 5 |
| 105 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CCTA CCTA CCTA TGAC | (11725- 11728) - tRNA-Ser | 4 |
| 106 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A G | (11730) - tRNA-Ser | 1 |
| 107 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CTGAG CTGAG CTGAG GGAGT | (11732- 11736) - tRNA-Ser | 5 |
| 108 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TA TA TA GG | (11739- 11740) - tRNA-Ser | 6 |
| 109 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TCCC TCCC TCCC AGGG | (11742- 11745) - tRNA-Ser | 4 |
| 110 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CA CA CA AT | (11747- 11748) - tRNA-Ser | 2 |
| 111 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | C C C G | (11750) - tRNA-Ser | 1 |

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| 112 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | CCCTCACTT CCCTCACTT CCCTCACTT AGTGTACAA | (11752- 11761) - tRNA-Ser; tRNA-Leu | 10 |
| 113 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AA AA AA TG | (11763- 11764) - tRNA-Leu | 2 |
| 114 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | G G G T | (11766) - tRNA-Leu | 1 |
| 115 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A C | (11768) - tRNA-Leu | 1 |
| 116 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | A A A G | (11770) - tRNA-Leu | 1 |
| 117 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | AA AA AA TG | (11772- 11773) - tRNA-Leu | 2 |
| 118 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TAATCCAC TAATCCAC TAATCCAC GTTGTTTG | (11775- 11782) - tRNA-Leu | 8 |
| 119 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | G G G T | (11784) - tRNA-Leu | 1 |
| 120 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | TT TT TT CG | (11788- 11789) - tRNA-Leu | 2 |
| 121 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | GG GG GG TT | (11791- 11792) - tRNA-Leu | 2 |
| 122 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 | ACCACG ACCACA ACCACA GATTTC | (11794- 11799) - tRNA-Leu | 6 |

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| | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | ACCCCTTG ACCCCTTG ACCCCTTG GTGGGGT ACCCCTTG ACCCCTTG | (11801- 11807) - tRNA-Leu | 7 |
| 123 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | TGCAAT TGCAAT TGCAAT GCATGG TGCAAT TGCAAT | (11809- 11814) - tRNA-Leu | 6 |
| 124 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | CCA CCA CCA TTT CCA CCA | (11816- 11818) - tRNA-Leu | 3 |
| 125 | secuencia referencia TRINITY_DN26669_c0_g1_i1 TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | G G T G G | (11820) - tRNA-Leu | 1 |
| 126 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | AAAGTAAT AAAGTAAT TTTTGGCA AAAGTAAT AAAGTAAT | (11823- 11830) - tRNA-Leu; NADH dehydrogenas e subunit 5 | 8 |
| 127 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | ACCACAC ACCACAC TTGGTTG ACCACAC ACCACAC | (11832- 11838) - NADH dehydrogenas e subunit 5 | 7 |
| 128 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | AA AA TT AA AA | (11840- 11841) - NADH dehydrogenas e subunit 5 | 2 |
| 129 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | A A T A A | (11843) - NADH dehydrogenas e subunit 5 | 1 |
| 130 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 | | | |

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| | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CA CA GC CA CA CA | (11848- 11849) - NADH dehydrogenas e subunit 5 | 2 |
| 131 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CC CC GG CC CC CC | (11851- 11852) - NADH dehydrogenas e subunit 5 | 2 |
| 132 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCTCT CC----- GAGGG CCTCT CCTCT CCTCT | (11853- 11859) - NADH dehydrogenas e subunit 5 | 5 |
| 133 | secuencia referencia TRINITY_DN26669_c0_g1_i2 TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AGCCC TATTG AGCCC AGCCC AGCCC | (11861- 11865) - NADH dehydrogenas e subunit 5 | 5 |
| 134 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AAT TGA AAT AAT AAT | (11867- 11869) - NADH dehydrogenas e subunit 5 | 3 |
| 135 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ACCC GTAA ACCC ACCC ACCC | (11871- 11874) - NADH dehydrogenas e subunit 5 | 4 |
| 136 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AATATT TTGTTG AATACT AATACT AATACT | (11876- 11881) - NADH dehydrogenas e subunit 5 | 6 |
| 137 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCACT GATGG CCACT CCACT CCACT | (11883- 11887) - NADH dehydrogenas e subunit 5 | 5 |
| 138 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | | | |

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| 139 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ACAAACAA GTTGGTT ACAAACAA ACAAACAA ACAAACAA | (11889- 11895) - NADH dehydrogenas e subunit 5 | 7 |
| 140 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T G T T T | (11897) - NADH dehydrogenas e subunit 5 | 1 |
| 141 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCCA GTTT CCCA CCCA CCCA | (11900- 11903) - NADH dehydrogenas e subunit 5 | 4 |
| 142 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AAA TCT AAA AAA AAA | (11907) - NADH dehydrogenas e subunit 5 | 3 |
| 143 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A G A A A | (11909) - NADH dehydrogenas e subunit 5 | 1 |
| 144 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AACAC GGGTA AACAC AACAC AACAC | (11912- 11916) - NADH dehydrogenas e subunit 5 | 5 |
| 145 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTAAAAACAA TCGTCCGATT ATAAAAACAA ATAAAAACAA ATAAAAACAA | (11919- 11928) - NADH dehydrogenas e subunit 5 | 10 |
| 146 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AACAGCTGT GGTGATGAG AACAGCTGT AACAGCTGT AACAGCTGT | (11930- 11938) - NADH dehydrogenas e subunit 5 | 9 |
| 147 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AA GT AA AA AA | (11941- 11942) - NADH dehydrogenas e subunit 5 | 2 |

