## User Information

<table>
<thead>
<tr>
<th>Name</th>
<th>Email</th>
<th>Dept./Inst.</th>
<th>P.I.</th>
<th>TEL</th>
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</thead>
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| *UC Funding Code/PO#: |

### Fund Manager Information

| Name | Email | TEL |

### Possible Biohazard

- Yes**/ No BSL1/BSL2 (no BSL3+)

### DNA/Tissue

- HMW Genomic gDNA
- Tissue for Extraction
  - mg: ______
  - Tissue: ______
  - Past Storage: ______
  - (include excel if many samples)

### IsoSeq Submissions

- Total RNA
- cDNA

### HMW Quality check only

- Pulse Field Gel only (Femto Pulse)
- FPGE, qubit, + Nanodrop

### Info about Library

- Amplicons
- Size range: ______

- HMW
- Desired insert size: <10kb, 20kb, 30kb

- Multiplexing
- Yes (requires 5-7kb lib)

### Info about Sequencing

- No. of SMRT cell/sample per each (~3-5Gb)

### Notes:

Submit the following info as an excel sheet with this form via email to vcgsq qb3@gmail.com

### User Affiliation

- UC / Affiliate / Academic / For-Profit

### Used other chemicals-

- Extraction
- gDNA QC
- Full length cDNA
- Lib prep
- SMRT cells

### Order Total:

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*Please note that we will **not process** without fund a source. **Please contact before submitting your samples to review your submission needs. Please also send a 5ul aliquot of gDNA samples so QC can be performed without thawing HMW gDNA. All samples should be shipped on dry ice and stay frozen in transit.*