

## **Alcohol QTL PDF set of three documents:**

- \*1. AlcoholQTLsDescriptions**
- 2. AlcoholQTLsByChromosome**
- 3. AlcoholQTLsByPhenotype**

### **This document is 1 of 3, \*AlcoholQTLsDescriptions**

This document lists individual alcohol-related QTLs by *QTL Code*, in alphabetical order.

Detailed information includes, name, alias(es), the drug (*Alcohol* = ethanol), the broad phenotype designation, and a somewhat more detailed description of how that phenotype was assessed (*Assay*). LOD scores are typically based on  $df = 2$  (i.e., two degrees of freedom). *Strains Tested* gives mapping populations for which data have been cumulated to test significance, which often includes selectively bred lines and/or congenic lines or strains. The sex specificity is given if known, the increaser allele identified as D2 (DBA/2J) or B6 (C57BL/6J), and dominance given if known. Where results are published, the references are given. Where unpublished, the PI's laboratory is stated. The time of the last update of the tabled information is indicated. QTL information for loss of righting reflex was obtained from Beth Bennett and Tom Johnson at the Institute for Behavior Genetics in Boulder.

### **About *QTL Code* - Names and Aliases**

There is now terminological confusion for many traits. We at PARC adopted early on (and have maintained) the practice of not "naming" our QTLs in the literature unless they reached Lander & Kruglyak "significant" levels. These lists also report our "suggestive" QTLs by that standard. Mouse Genome Informatics has of their own initiative named several of our published QTLs. They also assigned names to some QTLs regardless of statistical significance. They made assumptions about the phenotypes in several instances. For instance, they treated Acute QTLs mapped in the Hitzemann lab as representing the same trait as Ectact QTLs mapped in the Phillips lab. Both groups assessed Alcohol Stimulated Activity, but they used different doses of ethanol and different time periods after injections. To our way of very conservative thinking, these are not the same trait. So, we list a number of aliases in our table to try to indicate what is going on. The *QTL Code* name is our new local name, and sometimes combines two earlier names. For example, the Chronic Alcohol Withdrawal QTL on chromosome 14 we originally named Caws5 was renamed by MGI as Alcw7. We now also name all suggestive QTLs using the suffix "prov" for provisional, which allows us to keep them with the same basic name, and to upgrade them later if additional data render them significant.

For more information, contact Mark Rutledge-Gorman at [rutledgm@ohsu.edu](mailto:rutledgm@ohsu.edu)

# Aaq1/Etacc

|                |  |                  |                     |              |              |
|----------------|--|------------------|---------------------|--------------|--------------|
| QTL Code       | Aaq1/Etacc   | PARC Alias       | Etacc               | MGI Alias    | Aaq1         |
| Drug           | Alcohol  | Phenotype        | Alcohol Acceptance  |              |              |
| Assay          | Total amount of 10% (v/v) ethanol consumed over 24-h period following 24-h period of water deprivation |                  |                     |              |              |
| Chromosome     | 15   | Peak             | 30                  | Range        | 15-48        |
| Max Range (Mb) |  | Min Range (Mb)   |                     | Status       | Significant  |
| LOD score      | 3.8  | p value          |                     | Sex Specific | Male limited |
| Strains Tested | B6, D2, BXD RI, B6D2 F2  | Increaser Allele | B6                  | Dominance    |              |
| Comments       |  |                  |                     |              |              |
| Contact Person |  | Last Updated     | 2007-04-10 13:12:20 |              |              |

## References

McClearn GE, Tarantino LM, Rodriguez LA, Jones BC, Blizzard DA, Plomin R. Genotypic selection provides experimental confirmation for an alcohol consumption quantitative trait locus in mouse. Mol Psychiatry. 1997 Oct-Nov;2(6):486-9.

[Medline Link](#)

# Actre1/Actre1q

|                |                |                  |                             |              |             |
|----------------|----------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre1/Actre1q | PARC Alias       | Actre1q                     | MGI Alias    | Actre1      |
| Drug           | Alcohol        | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |                |                  |                             |              |             |
| Chromosome     | 1              | Peak             | 90 cM                       | Range        |             |
| Max Range (Mb) |                | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |                | p value          | 1e-12                       | Sex Specific |             |
| Strains Tested | BXD RI, F2, HS | Increaser Allele |                             | Dominance    |             |
| Comments       |                |                  |                             |              |             |
| Contact Person | R. Hitzemann   | Last Updated     | 2007-04-12 10:46:09         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. *Mamm Genome* 2006 Dec;17(12):1193-204.

[Medline Link](#)

Demarest K, Koyner J, McCaughran J Jr, Cipp L, Hitzemann R. Further characterization and high-resolution mapping of quantitative trait loci for ethanol-induced locomotor activity. *Behav Genet.* 2001 Jan;31(1):79-91.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre1q

|            |         |                |                             |
|------------|---------|----------------|-----------------------------|
| QTL Code   | Actre1q |                |                             |
| Chromosome | 1       | Class          | Alcohol                     |
| Peak       | 90 cM   | Phenotype      | Ethanol-Stimulated Activity |
| Range      |         | Assay          |                             |
| p Value    | 1e-12   | Strains Tested | BXD RI, F2, HS              |
| LOD Score  |         | Status         | Significant                 |

## References

Demarest K, Koyner J, McCaughran J Jr, Cipp L, Hitzemann R. Further characterization and high-resolution mapping of quantitative trait loci for ethanol-induced locomotor activity. Behav Genet. 2001 Jan;31(1):79-91.

[Medline Link](#)

# Actre2

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre2  | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol   | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |   |                  |                             |              |             |
| Chromosome     | 2   | Peak             | 56 cM                       | Range        |             |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |   | p value          |                             | Sex Specific |             |
| Strains Tested |   | Increaser Allele |                             | Dominance    |             |
| Comments       | May be the same QTL as Actre3/Actre3q and/or Actre4 |                  |                             |              |             |
| Contact Person | R. Hitzemann  | Last Updated     | 2007-04-09 13:56:34         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre3/Actre3q

|                |                |                  |                             |              |             |
|----------------|----------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre3/Actre3q | PARC Alias       | Actre3q                     | MGI Alias    | Actre3      |
| Drug           | Alcohol        | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |                |                  |                             |              |             |
| Chromosome     | 2              | Peak             | 60 cM                       | Range        |             |
| Max Range (Mb) |                | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |                | p value          | 1e-07                       | Sex Specific |             |
| Strains Tested | BXD RI, F2, HS | Increaser Allele |                             | Dominance    |             |
| Comments       |                |                  |                             |              |             |
| Contact Person | R. Hitzemann   | Last Updated     | 2007-04-09 13:56:34         |              |             |

## References

Demarest K, Koyner J, McCaughran J Jr, Cipp L, Hitzemann R. Further characterization and high-resolution mapping of quantitative trait loci for ethanol-induced locomotor activity. Behav Genet. 2001 Jan;31(1):79-91.

[Medline Link](#)

# Actre3q

|            |         |                |                             |
|------------|---------|----------------|-----------------------------|
| QTL Code   | Actre3q |                |                             |
| Chromosome | 2       | Class          | Alcohol                     |
| Peak       | 60 cM   | Phenotype      | Ethanol-Stimulated Activity |
| Range      |         | Assay          |                             |
| p Value    | 1e-07   | Strains Tested | BXD RI, F2, HS              |
| LOD Score  |         | Status         | Significant                 |

## References

Demarest K, Koyner J, McCaughran J Jr, Cipp L, Hitzemann R. Further characterization and high-resolution mapping of quantitative trait loci for ethanol-induced locomotor activity. Behav Genet. 2001 Jan;31(1):79-91.

[Medline Link](#)

# Actre4

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre4  | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol   | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |   |                  |                             |              |             |
| Chromosome     | 2   | Peak             | 65 cM                       | Range        |             |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |   | p value          |                             | Sex Specific |             |
| Strains Tested |   | Increaser Allele |                             | Dominance    |             |
| Comments       | May be the same QTL as Actre2 and/or Actre3/Actre3q |                  |                             |              |             |
| Contact Person | R. Hitzemann  | Last Updated     | 2007-04-09 13:54:44         |              |             |

## References

Demarest K, Koyner J, McCaughran J Jr, Cipp L, Hitzemann R. Further characterization and high-resolution mapping of quantitative trait loci for ethanol-induced locomotor activity. Behav Genet. 2001 Jan;31(1):79-91.

[Medline Link](#)

# Actre5

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre5       | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 3            | Peak             | 56 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:07:07         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre6

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre6       | PARC Alias       |                             | MGI Alias    | Actre6      |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 9            | Peak             | 52 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:51:21         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre7

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre7       | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 1            | Peak             | 113 cM                      | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 13:56:34         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre8

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre8       | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 4            | Peak             | 8 cM                        | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:16:02         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre9

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre9       | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 6            | Peak             | 17 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:21:28         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre10

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre10      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 7            | Peak             | 34 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:46:54         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre11

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre11      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 7            | Peak             | 61 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:47:32         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre12

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre12      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 8            | Peak             | 48 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:50:09         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre13

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre13      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 10           | Peak             | 51 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:55:45         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre14

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre14      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 11           | Peak             | 37 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 14:58:53         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre15

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre15      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 14           | Peak             | 37 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 15:08:35         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre16

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre16      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 15           | Peak             | 67 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 15:09:56         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre17

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre17      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | 19           | Peak             | 24 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 15:16:57         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre18

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre18      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | X            | Peak             | 33 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 15:17:28         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Actre19

|                |              |                  |                             |              |             |
|----------------|--------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Actre19      | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol      | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |              |                  |                             |              |             |
| Chromosome     | X            | Peak             | 50 cM                       | Range        |             |
| Max Range (Mb) |              | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      |              | p value          |                             | Sex Specific |             |
| Strains Tested |              | Increaser Allele |                             | Dominance    |             |
| Comments       |              |                  |                             |              |             |
| Contact Person | R. Hitzemann | Last Updated     | 2007-04-09 15:17:51         |              |             |

## References

Malmanger B, Lawler M, Coulombe S, Murray R, Cooper S, Polyakov Y, Belknap J, Hitzemann R. Further studies on using multiple-cross mapping (MCM) to map quantitative trait loci. Mamm Genome 2006 Dec;17(12):1193-204.