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| 148 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TAGCAT GTAATG TAGCAT TAGCAT TAGCAT | (11944- 11949) - NADH dehydrogenas e subunit 5 | 6 |
| 149 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CT TA CT CT CT | (11951- 11952) - NADH dehydrogenas e subunit 5 | 2 |
| 150 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAT TGA CAT CAT CAT | (11954- 11956) - NADH dehydrogenas e subunit 5 | 3 |
| 151 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCC GGA CCC CCC CCC | (11959- 11961) - NADH dehydrogenas e subunit 5 | 3 |
| 152 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A G A A A | (11963) - NADH dehydrogenas e subunit 5 | 1 |
| 153 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TCCC ATGA TCCC TCCC TCCC | (11965- 11968) - NADH dehydrogenas e subunit 5 | 4 |
| 154 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C C | (11970) - NADH dehydrogenas e subunit 5 | 1 |
| 155 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A T A A A | (11972) - NADH dehydrogenas e subunit 5 | 1 |
| 156 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CGCCT ATAGA CACCT CACCT CACCT | (11975- 11979) - NADH dehydrogenas e subunit 5 | 5 |

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| 157 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CA TT CA CA CA | (11981- 11982) - NADH dehydrogenas e subunit 5 | 2 |
| 158 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TTAT AATG TTAT TTAT TTAT | (11984- 11987) - NADH dehydrogenas e subunit 5 | 4 |
| 159 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAGACA GGCCTT CAGACA CAGACA CAGACA | (11989- 11994) - NADH dehydrogenas e subunit 5 | 6 |
| 160 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GAATC TGTGA GAATC GAATC GAATC | (11997- 12001) - NADH dehydrogenas e subunit 5 | 5 |
| 161 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TTATC CGGAT TTATC TTATC TTATC | (12004- 12008) - NADH dehydrogenas e subunit 5 | 5 |
| 162 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCAACC AAGTTG CCAACC CCAACC CCAACC | (12010- 12015) - NADH dehydrogenas e subunit 5 | 6 |
| 163 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CACT ATTG CACT CACT CACT | (12018- 12021) - NADH dehydrogenas e subunit 5 | 4 |
| 164 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ATCA ATAG ATCA ATCA ATCA | (12023- 12026) - NADH dehydrogenas e subunit 5 | 4 |
| 165 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCACATC TTTGTA CCACTAA CCACATC CCACATC | (12028- 12034) - NADH dehydrogenas e subunit 5 | 7 |

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| 166 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CACAT TTTGC CACCT CACAT CACAT | (12035- 12039) - NADH dehydrogenas e subunit 5 | 5 |
| 167 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C T C C C | (12041) - NADH dehydrogenas e subunit 5 | 1 |
| 168 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCATAAACAT GTCCTTTGG TT-TATACAG CCATAAACAT CCATAAACAT | (12043- 12052) - NADH dehydrogenas e subunit 5 | 10 |
| 169 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AGCTTTAACTTG CCAACGTTTCAT CATTGAATCTGTT AGCTTTAACTTG AGCTTTAACTTG | (12054- 12066) - NADH dehydrogenas e subunit 5 | 13 |
| 170 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CA TC TC CA CA | (12068- 12069) - NADH dehydrogenas e subunit 5 | 2 |
| 171 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GT GG CC GT GT | (12071- 12072) - NADH dehydrogenas e subunit 5 | 2 |
| 172 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTCCATCA CTGGTCAA ACCTACAC CTCCATCA CTCCATCA | (12074- 12081) - NADH dehydrogenas e subunit 5 | 8 |
| 173 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GTTC TAGA GATC GTTC GTTC | (12083- 12086) - NADH dehydrogenas e subunit 5 | 4 |
| 174 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g2_i1 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GTCCCAATGCC GTGTTGCGATGT AACCAC----- GCCCCAATCGC GCCCCAATCGC | (12087- 12098) - NADH dehydrogenas e subunit 5 | 12 |

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| 175 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ATA TGG ATA ATA | (12106- 12108) - NADH dehydrogenas e subunit 5 | 3 |
| 176 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GTC TAA GTC GTC | (12105- 12107) - NADH dehydrogenas e subunit 5 | 3 |
| 177 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CA TT CA CA | (12109- 12110) - NADH dehydrogenas e subunit 5 | 2 |
| 178 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | G T G G | (12112) - NADH dehydrogenas e subunit 5 | 1 |
| 179 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TC AG TC TC | (12114- 12115) - NADH dehydrogenas e subunit 5 | 2 |
| 180 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ATC TCA ATC ATC | (12118- 12119) - NADH dehydrogenas e subunit 5 | 3 |
| 181 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GAAT TTGG GAAT GAAT | (12123- 12126) - NADH dehydrogenas e subunit 5 | 4 |
| 182 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TACAC AGTGA TACAC TACAC | (12128- 12132) - NADH dehydrogenas e subunit 5 | 5 |
| 183 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T G T T | (12135) - NADH dehydrogenas e subunit 5 | 1 |

