

quantify_genes
sample: Sample1

quantify_genes
sample: Sample2

collate_outputs

all

```
graph TD; A["quantify_genes<br/>sample: Sample1"] --> C["collate_outputs"]; B["quantify_genes<br/>sample: Sample2"] --> C; C --> D["all"];
```

The diagram is a flowchart with three levels. At the top, there are two boxes with a light green border. The left box contains the text 'quantify_genes' followed by 'sample: Sample1' on a new line. The right box contains 'quantify_genes' followed by 'sample: Sample2' on a new line. Arrows from the bottom of both of these boxes point towards a central box below them. This central box has a light green border and contains the text 'collate_outputs'. An arrow points from the bottom of this central box to a final box at the bottom. This final box has a red border and contains the text 'all'.