# **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:**  
   Pan3608
2. **Panel description/name**

|  |  |  |
| --- | --- | --- |
| Category | SubCategory | Panel |
| VCP\_CNV\_Control\_Sites\_v1 | VCP CNV Control Sites | VCP\_CNV\_Control\_Sites\_v1 - VCP CNV Control Sites |

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

Yes

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

No – see attached 

chr10 90869951 90870151 CNV02\_1

chr10 90875132 90875332 CNV03

chr10 90905302 90905502 CNV07

chr11 126373957 126374157 CNV08

chr14 103715862 103716062 CNV09\_1

chr17 28745696 28745896 CNV10\_1

chr19 10869817 10870017 CNV11

chr1 55038099 55038299 CNV12\_1

chr20 10643041 10643241 CNV13

chr2 112805963 112806163 CNV14\_1

chr5 13763956 13764156 CNV15

chr6 160243267 160243467 CNV16\_1

* 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