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| 184 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ATACA TGTAT ATACA ATACA | (12137- 12141) - NADH dehydrogenas e subunit 5 | 5 |
| 185 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTAC GGTG CTAC CTAC | (12145- 12148) - NADH dehydrogenas e subunit 5 | 4 |
| 186 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GACCC TGAAG GACCC GACCC | (12150- 12154) - NADH dehydrogenas e subunit 5 | 5 |
| 187 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TATA ATGC TATA TATA | (12156- 12159) - NADH dehydrogenas e subunit 5 | 4 |
| 188 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CACA AGTG CACA CACA | (12161- 12164) - NADH dehydrogenas e subunit 5 | 4 |
| 189 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A T A A | (12167) - NADH dehydrogenas e subunit 5 | 1 |
| 190 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TTTC GAGA TTTC TTTC | (12170- 12173) - NADH dehydrogenas e subunit 5 | 4 |
| 191 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A T A A | (12176) - NADH dehydrogenas e subunit 5 | 1 |
| 192 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ACCTACTAATTTCC GTGGGTGTTGGAGGG ACCTACTAATTTCC ACCTACTAATTTCC | (12178- 12192) - NADH dehydrogenas e subunit 5 | 15 |

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| 193 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AG TT AG AG | (12194- 12195) - NADH dehydrogenas e subunit 5 | 2 |
| 194 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A T A A | (12197) - NADH dehydrogenas e subunit 5 | 1 |
| 195 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCA GGT CCA CCA | (12199- 12201) - NADH dehydrogenas e subunit 5 | 3 |
| 196 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AATAA TTGTT AATAA AATAA | (12203- 12207) - NADH dehydrogenas e subunit 5 | 5 |
| 197 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C A C C | (12209) - NADH dehydrogenas e subunit 5 | 1 |
| 198 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | G T G G | (12213) - NADH dehydrogenas e subunit 5 | 1 |
| 199 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AAC GTG AAC AAC | (12215- 12217) - NADH dehydrogenas e subunit 5 | 3 |
| 200 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCCAAAC TTAGTTCTA GCCAAAC GCCAAAC | (12219- 12227) - NADH dehydrogenas e subunit 5 | 9 |
| 201 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TAT GGC TAT TAT | (12229- 12231) - NADH dehydrogenas e subunit 5 | 3 |
| 202 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TCA AGT TCA TCA | (12233- 12235) - NADH | 3 |

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| | | | dehydrogenas e subunit 5 | |
| 203 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T G T T | (12237) - NADH dehydrogenas e subunit 5 | 1 |
| 204 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTTTA AGGCC CTTTA CTTTA | (12239- 12243) - NADH dehydrogenas e subunit 5 | 5 |
| 205 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T A T T | (12245) - NADH dehydrogenas e subunit 5 | 1 |
| 206 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C A C C | (12248) - NADH dehydrogenas e subunit 5 | 1 |
| 207 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GAGA TGTT GAGA GAGA | (12250- 12253) - NADH dehydrogenas e subunit 5 | 4 |
| 208 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GGAG CTGT GGAG GGAG | (12255- 12258) - NADH dehydrogenas e subunit 5 | 4 |
| 209 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GAA TGC GAA GAA | (12262- 12264) - NADH dehydrogenas e subunit 5 | 3 |
| 210 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAT GTG CAT CAT | (12266- 12268) - NADH dehydrogenas e subunit 5 | 3 |
| 211 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CC TT CC CC | (12271- 12272) - NADH dehydrogenas e subunit 5 | 2 |

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| 212 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TCCTCT AATGTA TCCTCT TCCTCT | (12274- 12279) - NADH dehydrogenas e subunit 5 | 6 |
| 213 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AA GT AA AA | (12281- 12282) - NADH dehydrogenas e subunit 5 | 2 |
| 214 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12284) - NADH dehydrogenas e subunit 5 | 1 |
| 215 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | G T G G | (12286) - NADH dehydrogenas e subunit 5 | 1 |
| 216 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GA AG GA GA | (12289- 12290) - NADH dehydrogenas e subunit 5 | 2 |
| 217 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GA TG GA GA | (12292- 12293) - NADH dehydrogenas e subunit 5 | 2 |
| 218 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CC TG CC CC | (12295- 12296) - NADH dehydrogenas e subunit 5 | 2 |
| 219 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCGAACAGA GGAGTTTT CCGAACAGA CCGAACAGA | (12299- 12307) - NADH dehydrogenas e subunit 5 | 9 |
| 220 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCAAAC ATGGTG GCAAAC GCAAAC | (12309- 12314) - NADH dehydrogenas e subunit 5 | 6 |

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| 221 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12316) - NADH dehydrogenas e subunit 5 | 1 |
| 222 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAGCCC ATTAG CAGCCC CAGCCC | (12319- 12324) - NADH dehydrogenas e subunit 5 | 6 |
| 223 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AC GA AC AC | (12326- 12327) - NADH dehydrogenas e subunit 5 | 2 |
| 224 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCCA TGAG GCCA GCCA | (12330- 12333) - NADH dehydrogenas e subunit 5 | 4 |
| 225 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T A T T | (12335) - NADH dehydrogenas e subunit 5 | 1 |
| 226 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TTTACAAC GCCTCGA TTTACAAC TTTACAAC | (12337- 12344) - NADH dehydrogenas e subunit 5 | 8 |
| 227 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TATCGGAGACA ATGCTTCCTAT TATTGGAGACG TATTGGAGACG | (12347- 12357) - NADH dehydrogenas e subunit 5 | 11 |
| 228 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CG GC CG CG | (12359- 12360) - NADH dehydrogenas e subunit 5 | 2 |
| 229 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTAACCTCAGTATAGCC GGCGGGTGATTGGGTGA CTAACCTCAGTATAGCC CTAACCTCAGTATAGCC | (12363- 12380) - NADH dehydrogenas e subunit 5 | 18 |

