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Re: Re-distribution of Blat

■ Jim Kent [kent@soe.ucsc.edu]

To:  Shepard, Samuel (CDC/OID/NCIRD)

Cc:

Tuesday, May 12, 2015 11:09 AM

Hi, that works. The change in 36 was minor, and only affected an edge case with a command line option:

36:

o (in 35x1) added repMatch default values for tileSize 16 to 18 in genoFind.c

On Mon, May 11, 2015 at 8:18 AM, Shepard, Samuel (CDC/OID/NCIRD) <vfn4@cdc.gov> wrote:

Dear Dr. Kent,

I have been using your fine software as one of the components in a deep sequencing assembly pipeline I have written. I would like to re-distribute the blat binary within the package. The pipeline is for *academic*, *non-profit*, and *government* use. I would:

- 1) cite Blat within our manuscript
- 2) package the Blat license information within an appropriate directory (including these correspondences)
- 3) make clear on our website the components of the pipeline that are for non-commercial use only

Please let me if this is acceptable.

Kind regards,

Sam Shepard

CDC/OID/NCIRD, Influenza Division

vfn4@cdc.gov,

P.S. I noticed Blat was updated to 3.6 recently from 3.5. Is there a change log available?