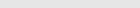
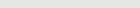


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Description	NC_016565.1: <i>Escherichia coli</i> O157:H7, complete genome, 4,855,221 bp, 100% GC content. This is a reference genome for a strain of <i>E. coli</i> O157:H7, a serotype associated with hemorrhagic colitis and hemolytic uremic syndrome. The genome is circular and contains 10,415 genes, including 10,388 protein-coding genes, 26 rRNA genes, and 1 tRNA gene. The genome is annotated with various features, including genes, CDS, and regulatory elements. NC_016565.1: <i>Escherichia coli</i> O157:H7, complete genome, 4,855,221 bp, 100% GC content. This is a reference genome for a strain of <i>E. coli</i> O157:H7, a serotype associated with hemorrhagic colitis and hemolytic uremic syndrome. The genome is circular and contains 10,415 genes, including 10,388 protein-coding genes, 26 rRNA genes, and 1 tRNA gene. The genome is annotated with various features, including genes, CDS, and regulatory elements.
Organism	<i>Escherichia coli</i>
Tissue	Intestine

XXXXXXXXXXXX

Morphology	
Growth properties	

XXXXXXXXXX XXXXXXXXXXXXXXXXXXXXXXX

Citation	L6565 (Cytion 305189)
Biosafety level	1
NCBI_TaxID	10090
CellosaurusAccession	CVCL_A9NB

XXXXXXXXXX

[illegible]

██████████ ████████████ ████████████, ████ ████████████ ████████████ ████████████ ████████████ (██████████ FBS) + 10% DMSO ████████████ ████████████ ████████████ ████████████ ████████████, ████ C

[illegible]37°C, 5% CO₂, 100% humidity

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[illegible][illegible][illegible]

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Sterility

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XX XXXXXXXX XXXX XXXXXXX XXXXXXX, XXXXXXXXXXX XX XXXX, XXXXXXXXXXX XXXXXXX XXXXXXX XXXXXXX XXXXXXXXXXX XXXXXXX.