```
"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 quotient
  "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 [%]
    },
    "DIAMOND" : {                                          // Prediction source
        "Coverage" : 98.07,                                // Coverage to closest reference [%]
        "Similarity" : 51.1,                               // Similarity to closest reference [%]
        "Evalue" : 1.5e-119                                // Evalue/Significance of hit
    }
  }
}
```