[Medline Link](#)

Hitzemann R, Boudreau E and Malmanger B; unpublished data

# Alcdpprov1

|                |   |                  |                            |              |              |
|----------------|---|------------------|----------------------------|--------------|--------------|
| QTL Code       | Alcdpprov1  | PARC Alias       | Calws                      | MGI Alias    |              |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |              |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |              |
| Chromosome     | 13  | Peak             | 37                         | Range        | 16-59        |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Suggestive   |
| LOD score      | 2.9   | p value          | 0.0003                     | Sex Specific | Male Limited |
| Strains Tested | BXD RI, B6D2 F2   | Increaser Allele | D2                         | Dominance    |              |
| Comments       |   |                  |                            |              |              |
| Contact Person | K. Buck   | Last Updated     | 2007-04-05 11:03:44        |              |              |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15  
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Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. Addiction. 2001 Jan;96(1):139-49.  
[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.  
[Medline Link](#)

# Alcdpprov2

|                |   |                  |                            |              |            |
|----------------|---|------------------|----------------------------|--------------|------------|
| QTL Code       | Alcdpprov2  | PARC Alias       | Calw4                      | MGI Alias    |            |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |            |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |            |
| Chromosome     | 1   | Peak             | 20                         | Range        | 12-35      |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Suggestive |
| LOD score      | 3.1   | p value          | 1e-04                      | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2   | Increaser Allele | B6                         | Dominance    |            |
| Comments       |   |                  |                            |              |            |
| Contact Person | K. Buck   | Last Updated     | 2007-04-04 13:13:58        |              |            |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15  
[Medline Link](#)

# Alcdpprov3

|                |   |                  |                            |              |            |
|----------------|---|------------------|----------------------------|--------------|------------|
| QTL Code       | Alcdpprov3  | PARC Alias       | Calw5                      | MGI Alias    |            |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |            |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |            |
| Chromosome     | 4   | Peak             | 37                         | Range        | 30-44      |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Suggestive |
| LOD score      | 2.0   | p value          | 0.003                      | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2   | Increaser Allele | D2                         | Dominance    |            |
| Comments       |   |                  |                            |              |            |
| Contact Person | K. Buck   | Last Updated     | 2007-04-04 13:22:55        |              |            |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15  
[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.  
[Medline Link](#)

# Alcp1

|                |         |                  |                             |              |             |
|----------------|---------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Alcp1   | PARC Alias       | Alcp1                       | MGI Alias    | Alcp1       |
| Drug           | Alcohol | Phenotype        | Alcohol Preference Drinking |              |             |
| Assay          |         |                  |                             |              |             |
| Chromosome     | 2       | Peak             | 28                          | Range        | 19-35       |
| Max Range (Mb) |         | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      | 5.3     | p value          |                             | Sex Specific |             |
| Strains Tested |         | Increaser Allele |                             | Dominance    |             |
| Comments       |         |                  |                             |              |             |
| Contact Person | JA Melo | Last Updated     | 2007-04-05 12:29:20         |              |             |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Melo JA, Shendure J, Pociask K, Silver LM. Identification of sex-specific quantitative trait loci controlling alcohol preference in C57BL/6 mice. *Nat Genet.* 1996 Jun;13(2):147-53.

[Medline Link](#)

# Alcw1

|                |  |                  |                              |              |             |
|----------------|--|------------------|------------------------------|--------------|-------------|
| QTL Code       | Alcw1  | PARC Alias       | Alcw1                        | MGI Alias    | Alcw1       |
| Drug           | Alcohol  | Phenotype        | Acute Alcohol Withdrawal     |              |             |
| Assay          | Area under the curve for handling-induced convulsions 2-12h after 4 g/kg ethanol injection |                  |                              |              |             |
| Chromosome     | 1  | Peak             | 96                           | Range        | 66-ter      |
| Max Range (Mb) | 172.3 - 174.0, 174.0 - 176.5   | Min Range (Mb)   | 172.9 - 174.0, 174.0 - 176.3 | Status       | Significant |
| LOD score      | 5.6  | p value          | 3e-07                        | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2, HAW/LAW   | Increaser Allele | D2                           | Dominance    |             |
| Comments       | Two qtls in this interval. Mb intervals based on congenics studies                         |                  |                              |              |             |
| Contact Person | K. Buck  | Last Updated     | 2007-04-04 13:08:20          |              |             |

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[Medline Link](#)

Kozell LM, Buck KJ; unpublished results

# Alcw2

|                |  |                  |                          |              |             |
|----------------|--|------------------|--------------------------|--------------|-------------|
| QTL Code       | Alcw2  | PARC Alias       | Alcw2                    | MGI Alias    | Alcw2       |
| Drug           | Alcohol  | Phenotype        | Acute Alcohol Withdrawal |              |             |
| Assay          | Area under the curve for handling-induced convulsions 2-12h after 4 g/kg ethanol injection |                  |                          |              |             |
| Chromosome     | 4  | Peak             | 38                       | Range        | 38-38.7     |
| Max Range (Mb) | 80.3 - 82.1  | Min Range (Mb)   | 80.5 - 82.0              | Status       | Significant |
| LOD score      | 8.8  | p value          | 2e-10                    | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2, HAW/LAW   | Increaser Allele | D2                       | Dominance    |             |
| Comments       | Mb intervals based on congenics studies  |                  |                          |              |             |
| Contact Person | K. Buck  | Last Updated     | 2007-04-04 13:11:36      |              |             |

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Buck KJ, Metten P, Belknap JK, Crabbe JC. Quantitative trait loci involved in genetic predisposition to acute alcohol withdrawal in mice. J Neurosci. 1997 May 15;17(10):3946-55.

[Medline Link](#)

# Alcw3

|                |  |                  |                          |              |             |
|----------------|--|------------------|--------------------------|--------------|-------------|
| QTL Code       | Alcw3  | PARC Alias       | Alcw3                    | MGI Alias    | Alcw3       |
| Drug           | Alcohol  | Phenotype        | Acute Alcohol Withdrawal |              |             |
| Assay          | Area under the curve for handling-induced convulsions 2-12h after 4 g/kg ethanol injection |                  |                          |              |             |
| Chromosome     | 11   | Peak             | 20                       | Range        | 16-28       |
| Max Range (Mb) | 34.7 - 55.9  | Min Range (Mb)   | 34.9 - 55.6              | Status       | Significant |
| LOD score      | 4.1  | p value          | 1e-05                    | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2, HAW/LAW   | Increaser Allele | D2                       | Dominance    |             |
| Comments       | Mb intervals based on D1D2 SNP mapping   |                  |                          |              |             |
| Contact Person | K. Buck  | Last Updated     | 2007-04-04 13:11:51      |              |             |

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Buck KJ, Metten P, Belknap JK, Crabbe JC. Quantitative trait loci involved in genetic predisposition to acute alcohol withdrawal in mice. *J Neurosci.* 1997 May 15;17(10):3946-55.

[Medline Link](#)

# Alcw4

|                |  |                  |                          |              |            |
|----------------|--|------------------|--------------------------|--------------|------------|
| QTL Code       | Alcw4  | PARC Alias       | Alcw4                    | MGI Alias    | Alcw4      |
| Drug           | Alcohol  | Phenotype        | Acute Alcohol Withdrawal |              |            |
| Assay          | Area under the curve for handling-induced convulsions 2-12h after 4 g/kg ethanol injection |                  |                          |              |            |
| Chromosome     | 2  | Peak             | 38                       | Range        | 28-53      |
| Max Range (Mb) |  | Min Range (Mb)   |                          | Status       | Suggestive |
| LOD score      | 2.3  | p value          | 0.001                    | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2, HAW/LAW   | Increaser Allele | D2                       | Dominance    |            |
| Comments       |  |                  |                          |              |            |
| Contact Person | K. Buck  | Last Updated     | 2007-04-04 13:08:51      |              |            |

## References

Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. *Addiction*. 2001 Jan;96(1):139-49.

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Buck KJ, Metten P, Belknap JK, Crabbe JC. Quantitative trait loci involved in genetic predisposition to acute alcohol withdrawal in mice. *J Neurosci*. 1997 May 15;17(10):3946-55.

[Medline Link](#)

# Alcw5

|            |       |                |  |
|------------|-------|----------------|--|
| QTL Code   | Alcw5 |                |  |
| Chromosome | 2     | Class          | Alcohol  |
| Peak       | 60 cM | Phenotype      | Acute Alcohol Withdrawal   |
| Range      | ND cM | Assay          | Area under the curve for handling-induced convulsions 2-12h after 4 g/kg ethanol injection |
| p Value    | 0.002 | Strains Tested | BXD RI, B6D2 F2, HAW/LAW   |
| LOD Score  | 2.1   | Status         | Suggestive   |

## References

Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. *Addiction*. 2001 Jan;96(1):139-49.

[Medline Link](#)

Buck KJ, Metten P, Belknap JK, Crabbe JC. Quantitative trait loci involved in genetic predisposition to acute alcohol withdrawal in mice. *J Neurosci*. 1997 May 15;17(10):3946-55.

[Medline Link](#)

# Alcwprov1

|                |  |                  |                          |              |            |
|----------------|--|------------------|--------------------------|--------------|------------|
| QTL Code       | Alcwprov1  | PARC Alias       | Alcw5                    | MGI Alias    |            |
| Drug           | Alcohol  | Phenotype        | Acute Alcohol Withdrawal |              |            |
| Assay          | Area under the curve for handling-induced convulsions 2-12h after 4 g/kg ethanol injection |                  |                          |              |            |
| Chromosome     | 2  | Peak             | 60                       | Range        | ND         |
| Max Range (Mb) |  | Min Range (Mb)   |                          | Status       | Suggestive |
| LOD score      | 2.1  | p value          | 0.002                    | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2, HAW/LAW   | Increaser Allele | B6                       | Dominance    |            |
| Comments       |  |                  |                          |              |            |
| Contact Person | K. Buck  | Last Updated     | 2007-04-04 13:10:20      |              |            |

## References

Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. *Addiction*. 2001 Jan;96(1):139-49.

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Buck KJ, Metten P, Belknap JK, Crabbe JC. Quantitative trait loci involved in genetic predisposition to acute alcohol withdrawal in mice. *J Neurosci*. 1997 May 15;17(10):3946-55.

[Medline Link](#)

# Ap1q

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Ap1q  | PARC Alias       | Ap1q                        | MGI Alias    | Ap1q        |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |             |
| Assay          |   |                  |                             |              |             |
| Chromosome     | 1   | Peak             | 76                          | Range        | 45-95       |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      | 4.5   | p value          |                             | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2   | Increaser Allele | B6                          | Dominance    | B6 dominant |
| Comments       | QTL was NOT captured in a congenic with minimal interval of 152.1 - 176.5. However this does not co |                  |                             |              |             |
| Contact Person | LM Tarantino  | Last Updated     | 2007-04-05 14:14:07         |              |             |

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Tarantino LM, McClearn GE, Rodriguez LA, Plomin R. Confirmation of quantitative trait loci for alcohol preference in mice. *Alcohol Clin Exp Res.* 1998 Aug;22(5):1099-105.

[Medline Link](#)

Phillips TJ; unpublished data

# Ap2q

|                |                 |                  |                             |              |            |
|----------------|-----------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Ap2q            | PARC Alias       | Ap2q                        | MGI Alias    | Ap2q       |
| Drug           | Alcohol         | Phenotype        | Alcohol Preference Drinking |              |            |
| Assay          |                 |                  |                             |              |            |
| Chromosome     | 2               | Peak             | 40                          | Range        | 30-50      |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      | 3.1             | p value          |                             | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele |                             | Dominance    |            |
| Comments       |                 |                  |                             |              |            |
| Contact Person | LM Tarantino    | Last Updated     | 2007-04-05 13:34:12         |              |            |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Tarantino LM, McClearn GE, Rodriguez LA, Plomin R. Confirmation of quantitative trait loci for alcohol preference in mice. *Alcohol Clin Exp Res.* 1998 Aug;22(5):1099-105.

