>Rhcg GENE: Ammonium transporter Rh type C ORF type: complete len:631 (+), score=195.01 TRINITY_DN731_c0_g1:336-2228(+)
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>Rhcg2 GENE: Ammonium transporter Rh type C 2 ORF type: complete len:440 (+), score=74.24 TRINITY_DN731_c0_g2: 36-1322(+)
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>Rhag GENE: Ammonium transporter Rh type A ORF type: complete len:325 (-), score=72.50
TRINITY_DN15891_c0_g1: 639-1613(-)
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>AQP3 GENE: Aquaporin-3 ORF type: complete len:156 (+), score=4.71 TRINITY_DN9467_c0_g1: 248-715(+)
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>AQP1 GENE: Aquaporin-1 ORF type: complete len:278 (+), score=18.30 TRINITY_DN9406_c0_g3: 26-835(+)
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>NHE9B2 GENE: sodium/hydrogen exchanger 9B2 ORF type: complete len:558 (+), score=118.86 TRINITY_DN30526_c1_g1: 134-1807(+)

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>NHE2 GENE: Sodium/hydrogen exchanger 2 ORF type: 3prime_partial len:489 (+), score=69.79 TRINITY_DN20285_c0_g1:64-1527(+)

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>CA4 GENE: Carbonic anhydrase 4 ORF type: complete len:319 (+), score=53.99 TRINITY_DN182_c1_g1:2761-3717(+)

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>CA2 GENE: Carbonic anhydrase 2 ORF type: complete len:261 (+), score=51.50 TRINITY_DN4375_c0_g1: 407-1189(+)

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>ANO8 GENE: anoctamin-8 ORF type: complete len:669 (-), score=102.92 TRINITY_DN1428_c2_g1: 1744-3750(-)

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>IRAK1 GENE: interleukin-1 receptor-associated kinase 1 ORF type: complete len:458 (+), score=83.44 TRINITY_DN2476_c0_g1:1109-2482(+)

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