# **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:**

Pan43450

1. **Panel description/name**

|  |  |  |
| --- | --- | --- |
| **Category** | **SubCategory** | **Panel** |
| VCP2 Viapath | R208 intronic CNV regions | VCP2 Viapath R208 intronic CNV regions |

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).

Yes

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

?

17 41322636 41322796 BRCA1\_PM\_5\_4

17 41297651 41297811 BRCA1\_PM\_5\_3

17 41286238 41286398 BRCA1\_PM\_5\_2

17 41281042 41281202 BRCA1\_PM\_5\_1

17 41235012 41235172 BRCA1\_IN11\_2

17 41234663 41234823 BRCA1\_IN11\_1

17 41234197 41234357 BRCA1\_IN12\_1

17 41233848 41234008 BRCA1\_IN12\_2

17 41196280 41196440 BRCA1\_3UTR

13 32973679 32973839 BRCA2\_3UTR

16 23614453 23614613 PALB2\_3UTR

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

No

* 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.**

N/A

* 1. **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?**

No

**Requested by: M Yau**

**Requested date:** 09/03/2021