[Medline Link](#)

# Ap3q

|                |                 |                  |                             |              |                         |
|----------------|-----------------|------------------|-----------------------------|--------------|-------------------------|
| QTL Code       | Ap3q            | PARC Alias       | Ap3q                        | MGI Alias    | Ap3q                    |
| Drug           | Alcohol         | Phenotype        | Alcohol Preference Drinking |              |                         |
| Assay          |                 |                  |                             |              |                         |
| Chromosome     | 4               | Peak             | 75                          | Range        | 65-ter                  |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Significant             |
| LOD score      | 4.6             | p value          |                             | Sex Specific |                         |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele | B6                          | Dominance    | B6 dominant or additive |
| Comments       |                 |                  |                             |              |                         |
| Contact Person | LM Tarantino    | Last Updated     | 2007-04-05 14:06:21         |              |                         |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Tarantino LM, McClearn GE, Rodriguez LA, Plomin R. Confirmation of quantitative trait loci for alcohol preference in mice. *Alcohol Clin Exp Res.* 1998 Aug;22(5):1099-105.

[Medline Link](#)

# Ap4q

|                |                 |                  |                             |              |            |
|----------------|-----------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Ap4q            | PARC Alias       | Ap4q                        | MGI Alias    | Ap4q       |
| Drug           | Alcohol         | Phenotype        | Alcohol Preference Drinking |              |            |
| Assay          |                 |                  |                             |              |            |
| Chromosome     | 10              | Peak             | 28                          | Range        | 18-57      |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      | 3.2             | p value          |                             | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele |                             | Dominance    |            |
| Comments       |                 |                  |                             |              |            |
| Contact Person | LM Tarantino    | Last Updated     | 2007-04-05 14:11:59         |              |            |

## References

Tarantino LM, McClearn GE, Rodriguez LA, Plomin R. Confirmation of quantitative trait loci for alcohol preference in mice. *Alcohol Clin Exp Res.* 1998 Aug;22(5):1099-105.

[Medline Link](#)

# Ap5q

|                |                 |                  |                             |              |                         |
|----------------|-----------------|------------------|-----------------------------|--------------|-------------------------|
| QTL Code       | Ap5q            | PARC Alias       | Ap5q                        | MGI Alias    | Ap5q                    |
| Drug           | Alcohol         | Phenotype        | Alcohol Preference Drinking |              |                         |
| Assay          |                 |                  |                             |              |                         |
| Chromosome     | 9               | Peak             | 25                          | Range        | 10-35                   |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Significant             |
| LOD score      | 4.8             | p value          |                             | Sex Specific |                         |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele | B6                          | Dominance    | B6 dominant or additive |
| Comments       |                 |                  |                             |              |                         |
| Contact Person | LM Tarantino    | Last Updated     | 2007-04-05 14:07:46         |              |                         |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

[Medline Link](#)

Tarantino LM, McClearn GE, Rodriguez LA, Plomin R. Confirmation of quantitative trait loci for alcohol preference in mice. *Alcohol Clin Exp Res.* 1998 Aug;22(5):1099-105.

[Medline Link](#)

# Ap6q

|                |                 |                  |                             |              |            |
|----------------|-----------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Ap6q            | PARC Alias       | Ap6q                        | MGI Alias    | Ap6q       |
| Drug           | Alcohol         | Phenotype        | Alcohol Preference Drinking |              |            |
| Assay          |                 |                  |                             |              |            |
| Chromosome     | 3               | Peak             | 45                          | Range        | 38-60      |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      | 3.4             | p value          |                             | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele |                             | Dominance    |            |
| Comments       |                 |                  |                             |              |            |
| Contact Person | LM Tarantino    | Last Updated     | 2007-04-05 13:34:59         |              |            |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Tarantino LM, McClearn GE, Rodriguez LA, Plomin R. Confirmation of quantitative trait loci for alcohol preference in mice. *Alcohol Clin Exp Res.* 1998 Aug;22(5):1099-105.

[Medline Link](#)

# Calw1

|            |             |                |   |
|------------|-------------|----------------|---|
| QTL Code   | Calw1       |                |   |
| Chromosome | 1           | Class          | Alcohol   |
| Peak       | 96 cM       | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 89-106.3 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 2e-09       | Strains Tested | BXD RI, B6D2F2, congenics   |
| LOD Score  | 7.6         | Status         | Significant   |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15

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Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. Addiction. 2001 Jan;96(1):139-49.

[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.

[Medline Link](#)

# Calw1/Alcdp1

|                |   |                  |                            |              |             |
|----------------|---|------------------|----------------------------|--------------|-------------|
| QTL Code       | Calw1/Alcdp1  | PARC Alias       | Calw1                      | MGI Alias    | Alcdp1      |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |             |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |             |
| Chromosome     | 1   | Peak             | 96                         | Range        | 89-106.3    |
| Max Range (Mb) | 172.3 - 191.3   | Min Range (Mb)   | 172.9 - 186.9              | Status       | Significant |
| LOD score      | 7.6   | p value          | 2e-09                      | Sex Specific |             |
| Strains Tested | BXD RI, B6D2F2, congenics   | Increaser Allele | D2                         | Dominance    |             |
| Comments       | Mb intervals based on congenics studies   |                  |                            |              |             |
| Contact Person | K. Buck   | Last Updated     | 2007-04-04 13:20:40        |              |             |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15  
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Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. Addiction. 2001 Jan;96(1):139-49.  
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Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.  
[Medline Link](#)

Kozell LM, Buck KJ; unpublished results

# Calw2

|            |             |                |   |
|------------|-------------|----------------|---|
| QTL Code   | Calw2       |                |   |
| Chromosome | 19          | Class          | Alcohol   |
| Peak       | 47 cM       | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 36.5-ter cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 6e-07       | Strains Tested | BXD RI, B6D2 F2   |
| LOD Score  | 5.3         | Status         | Significant   |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15

[Medline Link](#)

Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. Addiction. 2001 Jan;96(1):139-49.

[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.

[Medline Link](#)

# Calw2/Alcdp2

|                |   |                  |                            |              |             |
|----------------|---|------------------|----------------------------|--------------|-------------|
| QTL Code       | Calw2/Alcdp2  | PARC Alias       | Calw2                      | MGI Alias    | Alcdp2      |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |             |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |             |
| Chromosome     | 19  | Peak             | 47                         | Range        | 36.5-ter    |
| Max Range (Mb) | 10.4 - 55.2   | Min Range (Mb)   | 12.3 - 54.8                | Status       | Significant |
| LOD score      | 5.3   | p value          | 6e-07                      | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2   | Increaser Allele | D2                         | Dominance    |             |
| Comments       | Mb intervals based on congenics studies   |                  |                            |              |             |
| Contact Person | K. Buck   | Last Updated     | 2007-04-05 11:06:28        |              |             |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15  
[Medline Link](#)

Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. Addiction. 2001 Jan;96(1):139-49.  
[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.  
[Medline Link](#)

Shirley RL, Buck KJ; unpublished data

# Calw3

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Calw3    |                |   |
| Chromosome | 13       | Class          | Alcohol   |
| Peak       | 37 cM    | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 16-59 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 0.0003   | Strains Tested | BXD RI, B6D2 F2   |
| LOD Score  | 2.9      | Status         | Suggestive  |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15

[Medline Link](#)

Buck KJ, Finn DA. Genetic factors in addiction: QTL mapping and candidate gene studies implicate GABAergic genes in alcohol and barbiturate withdrawal in mice. Addiction. 2001 Jan;96(1):139-49.

[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.

[Medline Link](#)

# Calw4

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Calw4    |                |   |
| Chromosome | 1        | Class          | Alcohol   |
| Peak       | 20 cM    | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 12-35 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 0.0001   | Strains Tested | BXD RI, B6D2 F2   |
| LOD Score  | 3.1      | Status         | Suggestive  |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15

[Medline Link](#)

# Calw5

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Calw5    |                |   |
| Chromosome | 4        | Class          | Alcohol   |
| Peak       | 37 cM    | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 30-44 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 0.003    | Strains Tested | BXD RI, B6D2 F2   |
| LOD Score  | 2.0      | Status         | Suggestive  |

## References

Buck KJ, Rademacher BS, Metten P, Crabbe JC. Mapping murine loci for physical dependence on ethanol. Psychopharmacology (Berl). 2002 Apr;160(4):398-407. Epub 2002 Feb 15

[Medline Link](#)

Crabbe JC. Provisional mapping of quantitative trait loci for chronic ethanol withdrawal severity in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1998 Jul;286(1):263-71.

[Medline Link](#)

# Caws1

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Caws1    |                |   |
| Chromosome | 1        | Class          | Alcohol   |
| Peak       |          | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 28-47 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 3e-07    | Strains Tested | IWSP-2 x IWSR-1 F2  |
| LOD Score  | 6.5      | Status         | Significant   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. *Mamm Genome*. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws1/Alcw5

|                |   |                  |                            |              |                   |
|----------------|---|------------------|----------------------------|--------------|-------------------|
| QTL Code       | Caws1/Alcw5   | PARC Alias       | Caws1                      | MGI Alias    | Alcw5             |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |                   |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |                   |
| Chromosome     | 1   | Peak             |                            | Range        | 28-47             |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Significant       |
| LOD score      | 6.5   | p value          | 3e-07                      | Sex Specific |                   |
| Strains Tested | IWSP-2 x IWSR-1 F2  | Increaser Allele | P                          | Dominance    | Partial, R allele |
| Comments       |   |                  |                            |              |                   |
| Contact Person | J. Belknap  | Last Updated     | 2007-04-04 13:16:03        |              |                   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws2

|                |   |                  |                            |              |                   |
|----------------|---|------------------|----------------------------|--------------|-------------------|
| QTL Code       | Caws2   | PARC Alias       | Caws2                      | MGI Alias    |                   |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |                   |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |                   |
| Chromosome     | 4   | Peak             |                            | Range        | 33-51             |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Significant       |
| LOD score      | 2.3   | p value          | 0.005                      | Sex Specific |                   |
| Strains Tested | IWSP-2 x IWSR-1 F2  | Increaser Allele | P                          | Dominance    | Overdom, P allele |
| Comments       |   |                  |                            |              |                   |
| Contact Person | J. Belknap  | Last Updated     | 2007-04-04 13:23:53        |              |                   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws3

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Caws3    |                |   |
| Chromosome | 8        | Class          | Alcohol   |
| Peak       |          | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 16-43 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 0.0003   | Strains Tested | IWSP-2 x IWSR-1 F2  |
| LOD Score  | 3.5      | Status         | Significant   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws4