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| 230 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | G A G G | (12382) - NADH dehydrogenas e subunit 5 | 1 |
| 231 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C T C C | (12384) - NADH dehydrogenas e subunit 5 | 1 |
| 232 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAATAAAC TGGGTTGTT CAATAAAC CAATAAAC | (12388- 12396) - NADH dehydrogenas e subunit 5 | 9 |
| 233 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AAACAC TCGTTT AAACAC AAACAC | (12398- 12403) - NADH dehydrogenas e subunit 5 | 6 |
| 234 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TG GT TG TG | (12405- 12406) - NADH dehydrogenas e subunit 5 | 2 |
| 235 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A G A A | (12409) - NADH dehydrogenas e subunit 5 | 1 |
| 236 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12411) - NADH dehydrogenas e subunit 5 | 1 |
| 237 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCAACAAATCT GTGGAGGGTA CCAACAAATCT CCAACAAATCT | (12413- 12423) - NADH dehydrogenas e subunit 5 | 11 |
| 238 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TACCCACAC CGTGGGTGT TACCCACAC TACCCACAC | (12425- 12433) - NADH dehydrogenas e subunit 5 | 9 |
| 239 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AATC CTGT AATC AATC | (12435- 12438) - NADH dehydrogenas e subunit 5 | 4 |

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| 240 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CACC TGTA CACC CACC | (12440- 12443) - NADH dehydrogenas e subunit 5 | 4 |
| 241 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12445) - NADH dehydrogenas e subunit 5 | 1 |
| 242 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CT AG CT CT | (12447- 12448) - NADH dehydrogenas e subunit 5 | 2 |
| 243 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C T C C | (12450) - NADH dehydrogenas e subunit 5 | 1 |
| 244 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TCC AGG TCC TCC | (12452- 12454) - NADH dehydrogenas e subunit 5 | 3 |
| 245 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CT TG CT CT | (12456- 12457) - NADH dehydrogenas e subunit 5 | 2 |
| 246 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CT GA CT CT | (12459- 12460) - NADH dehydrogenas e subunit 5 | 2 |
| 247 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | G A G G | (12462) - NADH dehydrogenas e subunit 5 | 1 |
| 248 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A C A A | (12464) - NADH dehydrogenas e subunit 5 | 1 |
| 249 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TA GT TA TA | (12466- 12467) - NADH dehydrogenas e subunit 5 | 2 |

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| 250 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TCC AAT TCC TCC | (12469- 12471) - NADH dehydrogenas e subunit 5 | 3 |
| 251 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AGCCGC GCGGTG AGCCGC AGCCGC | (12473- 12478) - NADH dehydrogenas e subunit 5 | 6 |
| 252 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAG TGA CAG CAG | (12481- 12483) - NADH dehydrogenas e subunit 5 | 3 |
| 253 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AAAATCAGCCCCAA GTTGCGGTTAGTG AAAATCAGCCCCAA AAAATCAGCCCCAA | (12485- 12497) - NADH dehydrogenas e subunit 5 | 13 |
| 254 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C A C C | (12500) - NADH dehydrogenas e subunit 5 | 1 |
| 255 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CC AG CC CC | (12503- 12504) - NADH dehydrogenas e subunit 5 | 2 |
| 256 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCAC AGGG CCAC CCAC | (12506- 12509) - NADH dehydrogenas e subunit 5 | 4 |
| 257 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C T C C | (12511) - NADH dehydrogenas e subunit 5 | 1 |
| 258 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T A T T | (12513) - NADH dehydrogenas e subunit 5 | 1 |
| 259 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A G A A | (12515) - NADH dehydrogenas e subunit 5 | 1 |

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| 260 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TACC GTTT TACC TACC | (12517- 12520) - NADH dehydrogenas e subunit 5 | 4 |
| 261 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCAGCT TGTATG GCAGCT GCAGCT | (12522- 12527) - NADH dehydrogenas e subunit 5 | 6 |
| 262 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AG GT AG AG | (12530- 12531) - NADH dehydrogenas e subunit 5 | 2 |
| 263 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AGGCCCCACC TTTATGGTTT AGGCCCCACC AGGCCCCACC | (12533- 12542) - NADH dehydrogenas e subunit 5 | 10 |
| 264 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12544) - NADH dehydrogenas e subunit 5 | 1 |
| 265 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GTT TAA GTT GTT | (12546- 12548) - NADH dehydrogenas e subunit 5 | 3 |
| 266 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12550) - NADH dehydrogenas e subunit 5 | 1 |
| 267 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCAT TGTC GCAT GCAT | (12551- 12555) - NADH dehydrogenas e subunit 5 | 4 |
| 268 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ACTAC TTAGA ACTAC ACTAC | (12557- 12561) - NADH dehydrogenas e subunit 5 | 5 |

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| 269 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTC TAG CTC CTC | (12563- 12565) - NADH dehydrogenas e subunit 5 | 3 |
| 270 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | G T G G | (12568) - NADH dehydrogenas e subunit 5 | 1 |
| 271 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ACTA CAGT ACTA ACTA | (12570- 12573) - NADH dehydrogenas e subunit 5 | 4 |
| 272 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GT TA GT GT | (12576- 12577) - NADH dehydrogenas e subunit 5 | 2 |
| 273 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12581) - NADH dehydrogenas e subunit 5 | 1 |
| 274 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTG TAT CTG CTG | (12584- 12585) - NADH dehydrogenas e subunit 5 | 3 |
| 275 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AA CC AA AA | (12587- 12588) - NADH dehydrogenas e subunit 5 | 2 |
| 276 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C G C C | (12590) - NADH dehydrogenas e subunit 5 | 1 |
| 277 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TCC GAG TCC TCC | (12592- 12594) - NADH dehydrogenas e subunit 5 | 3 |
| 278 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A G A A | (12596) - NADH dehydrogenas e subunit 5 | 1 |

