Filter on Fields from hg19.snp142CodingDbSnp

bin   is [ignored ▼]   [0]
chrom   [does ▼] match [*]   AND
chromStart is [ignored ▼]   [0]   AND
chromEnd   is [ignored ▼]   [0]   AND
name   [does ▼] match [*]   AND
transcript   [does ▼] match [*]   AND
frame   [does ▼] match [☑] * [ ] 1 [ ] 2 [ ] 3 [ ] n/a   AND
alleleCount is [ignored ▼]   [0]   AND
funcCodes   [does ▼] match [41]
alleles   [does ▼] match [*]
codons   [does ▼] match [*]
peptides   [does ▼] match [*]
[AND ▼] Free-form query: [   ]

[submit] [cancel]

hg19.snp142CodingDbSnp based filters

bin   is [ignored ▼]   [0]
chrom   [does ▼] match [*]   AND
chromStart is [ignored ▼]   [0]   AND
chromEnd   is [ignored ▼]   [0]   AND
name   [does ▼] match [*]   AND
transcript   [does ▼] match [*]   AND
frame   [does ▼] match [☑] * [ ] 1 [ ] 2 [ ] 3 [ ] n/a   AND
alleleCount is [ignored ▼]   [0]   AND
funcCodes   [does ▼] match [*]
alleles   [does ▼] match [*]
codons   [does ▼] match [*]
peptides   [does ▼] match [*]
[AND ▼] Free-form query: [   ]

[submit] [cancel]

Linked Tables

[ ] hg19	snp142	Polymorphism data from dbSNP
[ ] hg19	snp142Common	Polymorphism data from dbSNP
[ ] hg19	snp142Flagged	Polymorphism data from dbSNP
[ ] hg19	snp142Mult	Polymorphism data from dbSNP
[ ] hg19	snp142OrthoPt4Pa2Rm3	Orthologous alleles in chimp, orangutan and rhesus macaque (human only)
[ ] hg19	snp142Seq	Contains file offsets for reference SNP flanking sequences in a single fasta file.

[allow filtering using fields in checked tables]