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Caws4    |                |   |
| Chromosome | 11       | Class          | Alcohol   |
| Peak       |          | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 15-36 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 2e-07    | Strains Tested | IWSP-2 x IWSR-1 F2  |
| LOD Score  | 6.7      | Status         | Significant   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws4/Alcw3

|                |   |                  |                            |              |                   |
|----------------|---|------------------|----------------------------|--------------|-------------------|
| QTL Code       | Caws4/Alcw3   | PARC Alias       | Caws4                      | MGI Alias    | Alcw3             |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |                   |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |                   |
| Chromosome     | 11  | Peak             |                            | Range        | 15-36             |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Significant       |
| LOD score      | 6.7   | p value          | 2e-07                      | Sex Specific | Stronger in males |
| Strains Tested | IWSP-2 x IWSR-1 F2  | Increaser Allele | P                          | Dominance    | Partial, R allele |
| Comments       |   |                  |                            |              |                   |
| Contact Person | J. Belknap  | Last Updated     | 2007-04-05 11:00:08        |              |                   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws5

|            |          |                |   |
|------------|----------|----------------|---|
| QTL Code   | Caws5    |                |   |
| Chromosome | 14       | Class          | Alcohol   |
| Peak       |          | Phenotype      | Chronic Alcohol Withdrawal  |
| Range      | 24-48 cM | Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |
| p Value    | 6e-05    | Strains Tested | IWSP-2 x IWSR-1 F2  |
| LOD Score  | 4.2      | Status         | Significant   |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Caws5/Alcw7

|                |   |                  |                            |              |             |
|----------------|---|------------------|----------------------------|--------------|-------------|
| QTL Code       | Caws5/Alcw7   | PARC Alias       | Caws5                      | MGI Alias    | Alcw7       |
| Drug           | Alcohol   | Phenotype        | Chronic Alcohol Withdrawal |              |             |
| Assay          | Area under the curve for handling-induced convulsions over 25 hr period following 72 hr exposure to ethanol vapor |                  |                            |              |             |
| Chromosome     | 14  | Peak             |                            | Range        | 24-48       |
| Max Range (Mb) |   | Min Range (Mb)   |                            | Status       | Significant |
| LOD score      | 4.2   | p value          | 6e-05                      | Sex Specific |             |
| Strains Tested | IWSP-2 x IWSR-1 F2  | Increaser Allele | P                          | Dominance    | None        |
| Comments       |   |                  |                            |              |             |
| Contact Person | J. Belknap  | Last Updated     | 2007-04-05 11:05:01        |              |             |

## References

Bergeson SE, Kyle Warren R, Crabbe JC, Metten P, Gene Erwin V, Belknap JK. Chromosomal loci influencing chronic alcohol withdrawal severity. Mamm Genome. 2003 Jul;14(7):454-63.

[Medline Link](#)

# Et2bec1

|            |         |                |                                    |
|------------|---------|----------------|------------------------------------|
| QTL Code   | Et2bec1 |                |                                    |
| Chromosome | 17      | Class          | Alcohol                            |
| Peak       | 6.27 cM | Phenotype      | Acute Ethanol Metabolism           |
| Range      |         | Assay          | BEC at 60 min after 2 g/kg ethanol |
| p Value    | 0.00013 | Strains Tested | B6, D2, BXD RI                     |
| LOD Score  |         | Status         | Suggestive                         |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et2becprov1

|                |                                    |                  |                          |              |            |
|----------------|------------------------------------|------------------|--------------------------|--------------|------------|
| QTL Code       | Et2becprov1                        | PARC Alias       | Et2bec1                  | MGI Alias    |            |
| Drug           | Alcohol                            | Phenotype        | Acute Ethanol Metabolism |              |            |
| Assay          | BEC at 60 min after 2 g/kg ethanol |                  |                          |              |            |
| Chromosome     | 17                                 | Peak             | 6.27                     | Range        |            |
| Max Range (Mb) |                                    | Min Range (Mb)   |                          | Status       | Suggestive |
| LOD score      |                                    | p value          | 0.00013                  | Sex Specific |            |
| Strains Tested | B6, D2, BXD RI                     | Increaser Allele | B6                       | Dominance    |            |
| Comments       |                                    |                  |                          |              |            |
| Contact Person | J. Crabbe                          | Last Updated     | 2007-04-10 12:33:43      |              |            |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et2met1

|            |          |                |                                     |
|------------|----------|----------------|-------------------------------------|
| QTL Code   | Et2met1  |                |                                     |
| Chromosome | 17       | Class          | Alcohol                             |
| Peak       | 6.27 cM  | Phenotype      | Acute Ethanol Metabolism            |
| Range      |          | Assay          | Metabolic rate after 2 g/kg ethanol |
| p Value    | 6.95e-06 | Strains Tested | B6, D2, BXD RI                      |
| LOD Score  |          | Status         | Significant                         |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et2met1/Etmr

|                |                                     |                  |                          |              |             |
|----------------|-------------------------------------|------------------|--------------------------|--------------|-------------|
| QTL Code       | Et2met1/Etmr                        | PARC Alias       | Et2met1                  | MGI Alias    | Etmr        |
| Drug           | Alcohol                             | Phenotype        | Acute Ethanol Metabolism |              |             |
| Assay          | Metabolic rate after 2 g/kg ethanol |                  |                          |              |             |
| Chromosome     | 17                                  | Peak             | 6.27                     | Range        |             |
| Max Range (Mb) |                                     | Min Range (Mb)   |                          | Status       | Significant |
| LOD score      |                                     | p value          | 6.95e-06                 | Sex Specific |             |
| Strains Tested | B6, D2, BXD RI                      | Increaser Allele | D2                       | Dominance    |             |
| Comments       |                                     |                  |                          |              |             |
| Contact Person | J. Crabbe                           | Last Updated     | 2007-04-10 12:32:04      |              |             |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et2met2

|            |         |                |                                     |
|------------|---------|----------------|-------------------------------------|
| QTL Code   | Et2met2 |                |                                     |
| Chromosome | 3       | Class          | Alcohol                             |
| Peak       | 49 cM   | Phenotype      | Acute Ethanol Metabolism            |
| Range      |         | Assay          | Metabolic rate after 2 g/kg ethanol |
| p Value    | 0.001   | Strains Tested | B6, D2, BXD RI                      |
| LOD Score  |         | Status         | Suggestive                          |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et2metprov1

|                |                                     |                  |                          |         |              |            |  |  |  |
|----------------|-------------------------------------|------------------|--------------------------|---------|--------------|------------|--|--|--|
| QTL Code       | Et2metprov1                         |                  | PARC Alias               | Et2met2 |              | MGI Alias  |  |  |  |
| Drug           | Alcohol                             | Phenotype        | Acute Ethanol Metabolism |         |              |            |  |  |  |
| Assay          | Metabolic rate after 2 g/kg ethanol |                  |                          |         |              |            |  |  |  |
| Chromosome     | 3                                   | Peak             | 49                       |         | Range        |            |  |  |  |
| Max Range (Mb) |                                     | Min Range (Mb)   |                          |         | Status       | Suggestive |  |  |  |
| LOD score      |                                     | p value          | 0.001                    |         | Sex Specific |            |  |  |  |
| Strains Tested | B6, D2, BXD RI                      | Increaser Allele | D2                       |         | Dominance    |            |  |  |  |
| Comments       |                                     |                  |                          |         |              |            |  |  |  |
| Contact Person | J. Crabbe                           | Last Updated     | 2007-04-10 12:27:53      |         |              |            |  |  |  |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et3bec1

|            |         |                |                                    |
|------------|---------|----------------|------------------------------------|
| QTL Code   | Et3bec1 |                |                                    |
| Chromosome | 17      | Class          | Alcohol                            |
| Peak       | 6.27 cM | Phenotype      | Acute Ethanol Metabolism           |
| Range      |         | Assay          | BEC at 60 min after 3 g/kg ethanol |
| p Value    | 0.00074 | Strains Tested | B6, D2, BXD RI                     |
| LOD Score  |         | Status         | Suggestive                         |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et3bec2

|            |         |                |                                     |
|------------|---------|----------------|-------------------------------------|
| QTL Code   | Et3bec2 |                |                                     |
| Chromosome | 15      | Class          | Alcohol                             |
| Peak       | 39 cM   | Phenotype      | Acute Ethanol Metabolism            |
| Range      |         | Assay          | BEC at 60 min. after 2 g/kg ethanol |
| p Value    | 0.001   | Strains Tested | B6, D2, BXD RI                      |
| LOD Score  |         | Status         | Suggestive                          |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et3bec3

|            |         |                |                                     |
|------------|---------|----------------|-------------------------------------|
| QTL Code   | Et3bec3 |                |                                     |
| Chromosome | 12      | Class          | Alcohol                             |
| Peak       | 52 cM   | Phenotype      | Acute Ethanol Metabolism            |
| Range      |         | Assay          | BEC at 60 min. after 3 g/kg ethanol |
| p Value    | 0.001   | Strains Tested | B6, D2, BXD RI                      |
| LOD Score  |         | Status         | Suggestive                          |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et3becprov1

|                |                                     |                  |                          |              |            |
|----------------|-------------------------------------|------------------|--------------------------|--------------|------------|
| QTL Code       | Et3becprov1                         | PARC Alias       | Et3bec3                  | MGI Alias    |            |
| Drug           | Alcohol                             | Phenotype        | Acute Ethanol Metabolism |              |            |
| Assay          | BEC at 60 min. after 3 g/kg ethanol |                  |                          |              |            |
| Chromosome     | 12                                  | Peak             | 52                       | Range        |            |
| Max Range (Mb) |                                     | Min Range (Mb)   |                          | Status       | Suggestive |
| LOD score      |                                     | p value          | 0.001                    | Sex Specific |            |
| Strains Tested | B6, D2, BXD RI                      | Increaser Allele | B6                       | Dominance    |            |
| Comments       |                                     |                  |                          |              |            |
| Contact Person | J. Crabbe                           | Last Updated     | 2007-04-10 12:28:41      |              |            |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et3becprov2

|                |                                     |                  |                          |              |            |           |  |  |  |
|----------------|-------------------------------------|------------------|--------------------------|--------------|------------|-----------|--|--|--|
| QTL Code       | Et3becprov2                         |                  | PARC Alias               | Et3bec2      |            | MGI Alias |  |  |  |
| Drug           | Alcohol                             | Phenotype        | Acute Ethanol Metabolism |              |            |           |  |  |  |
| Assay          | BEC at 60 min. after 2 g/kg ethanol |                  |                          |              |            |           |  |  |  |
| Chromosome     | 15                                  | Peak             | 39                       | Range        |            |           |  |  |  |
| Max Range (Mb) |                                     | Min Range (Mb)   |                          | Status       | Suggestive |           |  |  |  |
| LOD score      |                                     | p value          | 0.001                    | Sex Specific |            |           |  |  |  |
| Strains Tested | B6, D2, BXD RI                      | Increaser Allele |                          | Dominance    |            |           |  |  |  |
| Comments       |                                     |                  |                          |              |            |           |  |  |  |
| Contact Person | J. Crabbe                           | Last Updated     | 2007-04-10 12:30:12      |              |            |           |  |  |  |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et3becprov3

