

Test Results

Order #: **2022021226**

CHU de Quebec
(# 36408 - CHU de Quebec-Site CHUL bloc R
QC)

2705 Boulevard Laurier
Quebec, QC G1V 4G2 Canada

Charles River Research Animal Diagnostic Services
(CR RADS)

261 Ballardvale Street
Receiving Dock, Bldg 22
Wilmington MA 01887 USA

Billing Information

Payment Method

Standing Purchase PO#: 92171
Order Exp. 12/2022

Details

Sample(s) from: Multiple locations

Collection Date
28-Apr-2022

Arrival Date
02-May-2022

Approval Date
09-May-2022

Notes

secteur 40 trimestre 2 support B

Diagnostic Summary

Test	Colony	Tested	+	+/-	?	PDG
All results NEGATIVE						

+ = Positive, +/- = Equivocal, ? = Indeterminate, PDG = Pending

To assure the health status of your research animal colonies, it is essential that you understand the sources, pathobiology, diagnosis and control of pathogens and other adventitious infectious agents that may cause research interference. We have summarized this important information in infectious agent **Technical Sheets**, which you can view by visiting http://www.criver.com/info/disease_sheets.

Test Results

Order #: **2022021226**

CHU de Quebec
(# 36408 - CHU de Quebec-Site CHUL bloc R
QC)

2705 Boulevard Laurier
Quebec, QC G1V 4G2 Canada

Charles River Research Animal Diagnostic Services
(CR RADS)

261 Ballardvale Street
Receiving Dock, Bldg 22
Wilmington MA 01887 USA

Notes

secteur 40 trimestre 2 support B

Results approved by Muise, Delia on 09 May 2022

Molecular Diagnostics: Infectious Disease PCR

Mite and Pinworm Screen PRIA

	<u>3</u> 40.1 Ba+b PCR	<u>6</u> 40.2 Ba+b PCR	<u>9</u> 40.7 Ba+b PCR	<u>12</u> 40.8 Ba+b PCR
Pinworm PCR	-	-	-	-
Mite PCR	-	-	-	-

Remarks

- = Negative, +/- = Equivocal; + = Positive; I = Inconclusive.

An equivocal result indicates inconsistent amplification detected by real-time PCR.

Inconclusive indicates failure of control result.

Nucleic Acid Recovery Control (NRC)/Inhibition Control: A low copy exogenous nucleic acid was added to sample lysis prior to nucleic acid isolation to serve as both a control to monitor for nucleic acid recovery and PCR inhibition. An RNA NRC also monitors reverse transcription for RNA virus assays. Nucleic acid recovery and PCR inhibition is monitored by a PCR assay specific for the NRC template. Unless otherwise stated, the control results passed for this order.

Any samples reported as equivocal or positive result in this report has been confirmed by re-extracting nucleic acid and repeating real-time PCR amplification to confirm the initial testing result.

Recommended sample types are essential to accurate results. Missing or inappropriate sample types and/or expired buffer/additives can affect detection. If this report contains an unexpected result or are unsure of recommended sample types, please contact Lab Services@crl.com before taking any action. Additional or alternative testing may be essential to reaching an accurate diagnosis. We will be glad to test newly submitted samples for the positive agents up to the number of unexpected results in this order.

Test Results

Order #: **2022021226**

CHU de Quebec
(# 36408 - CHU de Quebec-Site CHUL bloc R
QC)

2705 Boulevard Laurier
Quebec, QC G1V 4G2 Canada

Charles River Research Animal Diagnostic Services
(CR RADS)

