

<pre> { "_id" : ObjectId("57c6b2e3e138233c43eed53"), "gene" : "XLOC_010191", "chromosome" : "chrA03", "start" : 6240007, "end" : 6240929, "measurements" : [{ "time" : 22, "fpkm" : 16.764, "hi" : 23.254, "lo" : 10.2741, "tissue" : "apex", "accession" : "tapidor" }, ... { "time" : 72, "fpkm" : 0.0671628, "hi" : 0.27532, "lo" : 0, "tissue" : "leaf", "accession" : "westar" }], "homology" : [{ "agi" : "AT5G10140.4", "symbols" : ["FLC"], "hsp_bit_score" : 260.971, "identity" : 91.3043478261, "length_of_hsp" : 184 }, ... { "agi" : "AT5G10140.1", "symbols" : ["FLC", "FLF", "AGL25"], "hsp_bit_score" : 260.971, "identity" : 91.3043478261, "length_of_hsp" : 184 }] } </pre>	<pre> "_id" "gene" "chromosome" "start" "end" </pre> <p>Identifier assigned by the database software Gene name assigned by Cufflinks Chromosome on which the gene is located Base number at which the gene model begins Base number at which the gene model ends</p>
	<pre> "measurements" "time" "fpkm" "hi" "lo" "tissue" "accession" </pre> <p>List of measurements The days post sowing the tissue was sampled FPKM value Upper bound of the FPKM confidence interval Lower bound of the FPKM confidence interval Tissue of origin <i>Brassica napus</i> variety of origin</p>
	<pre> "homology" "agi" "symbols" "hsp_bit_score" "identity" "length_of_hsp" </pre> <p>List of <i>Arabidopsis thaliana</i> gene models that show sequence conservation to the <i>Brassica napus</i> gene <i>Arabidopsis thaliana</i> gene identifier <i>Arabidopsis thaliana</i> gene symbols Highest scoring pair bit score Percentage sequence identity of highest scoring pair Length of highest scoring pair</p>