|                |                                    |                  |                          |              |            |
|----------------|------------------------------------|------------------|--------------------------|--------------|------------|
| QTL Code       | Et3becprov3                        | PARC Alias       | Et3bec1                  | MGI Alias    |            |
| Drug           | Alcohol                            | Phenotype        | Acute Ethanol Metabolism |              |            |
| Assay          | BEC at 60 min after 3 g/kg ethanol |                  |                          |              |            |
| Chromosome     | 17                                 | Peak             | 6.27                     | Range        |            |
| Max Range (Mb) |                                    | Min Range (Mb)   |                          | Status       | Suggestive |
| LOD score      |                                    | p value          | 0.00074                  | Sex Specific |            |
| Strains Tested | B6, D2, BXD RI                     | Increaser Allele | B6                       | Dominance    |            |
| Comments       |                                    |                  |                          |              |            |
| Contact Person | J. Crabbe                          | Last Updated     | 2007-04-10 12:34:17      |              |            |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et3met1

|            |          |                |                                     |
|------------|----------|----------------|-------------------------------------|
| QTL Code   | Et3met1  |                |                                     |
| Chromosome | 17       | Class          | Alcohol                             |
| Peak       | 6.27 cM  | Phenotype      | Acute Ethanol Metabolism            |
| Range      |          | Assay          | Metabolic rate after 3 g/kg ethanol |
| p Value    | 7.83e-05 | Strains Tested | B6, D2, BXD RI                      |
| LOD Score  |          | Status         | Significant                         |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et3met1/Etmr

|                |                                     |                  |                          |              |             |
|----------------|-------------------------------------|------------------|--------------------------|--------------|-------------|
| QTL Code       | Et3met1/Etmr                        | PARC Alias       | Et3met1                  | MGI Alias    | Etmr        |
| Drug           | Alcohol                             | Phenotype        | Acute Ethanol Metabolism |              |             |
| Assay          | Metabolic rate after 3 g/kg ethanol |                  |                          |              |             |
| Chromosome     | 17                                  | Peak             | 6.27                     | Range        |             |
| Max Range (Mb) |                                     | Min Range (Mb)   |                          | Status       | Significant |
| LOD score      |                                     | p value          | 7.83e-05                 | Sex Specific |             |
| Strains Tested | B6, D2, BXD RI                      | Increaser Allele | D2                       | Dominance    |             |
| Comments       |                                     |                  |                          |              |             |
| Contact Person | J. Crabbe                           | Last Updated     | 2007-04-10 12:32:34      |              |             |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Et3met2

|            |         |                |                                     |
|------------|---------|----------------|-------------------------------------|
| QTL Code   | Et3met2 |                |                                     |
| Chromosome | 1       | Class          | Alcohol                             |
| Peak       | 102 cM  | Phenotype      | Acute Ethanol Metabolism            |
| Range      |         | Assay          | Metabolic rate after 3 g/kg ethanol |
| p Value    | 0.001   | Strains Tested | B6, D2, BXD RI                      |
| LOD Score  |         | Status         | Suggestive                          |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res*. 2002 May;26(5):610-6.

[Medline Link](#)

# Et3metprov1

|                |                                     |                  |                          |              |            |
|----------------|-------------------------------------|------------------|--------------------------|--------------|------------|
| QTL Code       | Et3metprov1                         | PARC Alias       | Et3met2                  | MGI Alias    |            |
| Drug           | Alcohol                             | Phenotype        | Acute Ethanol Metabolism |              |            |
| Assay          | Metabolic rate after 3 g/kg ethanol |                  |                          |              |            |
| Chromosome     | 1                                   | Peak             | 102                      | Range        |            |
| Max Range (Mb) |                                     | Min Range (Mb)   |                          | Status       | Suggestive |
| LOD score      |                                     | p value          | 0.001                    | Sex Specific |            |
| Strains Tested | B6, D2, BXD RI                      | Increaser Allele | B6                       | Dominance    |            |
| Comments       |                                     |                  |                          |              |            |
| Contact Person | J. Crabbe                           | Last Updated     | 2007-04-10 12:27:12      |              |            |

## References

Grisel JE, Metten P, Wenger CD, Merrill CM, Crabbe JC. Mapping of quantitative trait loci underlying ethanol metabolism in BXD recombinant inbred mouse strains. *Alcohol Clin Exp Res.* 2002 May;26(5):610-6.

[Medline Link](#)

# Etacc

|            |          |                |  |
|------------|----------|----------------|--|
| QTL Code   | Etacc    |                |  |
| Chromosome | 15       | Class          | Alcohol  |
| Peak       | 30 cM    | Phenotype      | Alcohol Acceptance   |
| Range      | 15-48 cM | Assay          | Total amount of 10% (v/v) ethanol consumed over 24-h period following 24-h period of water deprivation |
| p Value    |          | Strains Tested | B6, D2, BXD RI, B6D2 F2  |
| LOD Score  | 3.8      | Status         | Significant  |

## References

McClearn GE, Tarantino LM, Rodriguez LA, Jones BC, Blizzard DA, Plomin R. Genotypic selection provides experimental confirmation for an alcohol consumption quantitative trait locus in mouse. Mol Psychiatry. 1997 Oct-Nov;2(6):486-9.

[Medline Link](#)

# Etact1

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Etact1                                  | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol                                 | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |   |                  |                             |              |             |
| Chromosome     | 2                                       | Peak             | 51 cM                       | Range        | 41-61       |
| Max Range (Mb) | 20.2 - 153.3                            | Min Range (Mb)   | 37.5 - 130.8                | Status       | Significant |
| LOD score      | 3.9                                     | p value          | 2.2e-05                     | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2                         | Increaser Allele |                             | Dominance    |             |
| Comments       | Mb intervals based on congenics studies |                  |                             |              |             |
| Contact Person | T. Phillips                             | Last Updated     | 2007-04-09 13:48:24         |              |             |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. *Genes Brain Behav.* 2006 Oct;5(7):506-17.

[Medline Link](#)

Phillips TJ, Huson M, Gwiazdon C, Burkhart-Kasch S, Shen EH. Effects of acute and repeated ethanol exposures on the locomotor activity of BXD recombinant inbred mice. *Alcohol Clin Exp Res.* 1995 Apr;19(2):269-78.

[Medline Link](#)

# Etact2

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Etact2                                  | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol                                 | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |   |                  |                             |              |             |
| Chromosome     | 9                                       | Peak             | 113.5 Mb                    | Range        | 51-71 cM    |
| Max Range (Mb) | 0 - 124                                 | Min Range (Mb)   | 32.6 - 101.4                | Status       | Significant |
| LOD score      | 3.6                                     | p value          | 4.3e-05                     | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2                         | Increaser Allele |                             | Dominance    |             |
| Comments       | Mb intervals based on congenics studies |                  |                             |              |             |
| Contact Person | T. Phillips                             | Last Updated     | 2007-04-09 14:54:31         |              |             |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. *Genes Brain Behav.* 2006 Oct;5(7):506-17.

[Medline Link](#)

Phillips TJ, Huson M, Gwiazdon C, Burkhart-Kasch S, Shen EH. Effects of acute and repeated ethanol exposures on the locomotor activity of BXD recombinant inbred mice. *Alcohol Clin Exp Res.* 1995 Apr;19(2):269-78.

[Medline Link](#)

Kamens HM, Phillips TJ; unpublished data

# Etact3

|                |                 |                  |                             |              |             |
|----------------|-----------------|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Etact3          | PARC Alias       |                             | MGI Alias    |             |
| Drug           | Alcohol         | Phenotype        | Alcohol Stimulated Activity |              |             |
| Assay          |                 |                  |                             |              |             |
| Chromosome     | 1               | Peak             | 62.6 Mb                     | Range        |             |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      | 0.0             | p value          | 0                           | Sex Specific |             |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele |                             | Dominance    |             |
| Comments       |                 |                  |                             |              |             |
| Contact Person | T. Phillips     | Last Updated     | 2007-04-09 13:37:52         |              |             |

## References

# Etact4

|            |         |                |                             |
|------------|---------|----------------|-----------------------------|
| QTL Code   | Etact4  |                |                             |
| Chromosome | 3       | Class          | Alcohol                     |
| Peak       | 19 cM   | Phenotype      | Ethanol-Stimulated Activity |
| Range      | 9-29 cM | Assay          |                             |
| p Value    | 0.0027  | Strains Tested | BXD RI, B6D2 F2             |
| LOD Score  | 2.0     | Status         | Suggestive                  |

## References

Phillips TJ, Huson M, Gwiazdon C, Burkhart-Kasch S, Shen EH. Effects of acute and repeated ethanol exposures on the locomotor activity of BXD recombinant inbred mice. *Alcohol Clin Exp Res.* 1995 Apr;19(2):269-78.

[Medline Link](#)

# Etactprov1

|                |             |                  |                             |              |            |
|----------------|-------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov1  | PARC Alias       | Etact3                      | MGI Alias    |            |
| Drug           | Alcohol     | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |             |                  |                             |              |            |
| Chromosome     | 2           | Peak             |                             | Range        | 73 - 93 cM |
| Max Range (Mb) |             | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |             | p value          |                             | Sex Specific |            |
| Strains Tested |             | Increaser Allele |                             | Dominance    |            |
| Comments       |             |                  |                             |              |            |
| Contact Person | T. Phillips | Last Updated     | 2007-04-09 13:58:53         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. *Genes Brain Behav.* 2006 Oct;5(7):506-17.

[Medline Link](#)

Phillips TJ, Huson M, Gwiazdon C, Burkhart-Kasch S, Shen EH. Effects of acute and repeated ethanol exposures on the locomotor activity of BXD recombinant inbred mice. *Alcohol Clin Exp Res.* 1995 Apr;19(2):269-78.

[Medline Link](#)

# Etactprov2

|                |                 |                  |                             |              |            |
|----------------|-----------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov2      | PARC Alias       | Etact4                      | MGI Alias    |            |
| Drug           | Alcohol         | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |                 |                  |                             |              |            |
| Chromosome     | 3               | Peak             | 19 cM                       | Range        | 9-29       |
| Max Range (Mb) |                 | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      | 2.0             | p value          | 0.0027                      | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2 | Increaser Allele |                             | Dominance    |            |
| Comments       |                 |                  |                             |              |            |
| Contact Person | T. Phillips     | Last Updated     | 2007-04-09 14:02:16         |              |            |

## References

Phillips TJ, Huson M, Gwiazdon C, Burkhardt-Kasch S, Shen EH. Effects of acute and repeated ethanol exposures on the locomotor activity of BXD recombinant inbred mice. *Alcohol Clin Exp Res.* 1995 Apr;19(2):269-78.