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| 279 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AT TG AT AT | (12600- 12601) - NADH dehydrogenas e subunit 5 | 2 |
| 280 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CGAAT TACCA CGAAT CGAAT | (12603- 12607) - NADH dehydrogenas e subunit 5 | 5 |
| 281 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C T C C | (12609) - NADH dehydrogenas e subunit 5 | 1 |
| 282 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCCCAT GTTAGG TCCCAT TCCCAT | (12611- 12616) - NADH dehydrogenas e subunit 5 | 6 |
| 283 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T A C C | (12618) - NADH dehydrogenas e subunit 5 | 1 |
| 284 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A G A A | (12620) - NADH dehydrogenas e subunit 5 | 1 |
| 285 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CC GG CC CC | (12622- 12623) - NADH dehydrogenas e subunit 5 | 2 |
| 286 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCA GGT CCA CCA | (12625- 12627) - NADH dehydrogenas e subunit 5 | 3 |
| 287 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAACAC ATTGGT CAACAC CAACAC | (12629- 12634) - NADH dehydrogenas e subunit 5 | 6 |
| 288 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCC GGA CCC CCC | (12637- 12639) - NADH | 3 |

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| | | | dehydrogenas e subunit 5 | |
| 289 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C T C C | (12641) - NADH dehydrogenas e subunit 5 | 1 |
| 290 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CAACCTG GTGTAGT CAACCTG CAACCTG | (12643- 12649) - NADH dehydrogenas e subunit 5 | 7 |
| 291 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TTGCC CCTAT TTGCC TTGCC | (12653- 12657) - NADH dehydrogenas e subunit 5 | 5 |
| 292 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AGGA TTTC AGGA AGGA | (12659- 12662) - NADH dehydrogenas e subunit 5 | 4 |
| 293 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCATCACCA AATATCTTGTT CCATCACCA CCATCACCA | (12664- 12674) - NADH dehydrogenas e subunit 5 | 11 |
| 294 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TA GT TA TA | (12676- 12677) - NADH dehydrogenas e subunit 5 | 2 |
| 295 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T A T T | (12679) - NADH dehydrogenas e subunit 5 | 1 |
| 296 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AC TT AC AC | (12681- 12682) - NADH dehydrogenas e subunit 5 | 2 |
| 297 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C A C C | (12685) - NADH dehydrogenas e subunit 5 | 1 |

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| 298 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | T A T A | (12688) - NADH dehydrogenas e subunit 5 | 1 |
| 299 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TGCGCCCTC GAATAATGG TGCGCCCTC TGCGCCCTC | (12690- 12698) - NADH dehydrogenas e subunit 5 | 9 |
| 300 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | C A C C | (12700) - NADH dehydrogenas e subunit 5 | 1 |
| 301 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AA TG AA AA | (12703- 12704) - NADH dehydrogenas e subunit 5 | 2 |
| 302 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ATGAT GCATA ATGAT ATGAT | (12706- 12710) - NADH dehydrogenas e subunit 5 | 5 |
| 303 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TC GA TC TC | (12712- 12713) - NADH dehydrogenas e subunit 5 | 2 |
| 304 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A T A A | (12715) - NADH dehydrogenas e subunit 5 | 1 |
| 305 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | AA GT AA AA | (12717- 12718) - NADH dehydrogenas e subunit 5 | 2 |
| 306 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | ATTA TGGC ATTA ATTA | (12720- 12723) - NADH dehydrogenas e subunit 5 | 4 |
| 307 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCCTTCTCCACATC TGAAGAATGCGTGT GCCTTCTCCACATC GCCTTCTCCACATC | (12726- 12739) - NADH | 14 |

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| | | | dehydrogenas e subunit 5 | |
| 308 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | A T A A | (12741) - NADH dehydrogenas e subunit 5 | 1 |
| 309 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CCA GAG CCA CCA | (12743- 12745) - NADH dehydrogenas e subunit 5 | 3 |
| 310 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | CTA TAT CTA CTA | (12747- 12749) - NADH dehydrogenas e subunit 5 | 3 |
| 311 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | GCCTT TAGGA GCCTT GCCTT | (12751- 12755) - NADH dehydrogenas e subunit 5 | 5 |
| 312 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 | TAA GGC TAA TAA | (12757- 12759) - NADH dehydrogenas e subunit 5 | 3 |
| 313 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | TGGCC TGGCC AAAAA TGGCC TGGCC TGGCC | (14218- 14222) - cytochrome b | 5 |
| 314 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | C C A C C C | (14224) - cytochrome b | 1 |
| 315 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | CCT CCT AAA CCT CCT CCT | (14227- 14230) - cytochrome b | 3 |

