

MATCH_INFO
<div> SOURCE_SEQ_seq-id VARCHAR(100) </div> <div> match-found-bool TINYINT(1) </div> <div> match-order VARCHAR(20) </div> <div> match-family VARCHAR(20) </div> <div> match-genus VARCHAR(20) </div> <div> match-species VARCHAR(20) </div>
<div> Indexes </div>

SOURCE_SEQ
<div> seq-id VARCHAR(100) </div> <div> match-found-bool TINYINT(1) </div> <div> nucleotide-sequence VARCHAR(5000) </div>
<div> Indexes </div>

BLAST_RESULT
<div> SOURCE_SEQ_seq-id VARCHAR(100) </div> <div> amount-hits INT(11) </div> <div> result1-accession-code VARCHAR(2000) </div> <div> result1-alignment-score FLOAT </div> <div> result1-query-coverage-percentage INT(11) </div> <div> result1-E-value FLOAT </div> <div> result1-ident-percentage INT(11) </div> <div> result2-accession-code VARCHAR(2000) </div> <div> result2-alignment-score FLOAT </div> <div> result2-query-coverage-percentage INT(11) </div> <div> result2-E-value FLOAT </div> <div> result2-ident-percentage INT(11) </div> <div> result3-accession-code VARCHAR(2000) </div> <div> result3-alignment-score FLOAT </div> <div> result3-query-coverage-percentage INT(11) </div> <div> result3-E-value FLOAT </div> <div> result3-ident-percentage INT(11) </div> <div> result4-accession-code VARCHAR(2000) </div> <div> result4-alignment-score FLOAT </div> <div> result4-query-coverage-percentage INT(11) </div> <div> result4-E-value FLOAT </div> <div> result4-ident-percentage INT(11) </div> <div> result5-accession-code VARCHAR(2000) </div> <div> result5-alignment-score FLOAT </div> <div> result5-query-coverage-percentage INT(11) </div> <div> result5-E-value FLOAT </div> <div> result5-ident-percentage INT(11) </div> <div> result6-accession-code VARCHAR(2000) </div> <div> result6-alignment-score FLOAT </div> <div> result6-query-coverage-percentage INT(11) </div> <div> 22 more... </div>
<div> Indexes </div>

