# **Request for bed file creation**

**Instructions:**

1. **Add genes symbols and/or transcripts to a panel in Moka**
2. **Add panel description to Moka**
3. **Answer following questions:**
4. **Pan number of bed file to be created:**

Pan2078

1. **Panel description/name**

EB Virtual Panel (-/+5bp) for Agilent Focused Exome

1. **Purpose of BED files:** (delete as appropriate)

* Calculating coverage

1. **Should specific transcripts be used to create the bed file?** (delete as appropriate)

* Yes - use specified transcripts (see question 4a)

**4a) If transcripts are to be used have you provided multiple transcripts for the same gene?**

* No (If yes, please state any genes with multiple transcripts)

1. **If bed file is to be used for coverage:**

* Exon padding:\_5bp

1. **If bed file is to be used for CNV RPKM analysis:**

* Control sites to include: Pan\_\_\_\_
* Exon padding: \_\_\_\_bp

1. **Inclusion of UTRs**:

* Include 5’ and 3’ UTRs for all genes: No
* Padding of UTRs: No
* If UTRs are only required for a subset of genes please see section 8

1. **Any additional sites to be included:**

* Please provide genomic coordinates for target regions (without padding)
* Please describe region eg UTR, intronic site.
* Padding will be added as specified above

Eg: Chr1:123456-234567 5’ UTR for Gene ABC

**Requested by:** Michael Yau **Requested date:** 11/04/2018