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| | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i3 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | CG CG AA CG CG CG | (14232- 14233) - cytochrome b | 2 |
| 316 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | A A A C A | (14849) - cytochrome b | 1 |
| 317 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | A A A T | (15369) - tRNA-Thr | 1 |
| 318 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g1_i1 | T T T C | (16205) - [D- loop] | 1 |
| 319 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN10253_c0_g1_i1 | CC AG AG CC CC | (16211- 16212) - [D- loop] | 2 |
| 320 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN10253_c0_g1_i1 | TAC CGG CGG TAC TAC | (16215- 16217) - [D- loop] | 3 |
| 321 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN10253_c0_g1_i1 | CC GA GA CC CC | (16220- 16221) - [D- loop] | 2 |
| 322 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN10253_c0_g1_i1 | A A A C A | (16215- 16217) - [D- loop] | 1 |
| 323 | secuencia referencia TRINITY_DN28133_c1_g1_i2 TRINITY_DN28133_c1_g1_i1 TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 | CC CC CC AT | (16239- 16240) - [D- loop] | 2 |
| 324 | secuencia referencia TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN71581_c0_g1_i1 | | | |

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| 325 | secuencia referencia TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN71581_c0_g1_i1 | CCAC CCAC CCAC GAGA | (16243-16246) - [D-loop] | 4 |
| 326 | secuencia referencia TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN71581_c0_g1_i1 | CCCA CCCA CCCA TAAC | (16248-16251) - [D-loop] | 4 |
| 327 | secuencia referencia TRINITY_DN28133_c0_g2_i1 TRINITY_DN28133_c0_g1_i1 TRINITY_DN71581_c0_g1_i1 | A A A G | (16258) - [D-loop] | 1 |
| TOTAL | | | | 1173 |

1.4. Heteroplasmia Cc3:

| Heteroplasmia | Cobertura | Nucleótidos | Región | Longitud |
|---------------|--|--|------------------------|----------|
| 1 | secuencia referencia TRINITY_DN5524_c0_g1_i1 TRINITY_DN990_c0_g1_i1 | GA GA TG | (32-33) - tRNA-Phe | 2 |
| 2 | secuencia referencia TRINITY_DN23496_c0_g1_i1 TRINITY_DN23786_c0_g1_i1 TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 | A C C A A | (3881) - tRNA-Gln | 1 |
| 3 | secuencia referencia TRINITY_DN23496_c0_g1_i1 TRINITY_DN23786_c0_g1_i1 TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 | T A A T T | (3889) - tRNA-Gln | 1 |
| 4 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | AAA AAA AAA GCC | (5354-5355) - tRNA-Tyr | 3 |
| 5 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | GGGACTACAGC GGGACTACAGC GGGACTACAGC CTATTCCTGCT | (5357-5367) - tRNA-Tyr | 11 |
| 6 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | CAA CAA CAA ATG | (5369-5371) - tRNA-Tyr | 3 |

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| 7 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | G G G C | (5373) - tRNA-Tyr | 1 |
| 8 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | TA TA TA CG | (5375-5376) - tRNA-Tyr | 2 |
| 9 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | TACACTCAGCC TACACTCAGCC TACACTCAGCC AGATTAGGTAT | (5378-5388) - tRNA-Tyr | 11 |
| 10 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | CCC CCC CCC GGG | (5390-5392) - tRNA-Tyr | 3 |
| 11 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 | A A A G | (5394) - tRNA-Tyr | 1 |
| 12 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 TRINITY_DN29176_c0_g1_i2 | C C C C G | (6694) - cytochrome c oxidase subunit I | 1 |
| 13 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN29198_c0_g1_i2 TRINITY_DN29176_c0_g1_i2 | TGGAATACCA TGGAATACCA TGGAATACCA TGGAATACCA GTCGTGGTAT | (6699-6708) - cytochrome c oxidase subunit I | 10 |
| 14 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN23610_c0_g1_i1 TRINITY_DN24191_c0_g1_i1 | GCTCCCT AAAAAAA AAAAAAA GCTCCCT GCTCCCT | (9463-9470) - tRNA-Gly | 8 |
| 15 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN23610_c0_g1_i1 TRINITY_DN24191_c0_g1_i1 | GT AA AA GT GT | (9472-9473) - tRNA-Gly | 2 |
| 16 | secuencia referencia TRINITY_DN29198_c0_g1_i1 TRINITY_DN29176_c0_g1_i1 TRINITY_DN23610_c0_g1_i1 TRINITY_DN24191_c0_g1_i1 | T A A T T | (9475) - tRNA-Gly | 1 |

| | | | | |
|-------|--|--|--|----|
| TOTAL | | | | 61 |
|-------|--|--|--|----|

1.3 RESUMEN DE LA HETEROPLASIA PARA CADA INDIVIDUO EVALUADO:

Ei1:

| Región | # de sitios heteroplásmicos | % de heteroplasmia de la región | % de la heteroplasmia total | % de heteroplasmia del total del genoma |
|----------------------------------|-----------------------------|---------------------------------|-----------------------------|---|
| rRNA-12S ribosomal RNA | 2 | 0,21 | 0,7 | 0,01 |
| rRNA-16S ribosomal RNA | 12 | 0,74 | 4,2 | 0,07 |
| tRNA-Tyr | 18 | 25,35 | 6,29 | 0,11 |
| tRNA-Asp | 26 | 37,14 | 9,09 | 0,16 |
| cytochrome c oxidase subunit II | 28 | 4,08 | 9,79 | 0,17 |
| tRNA-Lys | 48 | 65,75 | 16,78 | 0,29 |
| ATP synthase F0 subunit 8 | 47 | 28,48 | 16,43 | 0,29 |
| cytochrome c oxidase subunit III | 39 | 4,97 | 13,64 | 0,24 |
| NADH dehydrogenase subunit 3 | 2 | 0,57 | 0,7 | 0,01 |
| tRNA-Arg | 5 | 7,14 | 1,75 | 0,03 |
| tRNA-His | 5 | 7,14 | 1,75 | 0,03 |
| NADH dehydrogenase subunit 6 | 1 | 0,19 | 0,35 | 0,01 |
| tRNA-Glu | 32 | 47,06 | 11,19 | 0,19 |
| Zona intergénica | 1 | - | 0,35 | 0,01 |
| cytochrome b | 11 | 0,96 | 3,85 | 0,07 |
| tRNA-Thr | 9 | 12,68 | 3,15 | 0,05 |

| | | | | |
|-------|-----|---|-----|------|
| total | 286 | - | 100 | 1,74 |
|-------|-----|---|-----|------|