261 Ballardvale Street
Receiving Dock, Bldg 22
Wilmington MA 01887 USA

Notes

secteur 40 trimestre 2 support B

Serology

Results approved by Estevez, Rebecca on 04 May 2022

	<u>1</u> 40.1 Ba2 sero	<u>2</u> 40.1 Bb2 sero	<u>4</u> 40.2 Ba2 Sero	<u>5</u> 40.2 Bb2 Sero	<u>7</u> 40.7 Ba2 sero	<u>8</u> 40.7 Bb2 sero	<u>10</u> 40.8 Ba2 sero	<u>11</u> 40.8 Bb2 sero
MFIA MHV	-	-	-	-	-	-	-	-
MFIA MVM	-	-	-	-	-	-	-	-
MFIA MPV-1	-	-	-	-	-	-	-	-
MFIA MPV-2	-	-	-	-	-	-	-	-
MFIA NS-1	-	-	-	-	-	-	-	-
MFIA MNV	-	-	-	-	-	-	-	-
MFIA GDVII	-	-	-	-	-	-	-	-
MFIA EDIM (ROTA-A)	-	-	-	-	-	-	-	-
MFIA Anti-Ig	P	P	P	P	P	P	P	P

Serology Profile: QC MFIA Mouse Prevalent Profile

Remarks

MFIA/IFA/ELISA/WIB Results: - = Negative; +/- = Equivocal; + = Moderate to strong positive; TC = Non-specific reaction with tissue control; I = Indeterminate or Inconclusive; IN = result interpreted as non-specific because not confirmed by alternative serologic assay or diagnostic methodology for other serologic assays, PDG = pending, QNS = Quantity not sufficient. The anti-immunoglobulin (Anti-Ig) MFIA verifies that a serum specimen contains a sufficient concentration of immunoglobulin to be suitable for serologic testing. A result of P (for Pass) corresponds to a median fluorescence index (MFI) at or above the Anti-Ig assay cutoff, typically ≥ 7000 . An Anti-Ig assay result of F (for Fail), assigned if the MFI is below the cutoff, might occur because the sample was received too dilute, was collected from an immunocompromised host or was from a species other than the one for which the MFIA is intended. If a sample fails the Anti-Ig MFIA, then negative and borderline results in MFIA for microbial antibodies are considered I (for inconclusive).

Test Results

Order #: **2022021226**

CHU de Quebec
(# 36408 - CHU de Quebec-Site CHUL bloc R
QC)

2705 Boulevard Laurier
Quebec, QC G1V 4G2 Canada

Charles River Research Animal Diagnostic Services
(CR RADS)

261 Ballardvale Street
Receiving Dock, Bldg 22
Wilmington MA 01887 USA

Notes

secteur 40 trimestre 2 support B

Sample Information

Number	Code	Species	Colony
1	40.1 Ba2 sero	Mouse	R-3740.1 CD-1 Mice (Rack A, B, C, D, E, F)
2	40.1 Bb2 sero	Mouse	R-3740.1 CD-1 Mice (Rack A, B, C, D, E, F)
3	40.1 Ba+b PCR	Mouse	R-3740.1 CD-1 Mice (Rack A, B, C, D, E, F)
4	40.2 Ba2 Sero	Mouse	R-3740.2 CD-1 Mice (Rack A, B, C, D, E, F)
5	40.2 Bb2 Sero	Mouse	R-3740.2 CD-1 Mice (Rack A, B, C, D, E, F)
6	40.2 Ba+b PCR	Mouse	R-3740.2 CD-1 Mice (Rack A, B, C, D, E, F)
7	40.7 Ba2 sero	Mouse	R-3740.7 CD-1 Mice (Rack A, B, C, D, E, F)
8	40.7 Bb2 sero	Mouse	R-3740.7 CD-1 Mice (Rack A, B, C, D, E, F)
9	40.7 Ba+b PCR	Mouse	R-3740.7 CD-1 Mice (Rack A, B, C, D, E, F)
10	40.8 Ba2 sero	Mouse	R-3740.8 CD-1 Mice (Rack A, B, C, D, E, F)
11	40.8 Bb2 sero	Mouse	R-3740.8 CD-1 Mice (Rack A, B, C, D, E, F)
12	40.8 Ba+b PCR	Mouse	R-3740.8 CD-1 Mice (Rack A, B, C, D, E, F)