

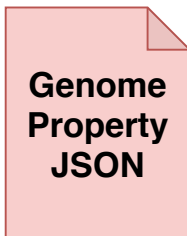
Endpoints

get_tree

get_fasta

**get_single
_genome
_property
_info**

**get_multiple
_genome
_property
_info**



to_json()

**write_supporting_
proteins_for_step_fasta()**

to_json()

to_json()

**Genome
Properties Results
With Matches**

**Genome
Properties Tree**