Cc1:

| Región | # de sitios heteroplásmicos | % de heteroplasmia de la región | % de la heteroplasmia total | % de heteroplasmia del total del genoma |
|----------------------------------|-----------------------------|---------------------------------|-----------------------------|---|
| rRNA-16S ribosomal RNA | 3 | 0,19 | 1 | 0,02 |
| NADH dehydrogenase subunit 2 | 32 | 3,07 | 10,7 | 0,19 |
| tRNA-Trp | 1 | 1,3 | 0,33 | 0,01 |
| tRNA-Tyr | 4 | 5,63 | 1,34 | 0,02 |
| cytochrome c oxidase subunit I | 33 | 2,13 | 11,04 | 0,2 |
| cytochrome c oxidase subunit III | 7 | 0,89 | 2,34 | 0,04 |
| tRNA-Gly | 10 | 14,71 | 3,34 | 0,06 |
| NADH dehydrogenase subunit 3 | 45 | 12,86 | 15,05 | 0,27 |
| tRNA-Arg | 48 | 68,57 | 16,05 | 0,29 |
| NADH dehydrogenase subunit 4L | 32 | 10,7 | 2,32 | 0,19 |
| cytochrome b | 76 | 6,64 | 25,42 | 0,46 |
| tRNA-Thr | 7 | 9,86 | 2,34 | 0,04 |
| tRNA-Pro | 1 | 1,41 | 0,33 | 0,01 |
| total | 299 | - | 100 | 1,8 |

Cc2:

| Región | # de sitios heteroplásmicos | % de heteroplasmia de la región | % de la heteroplasmia total | % de heteroplasmia del total del genoma |
|------------------------------|-----------------------------|---------------------------------|-----------------------------|---|
| NADH dehydrogenase subunit 1 | 1 | 0,1 | 0,09 | 0,01 |

| | | | | |
|----------------------------------|------|-------|-------|------|
| tRNA-Ile | 9 | 13,04 | 0,77 | 0,05 |
| tRNA-Met | 28 | 40,58 | 2,39 | 0,17 |
| NADH dehydrogenase subunit 2 | 5 | 0,48 | 0,43 | 0,03 |
| tRNA-Tyr | 18 | 25,35 | 1,53 | 0,11 |
| tRNA-Lys | 2 | 2,74 | 0,17 | 0,01 |
| ATP synthase F0 subunit 6 | 1 | 0,15 | 0,09 | 0,01 |
| cytochrome c oxidase subunit III | 2 | 0,26 | 0,17 | 0,01 |
| tRNA-Gly | 1 | 1,47 | 0,09 | 0,01 |
| NADH dehydrogenase subunit 4 | 221 | 16 | 18,84 | 1,34 |
| tRNA-His | 50 | 71,43 | 4,26 | 0,3 |
| tRNA-Ser | 55 | 83,33 | 4,69 | 0,33 |
| tRNA-Leu | 53 | 73,61 | 4,52 | 0,32 |
| NADH dehydrogenase subunit 5 | 694 | 38,62 | 59,16 | 4,22 |
| cytochrome b | 12 | 1,05 | 1,02 | 0,07 |
| tRNA-Thr | 1 | 1,41 | 0,09 | 0,01 |
| [D-loop] | 20 | 2,08 | 1,71 | 0,12 |
| total | 1173 | - | 100 | 7,13 |

Cc3:

| Región | # de sitios heteroplásmicos | % de heteroplasmia de la región | % de la heteroplasmia total | % de heteroplasmia del total del genoma |
|--------------------------------|-----------------------------|---------------------------------|-----------------------------|---|
| tRNA-Phe | 2 | 2,9 | 3,28 | 0,01 |
| tRNA-Gln | 2 | 2,82 | 3,28 | 0,01 |
| tRNA-Tyr | 35 | 49,3 | 57,38 | 0,21 |
| cytochrome c oxidase subunit I | 11 | 0,71 | 18,03 | 0,07 |
| tRNA-Gly | 11 | 16,18 | 18,03 | 0,07 |
| total | 61 | - | 100 | 0,37 |

1.4 MUTACIONES DE LAS TORTUGAS RELACIONADAS CON MUTACIONES PATOLÓGICAS EN HUMANOS. LOS CAMBIOS NUCLEOTÍDICOS EN NEGRO HACEN REFERENCIA A LOS REPORTADOS EN EL MTDNA HUMANO. LOS NUCLEÓTIDOS EN ROJO HACEN REFERENCIA A LOS CAMBIOS EVIDENCIADOS EN EL MTDNA DE LAS TORTUGAS ESTUDIADAS. LAS LECTURAS REVELAN EL NÚMERO DE SECUENCIAS QUE FUERON TOMADAS PARA DETERMINAR EL UMBRAL:

