```
"Contig_213_101.679:1.6825_3" : {
                                                   // Entry ID or primary key
"Contig": "Contig_213_101.679:1.6825",
                                                   // Prediction template
"Protein_id": "Contig_213_101.679:1.6825_3",
                                                   // Protein | ORF ID
"Start" : 1576,
                                                   // Predicted start position [nt]
"Stop" : 2980,
                                                   // Predicted stop position [nt]
"Strand" : 1,
                                                   // Direction: 1=fwd, -1=rev
"Length aa" : 467.
                                                   // Length of predicted protein [aa]
"Orflength" : 1401,
                                                   // Length of predicted protein [nt]
"pI" : 6.09124755859,
                                                   // Estimated Isoelectric point
"Molecular_weight" : 53.5625,
                                                   // Estimated molecular weight [kDa]
"Instability_index" : 36.6983083512,
                                                   // Estimated Instability Index
"SRR" : "SRR6659510",
                                                   // SRR accession of contig
"Organism" : "Edwardsiella phage PEi21",
                                                   // Source organism of prediction source
"Hitscore": 0.736478082660833.
                                                   // Score of prediction
"Virus_quotient" : 0.832707055,
                                                   // Viral qutient
"Description": "phage terminase, large subunit",
"Sources" : {
                                                   // Sources used in VIGA
  "VOG4544" : {
                                                   // HMM hit (pVOG or RVDBs)
      "Evalue" : 7.4e-28,
                                                   // Evalue/Significance of hit
       "Coverage" : 70.1754385965,
                                                   // Coverage to closest reference [%]
      "Similarity" : 56
                                                   // Similarity to closest reference [%]
  },
                                                   // Prediction source
   "DIAMOND" : {
      "Coverage" : 98.07,
                                                   // Coverage to closest reference [%]
      "Similarity" : 51.1,
                                                   // Similarity to closest reference [%]
       "Evalue" : 1.5e-119
                                                   // Evalue/Significance of hit
```