

<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" Identifier assigned by the database software "gene" Gene name assigned by Cufflinks "chromosome" Chromosome on which the gene is located "start" Base number at which the gene model begins "end" Base number at which the gene model ends </pre>
	<pre> "measurements" List of measurements "time" Gene name assigned by Cufflinks "fpkm" Chromosome on which the gene is located "hi" Base number at which the gene model begins "lo" Base number at which the gene model ends "tissue" Tissue of origin "accession" <i>Brassica napus</i> variety of origin </pre>
	<pre> "homology" List of <i>Arabidopsis thaliana</i> gene models that show sequence conservation to the <i>Brassica napus</i> gene "agi" <i>Arabidopsis thaliana</i> gene identifier "symbols" <i>Arabidopsis thaliana</i> gene symbols "hsp_bit_score" Highest scoring pair bit score "identity" Percentage of highest scoring pair sequence identity "length_of_hsp" Length of highest scoring pair </pre>