# **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:**  
   Pan3609
2. **Panel description/name**

|  |  |  |
| --- | --- | --- |
| Category | SubCategory | Panel |
| HID\_Variant\_v1 | In\_Del variants | HID\_Variant\_v1 - In\_Del 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 

NC\_000021.8:g.31372337\_31372338 rs10629077

NC\_000011.9:g.268180\_268181 rs10688868

NC\_000010.10:g.54442383\_54442385 rs1160886

NC\_000005.9:g.65378462\_65378464 rs1160956

NC\_000012.11:g.14909997\_14909998 rs1610919

NC\_000009.11:g.38406791\_38406794 rs16402

NC\_000002.11:g.235016390\_235016391 rs16624

NC\_000016.9:g.84582282\_84582286 rs2067208

NC\_000012.11:g.115288548\_115288549 rs2067238

NC\_000009.11:g.71314422\_71314424 rs2067294

NC\_000005.9:g.5125113\_5125114 rs2307526

NC\_000001.10:g.247812085\_247812086 rs2307579

NC\_000009.11:g.105586196\_105586199 rs2307580

NC\_000019.9:g.44204340\_44204342 rs2307689

NC\_000022.10:g.26790904\_26790905 rs2307700

NC\_000006.11:g.47821256\_47821259 rs2307710

NC\_000006.11:g.117093555\_117093556 rs2307839

NC\_000007.13:g.83283914\_83283915 rs2307978

NC\_000015.9:g.53481516\_53481517 rs2308020

NC\_000004.11:g.119185405\_119185406 rs2308026

NC\_000006.11:g.149614198\_149614199 rs2308137

NC\_000013.10:g.44880155\_44880156 rs2308171

NC\_000014.8:g.29036752\_29036756 rs2308189

NC\_000003.11:g.8616708\_8616709 rs2308242

NC\_000001.10:g.162810826\_162810829 rs3047269

NC\_000017.10:g.10135938\_10135941 rs3051300

NC\_000018.9:g.23253207\_23253208 rs3080855

NC\_000020.10:g.11695625\_11695626 rs33917182

NC\_000011.9:g.126288872\_126288873 rs33972805

NC\_000018.9:g.36423040\_36423044 rs34511541

NC\_000011.9:g.30177690\_30177691 rs34811743

NC\_000008.10:g.76518681\_76518682 rs35769550

NC\_000019.9:g.1402662\_1402663 rs36040336

NC\_000008.10:g.138420592\_138420593 rs5895447

* 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