Ei1:

| Mutación | Locus | Enfermedad | Cambio nucleotídico | Umbra (%) | Lecturas |
|------------|---------|------------------------|---------------------|-----------|----------|
| A7877C | MT-COX2 | PEG glaucoma | A-C C-A | 25 | 4 |
| G8719A | MT-ATP6 | Suspected mito disease | G-A G-A | 33 | 3 |
| T8741G | MT-ATP6 | MILS | T-G A-G | 33 | 3 |
| TA11622del | MT-ND4 | CPEO | TA-del G-A | 50 | 2 |

Cc1:

| Mutación | Locus | Enfermedad | Cambio nucleotídico | Umbra (%) | Lecturas |
|----------|---------|-----------------------|---------------------|-----------|----------|
| T4648C | MT-ND2 | PEG | T-C A-T | 33 | 3 |
| T9861C | MT-COX3 | AD | T-C A-T | 25 | 4 |
| T9957C | MT-COX3 | MELAS | T-C C-A | 33 | 3 |
| T15209C | MT-CYB | Prader-Willi syndrome | T-C T-C | 33 | 3 |
| T15237C | MT-CYB | LHON | T-C G-A | 33 | 3 |
| T15287C | MT-CYB | DEAF | T-C C-A | 33 | 3 |

Cc2:

| Posición | Locus | Enfermedad | Cambio nucleotídico | Umbral (%) | Lecturas |
|----------|----------|--------------------------|---------------------|------------|----------|
| C3551T | MT-ND1 | LHON | C-T C-T | 33 | 3 |
| C3769G | MT-ND1 | LHON | C-G C-A | 66 | 3 |
| G3946A | MT-ND1 | MELAS | G-A G-C | 33 | 3 |
| T3949C | MT-ND1 | MELAS | T-C G-C | 33 | 3 |
| G3958A | MT-ND1 | LHON | G-A C-A | 33 | 3 |
| G3959A | MT-ND1 | MELAS | G-A C-T | 33 | 3 |
| A11467G | MT-ND4 | Altered brain pH | A-G A-T | 33 | 3 |
| G11696A | MT-ND4 | LHON | G-A G-T | 33 | 3 |
| C11777A | MT-ND4 | Leigh Disease | C-A A-T | 33 | 33 |
| G11778A | MT-ND4 | LHON | G-A T-G | 33 | 3 |
| G11832A | MT-ND4 | EXIT | G-A A-T | 25 | 4 |
| C11874A | MT-ND4 | LHON | C-A C-A | 25 | 4 |
| C11919T | MT-ND4 | Thyroid Cancer Cell Line | C-T A-T | 25 | 4 |
| C11994T | MT-ND4 | OAT | C-T A-T | 25 | 4 |
| A12026G | MT-ND4 | DM | A-G A-G | 25 | 4 |
| A12146G | tRNA His | MELAS | A-G T-G | 33 | 3 |
| G12147A | tRNA His | Encephalopathy | G-A A-T | 33 | 3 |
| T12148C | tRNA His | Developmental delay | T-C C-G | 33 | 3 |

| | | | | | |
|------------|----------|--|--------------|----|---|
| G12183A | tRNA His | DEAF | G-A C-T | 33 | 3 |
| C12187A | tRNA His | Asthenozoospermia | C-A T-G | 33 | 3 |
| G12192A | tRNA His | MICM | G-A C-G | 33 | 3 |
| C12206T | tRNA His | MELAS | C-T A-T | 33 | 3 |
| G12207A | tRNA Ser | Encephalopathy | G-A A-T | 33 | 3 |
| C12224T | tRNA Ser | DEAF | C-T C-T | 33 | 3 |
| C12258A | tRNA Ser | DMDF | C-A G-T | 33 | 3 |
| C12262A | tRNA Ser | Deafness | C-A G-T | 33 | 3 |
| C12264T | tRNA Ser | DEAF | C-T A-C | 33 | 3 |
| G12276A | tRNA Leu | CPEO | G-A C-T | 33 | 3 |
| A12299C | tRNA Leu | MELAS | A-C C-G | 33 | 3 |
| T12311C | tRNA Leu | CPEO | T-C A-G | 33 | 3 |
| T12313C | tRNA Leu | FSHD | T-C A-T | 33 | 3 |
| G12316A | tRNA Leu | CPEO | G-A C-G | 33 | 3 |
| A12320G | tRNA Leu | MM | A-G A-T | 33 | 3 |
| T12338C | MT-ND5 | LHON | T-C T-C | 33 | 3 |
| G12372A | MT-ND5 | Altered brain pH | G-A A-T | 33 | 3 |
| A12425de I | MT-ND5 | Mitochondrial Myopathy & Renal Failure | A-del T-C | 33 | 3 |
| T12477C | MT-ND5 | MHCM | T-C G-T | 33 | 3 |

| | | | | | |
|---------|--------|-----------------------------|------------|----|---|
| G12622A | MT-ND5 | Leigh Disease | G-A C-G | 33 | 3 |
| A12634G | MT-ND5 | Thyroid Cancer Cell Line | A-G C-T | 33 | 3 |
| T12706C | MT-ND5 | Leigh Disease | T-C A-G | 33 | 3 |
| T14849C | MT-CYB | EXIT | T-C A-C | 25 | 4 |

Cc3:

| Posición | Locus | Enfermedad | Cambio nucleotídico | Umbral (%) | Lecturas |
|----------|------------|------------------------|------------------------|------------|----------|
| G6708A | MT- CO1 | Rhabdomyolysis y MM | G-A A-T | 25 | 4 |