[Medline Link](#)

# Etactprov3

|                |             |                  |                             |              |            |
|----------------|-------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov3  | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol     | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |             |                  |                             |              |            |
| Chromosome     | 3           | Peak             | 18.8 Mb                     | Range        |            |
| Max Range (Mb) |             | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |             | p value          |                             | Sex Specific |            |
| Strains Tested |             | Increaser Allele |                             | Dominance    |            |
| Comments       |             |                  |                             |              |            |
| Contact Person | T. Phillips | Last Updated     | 2007-04-09 14:05:09         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. Genes Brain Behav. 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov4

|                |             |                  |                             |              |            |
|----------------|-------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov4  | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol     | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |             |                  |                             |              |            |
| Chromosome     | 5           | Peak             | 124.2 Mb                    | Range        |            |
| Max Range (Mb) |             | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |             | p value          |                             | Sex Specific |            |
| Strains Tested |             | Increaser Allele |                             | Dominance    |            |
| Comments       |             |                  |                             |              |            |
| Contact Person | T. Phillips | Last Updated     | 2007-04-09 14:17:51         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. Genes Brain Behav. 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov5

|                |             |                  |                             |              |            |
|----------------|-------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov5  | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol     | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |             |                  |                             |              |            |
| Chromosome     | 6           | Peak             | 36.5 Mb                     | Range        |            |
| Max Range (Mb) |             | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |             | p value          |                             | Sex Specific | Males only |
| Strains Tested |             | Increaser Allele |                             | Dominance    |            |
| Comments       |             |                  |                             |              |            |
| Contact Person | T. Phillips | Last Updated     | 2007-04-09 14:20:12         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. Genes Brain Behav. 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov6

|                |                            |                  |                             |              |            |
|----------------|----------------------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov6                 | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol                    | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |                            |                  |                             |              |            |
| Chromosome     | 7                          | Peak             | 91.1 Mb                     | Range        |            |
| Max Range (Mb) |                            | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |                            | p value          |                             | Sex Specific |            |
| Strains Tested |                            | Increaser Allele |                             | Dominance    |            |
| Comments       | Sex specific: females only |                  |                             |              |            |
| Contact Person | T. Phillips                | Last Updated     | 2007-04-09 14:49:17         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. *Genes Brain Behav.* 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov7

|                |                            |                  |                             |              |            |
|----------------|----------------------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov7                 | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol                    | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |                            |                  |                             |              |            |
| Chromosome     | 10                         | Peak             | 93.3 Mb                     | Range        |            |
| Max Range (Mb) |                            | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |                            | p value          |                             | Sex Specific |            |
| Strains Tested |                            | Increaser Allele |                             | Dominance    |            |
| Comments       | Sex specific: females only |                  |                             |              |            |
| Contact Person | T. Phillips                | Last Updated     | 2007-04-09 14:58:11         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. Genes Brain Behav. 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov8

|                |                            |                  |                             |              |            |
|----------------|----------------------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov8                 | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol                    | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |                            |                  |                             |              |            |
| Chromosome     | 13                         | Peak             | 85.5 Mb                     | Range        |            |
| Max Range (Mb) |                            | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |                            | p value          |                             | Sex Specific |            |
| Strains Tested |                            | Increaser Allele |                             | Dominance    |            |
| Comments       | Sex specific: females only |                  |                             |              |            |
| Contact Person | T. Phillips                | Last Updated     | 2007-04-09 15:07:43         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. *Genes Brain Behav.* 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov9

|                |                          |                  |                             |              |            |
|----------------|--------------------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov9               | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol                  | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |                          |                  |                             |              |            |
| Chromosome     | 16                       | Peak             | 6.1 Mb                      | Range        |            |
| Max Range (Mb) |                          | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |                          | p value          |                             | Sex Specific |            |
| Strains Tested |                          | Increaser Allele |                             | Dominance    |            |
| Comments       | Sex specific: males only |                  |                             |              |            |
| Contact Person | T. Phillips              | Last Updated     | 2007-04-09 15:11:00         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. *Genes Brain Behav.* 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etactprov10

|                |                            |                  |                             |              |            |
|----------------|----------------------------|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etactprov10                | PARC Alias       |                             | MGI Alias    |            |
| Drug           | Alcohol                    | Phenotype        | Alcohol Stimulated Activity |              |            |
| Assay          |                            |                  |                             |              |            |
| Chromosome     | 18                         | Peak             | 25 Mb                       | Range        |            |
| Max Range (Mb) |                            | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      |                            | p value          |                             | Sex Specific |            |
| Strains Tested |                            | Increaser Allele |                             | Dominance    |            |
| Comments       | Sex specific: females only |                  |                             |              |            |
| Contact Person | T. Phillips                | Last Updated     | 2007-04-09 15:13:01         |              |            |

## References

Palmer AA, Lessov-Schlaggar CN, Ponder CA, McKinnon CS, Phillips TJ. Sensitivity to the locomotor-stimulant effects of ethanol and allopregnanolone: a quantitative trait locus study of common genetic influence. Genes Brain Behav. 2006 Oct;5(7):506-17.

[Medline Link](#)

# Etcpp1

|            |        |                |                                      |
|------------|--------|----------------|--------------------------------------|
| QTL Code   | Etcpp1 |                |                                      |
| Chromosome | 7      | Class          | Alcohol                              |
| Peak       | 51 cM  | Phenotype      | Alcohol Conditioned Place Preference |
| Range      |        | Assay          |                                      |
| p Value    | 0.0004 | Strains Tested | BXD RI, B6 D2 F2                     |
| LOD Score  | 2.7    | Status         | Suggestive                           |

## References

Cunningham CL. Localization of genes influencing ethanol-induced conditioned place preference and locomotor activity in BXD recombinant inbred mice. Psychopharmacology (Berl). 1995 Jul;120(1):28-41.

[Medline Link](#)

# Etcpp2

|                |                  |                  |                                   |              |             |
|----------------|------------------|------------------|-----------------------------------|--------------|-------------|
| QTL Code       | Etcpp2           | PARC Alias       |                                   | MGI Alias    |             |
| Drug           | Alcohol          | Phenotype        | Alcohol Conditioned Place Prefer. |              |             |
| Assay          |                  |                  |                                   |              |             |
| Chromosome     | 8                | Peak             | 43                                | Range        |             |
| Max Range (Mb) |                  | Min Range (Mb)   |                                   | Status       | Provisional |
| LOD score      | 2.0              | p value          | 0.002                             | Sex Specific |             |
| Strains Tested | BXD RI, B6 D2 F2 | Increaser Allele |                                   | Dominance    |             |
| Comments       |                  |                  |                                   |              |             |
| Contact Person | C. Cunningham    | Last Updated     | 2007-01-17 00:00:00               |              |             |

## References

Cunningham CL. Localization of genes influencing ethanol-induced conditioned place preference and locomotor activity in BXD recombinant inbred mice. Psychopharmacology (Berl). 1995 Jul;120(1):28-41.

[Medline Link](#)

# Etcta1

|            |        |                |                                    |
|------------|--------|----------------|------------------------------------|
| QTL Code   | Etcta1 |                |                                    |
| Chromosome | 1      | Class          | Alcohol                            |
| Peak       | 73 cM  | Phenotype      | Alcohol Conditioned Taste Aversion |
| Range      |        | Assay          |                                    |
| p Value    | 6e-09  | Strains Tested | BXD RI, B6 D2 F2, selected lines   |
| LOD Score  | 7.0    | Status         | Significant                        |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etcta1/Etohcta1

|                |                                  |                  |                                    |              |             |
|----------------|----------------------------------|------------------|------------------------------------|--------------|-------------|
| QTL Code       | Etcta1/Etohcta1                  | PARC Alias       | Etcta1                             | MGI Alias    | Etohcta1    |
| Drug           | Alcohol                          | Phenotype        | Alcohol Conditioned Taste Aversion |              |             |
| Assay          |                                  |                  |                                    |              |             |
| Chromosome     | 1                                | Peak             | 73                                 | Range        |             |
| Max Range (Mb) |                                  | Min Range (Mb)   |                                    | Status       | Significant |
| LOD score      | 7.0                              | p value          | 6e-09                              | Sex Specific |             |
| Strains Tested | BXD RI, B6 D2 F2, selected lines | Increaser Allele |                                    | Dominance    |             |
| Comments       |                                  |                  |                                    |              |             |
| Contact Person | C. Cunningham                    | Last Updated     | 2007-04-10 12:54:22                |              |             |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etcta2

|            |        |                |                                    |
|------------|--------|----------------|------------------------------------|
| QTL Code   | Etcta2 |                |                                    |
| Chromosome | 4      | Class          | Alcohol                            |
| Peak       | 82 cM  | Phenotype      | Alcohol Conditioned Taste Aversion |
| Range      |        | Assay          |                                    |
| p Value    | 2e-09  | Strains Tested | BXD RI, B6 D2 F2, selected lines   |
| LOD Score  | 7.5    | Status         | Significant                        |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etcta2/Etohcta2

|                |                                  |                  |                                    |              |             |
|----------------|----------------------------------|------------------|------------------------------------|--------------|-------------|
| QTL Code       | Etcta2/Etohcta2                  | PARC Alias       | Etcta2                             | MGI Alias    | Etohcta2    |
| Drug           | Alcohol                          | Phenotype        | Alcohol Conditioned Taste Aversion |              |             |
| Assay          |                                  |                  |                                    |              |             |
| Chromosome     | 4                                | Peak             | 82                                 | Range        |             |
| Max Range (Mb) |                                  | Min Range (Mb)   |                                    | Status       | Significant |
| LOD score      | 7.5                              | p value          | 2e-09                              | Sex Specific |             |
| Strains Tested | BXD RI, B6 D2 F2, selected lines | Increaser Allele |                                    | Dominance    |             |
| Comments       |                                  |                  |                                    |              |             |
| Contact Person | C. Cunningham                    | Last Updated     | 2007-04-10 12:55:38                |              |             |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etcta3

|            |        |                |                                    |
|------------|--------|----------------|------------------------------------|
| QTL Code   | Etcta3 |                |                                    |
| Chromosome | 6      | Class          | Alcohol                            |
| Peak       | 57 cM  | Phenotype      | Alcohol Conditioned Taste Aversion |
| Range      |        | Assay          |                                    |
| p Value    | 0.0007 | Strains Tested | BXD RI, B6 D2 F2, selected lines   |
| LOD Score  | 2.2    | Status         | Suggestive                         |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etcta4

|            |        |                |                                    |
|------------|--------|----------------|------------------------------------|
| QTL Code   | Etcta4 |                |                                    |
| Chromosome | 7      | Class          | Alcohol                            |
| Peak       | 36 cM  | Phenotype      | Alcohol Conditioned Taste Aversion |
| Range      |        | Assay          |                                    |
| p Value    | 0.0003 | Strains Tested | BXD RI, B6 D2 F2, selected lines   |
| LOD Score  | 2.6    | Status         | Suggestive                         |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Ethyp2

|            |        |                |  |
|------------|--------|----------------|--|
| QTL Code   | Ethyp2 |                |  |
| Chromosome | 5      | Class          | Alcohol  |
| Peak       | 59 cM  | Phenotype      | Hypothermic Sensitivity  |
| Range      |        | Assay          | Average difference from baseline of rectal body temperature measured at 30 and 60 minutes after 3 g/kg ethanol on day 1 of testing |
| p Value    | 0.002  | Strains Tested | BXD RI, B6D2 F2  |
| LOD Score  | 2.1    | Status         | Suggestive   |

## References

Crabbe JC, Phillips TJ, Gallaher EJ, Crawshaw LI, Mitchell SR. Common genetic determinants of the ataxic and hypothermic effects of ethanol in BXD/Ty recombinant inbred mice: genetic correlations and quantitative trait loci. *J Pharmacol Exp Ther.* 1996 M

[Medline Link](#)

Crabbe JC, Belknap JK, Mitchell SR, Crawshaw LI. Quantitative trait loci mapping of genes that influence the sensitivity and tolerance to ethanol-induced hypothermia in BXD recombinant inbred mice. *J Pharmacol Exp Ther.* 1994 Apr;269(1):184-92.

