

From: Wen Yao <ywhzau@gmail.com>
Time: 2014/3/1 21:49
To: 'Joep de Ligt'
Subject: Re: Read functions for new tools

Dear Joep de Ligt,

I have released a new version of intansv which now support read in the result of Lumpy and SoftSearch. You can check it on the Bionconductor site.

If you have any suggestions, please contact me.

Best regards,
Wen

From: Joep de Ligt [[mailto:](mailto:