# **Request for bed file creation**

Use this form to request the creation of a single BED file (one Pan number). If RPKM and coverage bedfiles are required please submit these on seperate forms.

**Instructions:**

* **Add genes symbols and/or transcripts to a panel in Moka**
* **Add panel description to Moka**
* **Answer following questions:**

1. **Pan number of bed file to be created:**  
   Pan3611
2. **Panel description/name**

|  |  |  |
| --- | --- | --- |
| Category | SubCategory | Panel |
| LS\_MSI\_v1 | MSI Homopolymer Variants | LS\_MSI\_v1 - MSI Homopolymer Variants |

1. **Is this BED file for RPKM?** (delete as appropriate)

No

1. **Is this BED file exclusively based on coordinates provided by a supplier** (e.g SNP identity kits).

No – see attached 

chr14 23652257 23652457 NR-21

chr2 95849238 95849438 NR-24

chr11 102193421 102193621 NR-27

chr4 55598115 55598315 BAT-25

chr2 47641466 47641666 BAT-26

* 1. If yes, does this need further padding?

No

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

N/A

* 1. **If specific transcripts are to be used have you provided multiple transcripts for the same gene?** (if Yes please state)

N/A

1. **What padding is required around the exons?**

N/A

1. **Should UTRs be included for ALL genes?** (delete as appropriate)

N/A

* 1. **Should UTRs be padded?**

N/A

1. **Are UTRs required for SOME genes?** (delete as appropriate)

N/A

* 1. **Please list genes/transcripts.**
  2. **Should UTRs be padded?**

N/A

1. **Any additional BED files to be included in this BED file?**

N/A

1. **Are any further regions required to be added?**

N/A

**Requested by:** Michael Yau

**Requested date:** 21/04/2020