[Medline Link](#)

# Ethypprov1

|                |  |                  |                         |              |            |
|----------------|--|------------------|-------------------------|--------------|------------|
| QTL Code       | Ethypprov1   | PARC Alias       | Ethyp2                  | MGI Alias    |            |
| Drug           | Alcohol  | Phenotype        | Hypothermic Sensitivity |              |            |
| Assay          | Average difference from baseline of rectal body temperature measured at 30 and 60 minutes after 3 g/kg ethanol on day 1 of testing |                  |                         |              |            |
| Chromosome     | 5  | Peak             | 59                      | Range        |            |
| Max Range (Mb) |  | Min Range (Mb)   |                         | Status       | Suggestive |
| LOD score      | 2.1  | p value          | 0.002                   | Sex Specific |            |
| Strains Tested | BXD RI, B6D2 F2  | Increaser Allele |                         | Dominance    |            |
| Comments       |  |                  |                         |              |            |
| Contact Person | J Crabbe   | Last Updated     | 2007-04-10 13:10:23     |              |            |

## References

Crabbe JC, Phillips TJ, Gallaher EJ, Crawshaw LI, Mitchell SR. Common genetic determinants of the ataxic and hypothermic effects of ethanol in BXD/Ty recombinant inbred mice: genetic correlations and quantitative trait loci. J Pharmacol Exp Ther. 1996 M

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Crabbe JC, Belknap JK, Mitchell SR, Crawshaw LI. Quantitative trait loci mapping of genes that influence the sensitivity and tolerance to ethanol-induced hypothermia in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1994 Apr;269(1):184-92.

[Medline Link](#)

# Ethypt2

|            |         |                |  |
|------------|---------|----------------|--|
| QTL Code   | Ethypt2 |                |  |
| Chromosome | 7       | Class          | Alcohol  |
| Peak       | 44 cM   | Phenotype      | Hypothermic Tolerance  |
| Range      |         | Assay          | Difference in hypothermic response between day 3 and day 1 after 3 g/kg ethanol each day |
| p Value    | 0.01    | Strains Tested | BXD RI, B6D2 F2  |
| LOD Score  | 1.4     | Status         | Provisional  |

## References

Crabbe JC, Phillips TJ, Gallaher EJ, Crawshaw LI, Mitchell SR. Common genetic determinants of the ataxic and hypothermic effects of ethanol in BXD/Ty recombinant inbred mice: genetic correlations and quantitative trait loci. J Pharmacol Exp Ther. 1996 M

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Crabbe JC, Belknap JK, Mitchell SR, Crawshaw LI. Quantitative trait loci mapping of genes that influence the sensitivity and tolerance to ethanol-induced hypothermia in BXD recombinant inbred mice. J Pharmacol Exp Ther. 1994 Apr;269(1):184-92.

[Medline Link](#)

# Etohcппрov1

|                |                  |                  |                                      |              |            |
|----------------|------------------|------------------|--------------------------------------|--------------|------------|
| QTL Code       | Etohcппрov1      | PARC Alias       | Etcpp1                               | MGI Alias    |            |
| Drug           | Alcohol          | Phenotype        | Alcohol Conditioned Place Preference |              |            |
| Assay          |                  |                  |                                      |              |            |
| Chromosome     | 7                | Peak             | 51                                   | Range        |            |
| Max Range (Mb) |                  | Min Range (Mb)   |                                      | Status       | Suggestive |
| LOD score      | 2.7              | p value          | 0.0004                               | Sex Specific |            |
| Strains Tested | BXD RI, B6 D2 F2 | Increaser Allele |                                      | Dominance    |            |
| Comments       |                  |                  |                                      |              |            |
| Contact Person | C. Cunningham    | Last Updated     | 2007-04-10 13:04:39                  |              |            |

## References

Cunningham CL. Localization of genes influencing ethanol-induced conditioned place preference and locomotor activity in BXD recombinant inbred mice. Psychopharmacology (Berl). 1995 Jul;120(1):28-41.

[Medline Link](#)

# Etohcprov1

|                |   |                  |                             |              |            |
|----------------|---|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etohcprov1  | PARC Alias       | Etp4                        | MGI Alias    |            |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |            |
| Assay          | Consumption of 10% ethanol with two-bottle choice |                  |                             |              |            |
| Chromosome     | 4   | Peak             | 59                          | Range        |            |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      | 2.4   | p value          | 0.0009                      | Sex Specific |            |
| Strains Tested | BXDRI, B6D2 F2, STSL                              | Increaser Allele |                             | Dominance    |            |
| Comments       |   |                  |                             |              |            |
| Contact Person | T. Phillips                                       | Last Updated     | 2007-05-11 11:02:33         |              |            |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

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Belknap JK, Richards SP, O'Toole LA, Helms ML, Phillips TJ. Short-term selective breeding as a tool for QTL mapping: ethanol preference drinking in mice. *Behav Genet.* 1997 Jan;27(1):55-66.

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Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

# Etohcprov2

|                |   |                  |                             |              |            |
|----------------|---|------------------|-----------------------------|--------------|------------|
| QTL Code       | Etohcprov2  | PARC Alias       | Etp6                        | MGI Alias    |            |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |            |
| Assay          | Consumption of 10% ethanol with two-bottle choice |                  |                             |              |            |
| Chromosome     | 15  | Peak             | 43                          | Range        |            |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Suggestive |
| LOD score      | 2.6   | p value          | 0.0005                      | Sex Specific |            |
| Strains Tested | BXDRI, B6D2 F2, STSL                              | Increaser Allele |                             | Dominance    |            |
| Comments       |   |                  |                             |              |            |
| Contact Person | T. Phillips                                       | Last Updated     | 2007-04-05 14:13:29         |              |            |

## References

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

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Belknap JK, Richards SP, O'Toole LA, Helms ML, Phillips TJ. Short-term selective breeding as a tool for QTL mapping: ethanol preference drinking in mice. *Behav Genet.* 1997 Jan;27(1):55-66.

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Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

# Etohctaprov1

|                |                                  |                  |                                    |              |            |
|----------------|----------------------------------|------------------|------------------------------------|--------------|------------|
| QTL Code       | Etohctaprov1                     | PARC Alias       | Etcta3                             | MGI Alias    |            |
| Drug           | Alcohol                          | Phenotype        | Alcohol Conditioned Taste Aversion |              |            |
| Assay          |                                  |                  |                                    |              |            |
| Chromosome     | 6                                | Peak             | 57                                 | Range        |            |
| Max Range (Mb) |                                  | Min Range (Mb)   |                                    | Status       | Suggestive |
| LOD score      | 2.2                              | p value          | 0.0007                             | Sex Specific |            |
| Strains Tested | BXD RI, B6 D2 F2, selected lines | Increaser Allele |                                    | Dominance    |            |
| Comments       |                                  |                  |                                    |              |            |
| Contact Person | C. Cunningham                    | Last Updated     | 2007-04-10 12:56:57                |              |            |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etohctaprov2

|                |                                  |                  |                                    |              |            |
|----------------|----------------------------------|------------------|------------------------------------|--------------|------------|
| QTL Code       | Etohctaprov2                     | PARC Alias       | Etcta4                             | MGI Alias    |            |
| Drug           | Alcohol                          | Phenotype        | Alcohol Conditioned Taste Aversion |              |            |
| Assay          |                                  |                  |                                    |              |            |
| Chromosome     | 7                                | Peak             | 36                                 | Range        |            |
| Max Range (Mb) |                                  | Min Range (Mb)   |                                    | Status       | Suggestive |
| LOD score      | 2.6                              | p value          | 0.0003                             | Sex Specific |            |
| Strains Tested | BXD RI, B6 D2 F2, selected lines | Increaser Allele |                                    | Dominance    |            |
| Comments       |                                  |                  |                                    |              |            |
| Contact Person | C. Cunningham                    | Last Updated     | 2007-04-10 12:57:17                |              |            |

## References

Risinger FO, Cunningham CL. Ethanol-induced conditioned taste aversion in BXD recombinant inbred mice. Alcohol Clin Exp Res. 1998 Sep;22(6):1234-44.

[Medline Link](#)

# Etp1

|            |       |                |   |
|------------|-------|----------------|---|
| QTL Code   | Etp1  |                |   |
| Chromosome | 2     | Class          | Alcohol   |
| Peak       | 28 cM | Phenotype      | Alcohol Preference Drinking                       |
| Range      |       | Assay          | Consumption of 10% ethanol with two-bottle choice |
| p Value    | 3e-05 | Strains Tested | BXDRI, B6D2 F2, STSL                              |
| LOD Score  | 3.8   | Status         | Significant                                       |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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[Medline Link](#)

# Etp1/Etohc1

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Etp1/Etohc1                                       | PARC Alias       | Etp1                        | MGI Alias    | Etohc1      |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |             |
| Assay          | Consumption of 10% ethanol with two-bottle choice |                  |                             |              |             |
| Chromosome     | 2   | Peak             | 28                          | Range        |             |
| Max Range (Mb) | 20.2 - 153.3                                      | Min Range (Mb)   | 37.5 - 130.8                | Status       | Significant |
| LOD score      | 3.8   | p value          | 3e-05                       | Sex Specific |             |
| Strains Tested | BXDRI, B6D2 F2, STSL                              | Increaser Allele | B6                          | Dominance    | Additive    |
| Comments       | Mb intervals based on congenics studies           |                  |                             |              |             |
| Contact Person | T. Phillips                                       | Last Updated     | 2007-04-05 12:28:06         |              |             |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

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Belknap JK, Richards SP, O'Toole LA, Helms ML, Phillips TJ. Short-term selective breeding as a tool for QTL mapping: ethanol preference drinking in mice. *Behav Genet.* 1997 Jan;27(1):55-66.

