Phase 1: simple contig browser

Pan Left   Zoom In/Out   Pan Right

Scale

0k   1k   2k   3k   4k   5k   6k   7k   8k   9k   10k

Escherichia coli str. K-12 substr. MG1655 (- strand, 3876051..3886051)

trmE   yidC   rnpA   dnaA   recF   gyrB

yidD   rpmH   dnaN

+
-

Export to SVG

Genome strand

Save frame
image/data to file

Genome name

Gene orientation
(start/stop)

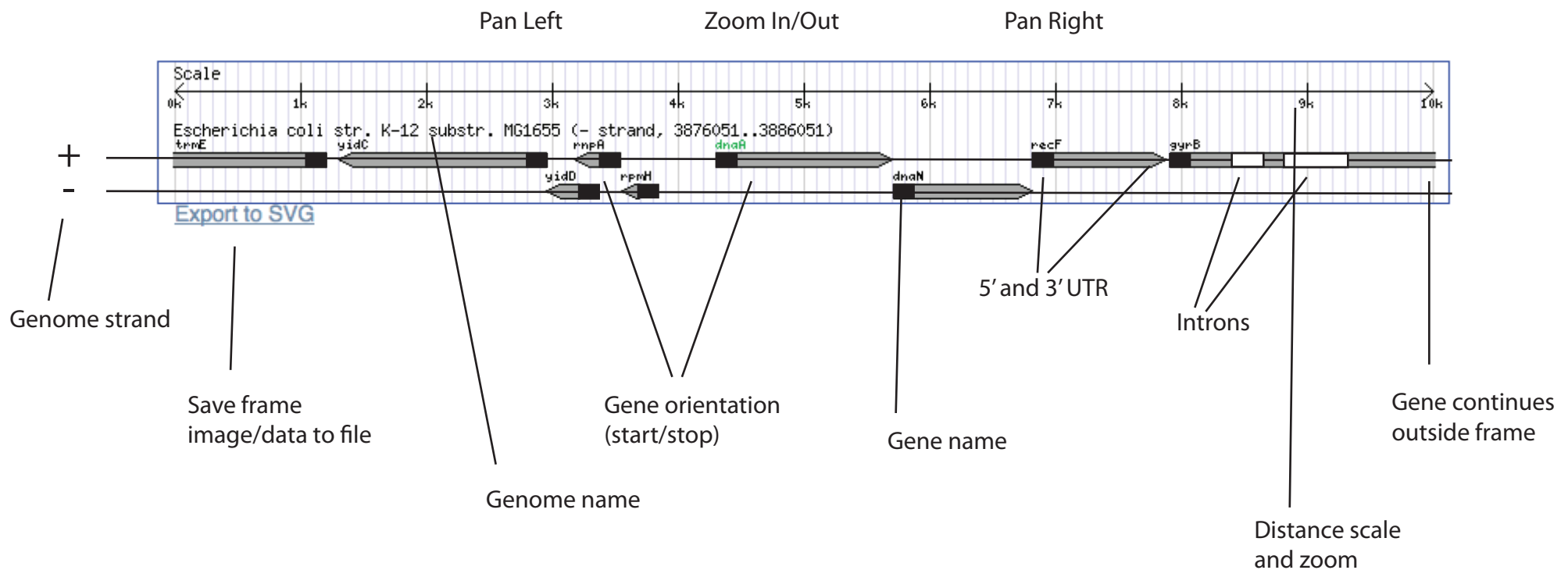Gene name

5' and 3' UTR

Introns

Distance scale
and zoom

Gene continues
outside frame

*Note, this does not represent alternative splicing yet.
**Even this simple contig browser could show a single track of data by the thickness of the gene line, e.g. gene expression.