

find\_differential\_genes  
mirna: mir197  
organism: human

find\_differential\_genes  
mirna: mir769  
organism: human

find\_differential\_genes  
mirna: mir197  
organism: mouse

find\_differential\_genes  
mirna: mir769  
organism: mouse

all

```
graph TD; A["find_differential_genes<br/>mirna: mir197<br/>organism: human"] --> D(all); B["find_differential_genes<br/>mirna: mir769<br/>organism: human"] --> D; C["find_differential_genes<br/>mirna: mir197<br/>organism: mouse"] --> D; E["find_differential_genes<br/>mirna: mir769<br/>organism: mouse"] --> D;
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The diagram illustrates a process where four separate search queries are combined into a single result set. Each query is contained within a green-bordered rounded rectangle. The queries are: 1) find\_differential\_genes mirna: mir197 organism: human, 2) find\_differential\_genes mirna: mir769 organism: human, 3) find\_differential\_genes mirna: mir197 organism: mouse, and 4) find\_differential\_genes mirna: mir769 organism: mouse. Arrows from each of these four boxes point towards a central red-bordered rounded rectangle labeled 'all', indicating that the results of all four searches are being aggregated.