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Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

Phillips TJ; unpublished data

# Etp2

|            |        |                |   |
|------------|--------|----------------|---|
| QTL Code   | Etp2   |                |   |
| Chromosome | 2      | Class          | Alcohol   |
| Peak       | 49 cM  | Phenotype      | Alcohol Preference Drinking                       |
| Range      |        | Assay          | Consumption of 10% ethanol with two-bottle choice |
| p Value    | 0.0003 | Strains Tested | BXDRI, B6D2 F2, STSL                              |
| LOD Score  | 2.9    | Status         | Significant                                       |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

# Etp2/Etohc2

|                |   |                  |                             |              |              |
|----------------|---|------------------|-----------------------------|--------------|--------------|
| QTL Code       | Etp2/Etohc2                                       | PARC Alias       | Etp2                        | MGI Alias    | Etohc2       |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |              |
| Assay          | Consumption of 10% ethanol with two-bottle choice |                  |                             |              |              |
| Chromosome     | 2   | Peak             | 49                          | Range        |              |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Significant  |
| LOD score      | 2.9   | p value          | 0.0003                      | Sex Specific |              |
| Strains Tested | BXDRI, B6D2 F2, STSL                              | Increaser Allele |                             | Dominance    | B6 recessive |
| Comments       |   |                  |                             |              |              |
| Contact Person | T. Phillips                                       | Last Updated     | 2007-04-05 13:33:09         |              |              |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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[Medline Link](#)

Phillips TJ; unpublished data

# Etp3

|                |   |                  |                             |              |             |
|----------------|---|------------------|-----------------------------|--------------|-------------|
| QTL Code       | Etp3  | PARC Alias       | Etp3                        | MGI Alias    |             |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |             |
| Assay          | Consumption of 10% ethanol with two-bottle choice |                  |                             |              |             |
| Chromosome     | 3   | Peak             | 75                          | Range        |             |
| Max Range (Mb) |   | Min Range (Mb)   |                             | Status       | Significant |
| LOD score      | 3.0   | p value          | 0.0002                      | Sex Specific |             |
| Strains Tested | BXDRI, B6D2 F2, STSL                              | Increaser Allele |                             | Dominance    |             |
| Comments       |   |                  |                             |              |             |
| Contact Person | T. Phillips                                       | Last Updated     | 2007-04-05 13:35:40         |              |             |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

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Belknap JK, Richards SP, O'Toole LA, Helms ML, Phillips TJ. Short-term selective breeding as a tool for QTL mapping: ethanol preference drinking in mice. *Behav Genet.* 1997 Jan;27(1):55-66.

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Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

# Etp4

|            |        |                |   |
|------------|--------|----------------|---|
| QTL Code   | Etp4   |                |   |
| Chromosome | 4      | Class          | Alcohol   |
| Peak       | 59 cM  | Phenotype      | Alcohol Preference Drinking                       |
| Range      |        | Assay          | Consumption of 10% ethanol with two-bottle choice |
| p Value    | 0.0009 | Strains Tested | BXDRI, B6D2 F2, STSL                              |
| LOD Score  | 2.4    | Status         | Suggestive  |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

[Medline Link](#)

Belknap JK, Richards SP, O'Toole LA, Helms ML, Phillips TJ. Short-term selective breeding as a tool for QTL mapping: ethanol preference drinking in mice. *Behav Genet.* 1997 Jan;27(1):55-66.

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Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

# Etp5

|            |       |                |   |
|------------|-------|----------------|---|
| QTL Code   | Etp5  |                |   |
| Chromosome | 9     | Class          | Alcohol   |
| Peak       | 29 cM | Phenotype      | Alcohol Preference Drinking                       |
| Range      |       | Assay          | Consumption of 10% ethanol with two-bottle choice |
| p Value    | 3e-10 | Strains Tested | BXDRI, B6D2 F2, STSL                              |
| LOD Score  | 8.0   | Status         | Significant                                       |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

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Belknap JK, Richards SP, O'Toole LA, Helms ML, Phillips TJ. Short-term selective breeding as a tool for QTL mapping: ethanol preference drinking in mice. *Behav Genet.* 1997 Jan;27(1):55-66.

[Medline Link](#)

Phillips TJ, Crabbe JC, Metten P, Belknap JK. Localization of genes affecting alcohol drinking in mice. *Alcohol Clin Exp Res.* 1994 Aug;18(4):931-41.

[Medline Link](#)

# Etp5/Etohc3

|                |   |                  |                             |              |               |
|----------------|---|------------------|-----------------------------|--------------|---------------|
| QTL Code       | Etp5/Etohc3                                       | PARC Alias       | Etp5                        | MGI Alias    | Etohc3        |
| Drug           | Alcohol   | Phenotype        | Alcohol Preference Drinking |              |               |
| Assay          | Consumption of 10% ethanol with two-bottle choice |                  |                             |              |               |
| Chromosome     | 9   | Peak             | 29                          | Range        |               |
| Max Range (Mb) | 0 - 124   | Min Range (Mb)   | 32.6 - 101.4                | Status       | Significant   |
| LOD score      | 8.0   | p value          | 3e-10                       | Sex Specific |               |
| Strains Tested | BXDRI, B6D2 F2, STSL                              | Increaser Allele | B6                          | Dominance    | Additive/Free |
| Comments       | Mb intervals based on congenics studies           |                  |                             |              |               |
| Contact Person | T. Phillips                                       | Last Updated     | 2007-04-05 14:10:45         |              |               |

## References

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[Medline Link](#)

Kamens HM, Phillips TJ; unpublished data

# Etp6

|            |        |                |   |
|------------|--------|----------------|---|
| QTL Code   | Etp6   |                |   |
| Chromosome | 15     | Class          | Alcohol   |
| Peak       | 43 cM  | Phenotype      | Alcohol Preference Drinking                       |
| Range      |        | Assay          | Consumption of 10% ethanol with two-bottle choice |
| p Value    | 0.0005 | Strains Tested | BXDRI, B6D2 F2, STSL                              |
| LOD Score  | 2.6    | Status         | Suggestive  |

## References

Belknap JK, Atkins AL. The replicability of QTLs for murine alcohol preference drinking behavior across eight independent studies. *Mamm Genome.* 2001 Dec;12(12):893-9.

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Phillips TJ, Belknap JK, Buck KJ, Cunningham CL. Genes on mouse chromosomes 2 and 9 determine variation in ethanol consumption. *Mamm Genome.* 1998 Dec;9(12):936-41.

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[Medline Link](#)

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[Medline Link](#)

# Lore1

|                |                |                  |   |              |             |
|----------------|----------------|------------------|---|--------------|-------------|
| QTL Code       | Lore1          | PARC Alias       |   | MGI Alias    | Lore1       |
| Drug           | Alcohol        | Phenotype        | Alcohol-induced Loss of Righting Reflex |              |             |
| Assay          |                |                  |   |              |             |
| Chromosome     | 1              | Peak             |   | Range        | 36.9-41     |
| Max Range (Mb) |                | Min Range (Mb)   |   | Status       | Significant |
| LOD score      | 5.4            | p value          |   | Sex Specific |             |
| Strains Tested | ILS, ISS, ISCR | Increaser Allele |   | Dominance    |             |
| Comments       |                |                  |   |              |             |
| Contact Person | B. Bennett     | Last Updated     | 2007-04-10 12:35:59                     |              |             |

## References

Markel PD, Bennett B, Beeson M, Gordon L, Johnson TE. Confirmation of quantitative trait loci for ethanol sensitivity in long-sleep and short-sleep mice. Genome Res. 1997 Feb;7(2):92-9.

[Medline Link](#)

# Lore2

|                |                |                  |   |              |             |
|----------------|----------------|------------------|---|--------------|-------------|
| QTL Code       | Lore2          | PARC Alias       |   | MGI Alias    | Lore2       |
| Drug           | Alcohol        | Phenotype        | Alcohol-induced Loss of Righting Reflex |              |             |
| Assay          |                |                  |   |              |             |
| Chromosome     | 2              | Peak             |   | Range        | 75.6-81.7   |
| Max Range (Mb) |                | Min Range (Mb)   |   | Status       | Significant |
| LOD score      | 6.6            | p value          |   | Sex Specific |             |
| Strains Tested | ILS, ISS, ISCR | Increaser Allele |   | Dominance    |             |
| Comments       |                |                  |   |              |             |
| Contact Person | B. Bennett     | Last Updated     | 2007-04-10 12:36:10                     |              |             |

## References

Markel PD, Bennett B, Beeson M, Gordon L, Johnson TE. Confirmation of quantitative trait loci for ethanol sensitivity in long-sleep and short-sleep mice. Genome Res. 1997 Feb;7(2):92-9.

[Medline Link](#)

# Lore3

|                |                |                  |   |              |            |
|----------------|----------------|------------------|---|--------------|------------|
| QTL Code       | Lore3          | PARC Alias       |   | MGI Alias    | Lore3      |
| Drug           | Alcohol        | Phenotype        | Alcohol-induced Loss of Righting Reflex |              |            |
| Assay          |                |                  |   |              |            |
| Chromosome     | 8              | Peak             |   | Range        | 44-71      |
| Max Range (Mb) |                | Min Range (Mb)   |   | Status       | Suggestive |
| LOD score      | 3.4            | p value          |   | Sex Specific |            |
| Strains Tested | ILS, ISS, ISCR | Increaser Allele |   | Dominance    |            |
| Comments       |                |                  |   |              |            |
| Contact Person | B. Bennett     | Last Updated     | 2007-04-10 12:36:16                     |              |            |

## References

Markel PD, Bennett B, Beeson M, Gordon L, Johnson TE. Confirmation of quantitative trait loci for ethanol sensitivity in long-sleep and short-sleep mice. Genome Res. 1997 Feb;7(2):92-9.

[Medline Link](#)

# Lore4

|                |                |                  |   |              |             |
|----------------|----------------|------------------|---|--------------|-------------|
| QTL Code       | Lore4          | PARC Alias       |   | MGI Alias    | Lore4       |
| Drug           | Alcohol        | Phenotype        | Alcohol-induced Loss of Righting Reflex |              |             |
| Assay          |                |                  |   |              |             |
| Chromosome     | 11             | Peak             |   | Range        | 36-42       |
| Max Range (Mb) |                | Min Range (Mb)   |   | Status       | Significant |
| LOD score      | 6.5            | p value          |   | Sex Specific |             |
| Strains Tested | ILS, ISS, ISCR | Increaser Allele |   | Dominance    |             |
| Comments       |                |                  |   |              |             |
| Contact Person | B. Bennett     | Last Updated     | 2007-04-10 12:36:24                     |              |             |

## References

Markel PD, Bennett B, Beeson M, Gordon L, Johnson TE. Confirmation of quantitative trait loci for ethanol sensitivity in long-sleep and short-sleep mice. Genome Res. 1997 Feb;7(2):92-9.

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# Lore5

|                |                |                  |   |              |             |
|----------------|----------------|------------------|---|--------------|-------------|
| QTL Code       | Lore5          | PARC Alias       |   | MGI Alias    | Lore5       |
| Drug           | Alcohol        | Phenotype        | Alcohol-induced Loss of Righting Reflex |              |             |
| Assay          |                |                  |   |              |             |
| Chromosome     | 15             | Peak             |   | Range        | 18.8-61.7   |
| Max Range (Mb) |                | Min Range (Mb)   |   | Status       | Significant |
| LOD score      | 4.0            | p value          |   | Sex Specific |             |
| Strains Tested | ILS, ISS, ISCR | Increaser Allele |   | Dominance    |             |
| Comments       |                |                  |   |              |             |
| Contact Person | B. Bennett     | Last Updated     | 2007-04-10 12:36:30                     |              |             |

## References

Markel PD, Bennett B, Beeson M, Gordon L, Johnson TE. Confirmation of quantitative trait loci for ethanol sensitivity in long-sleep and short-sleep mice. Genome Res. 1997 Feb;7(2):92-9.

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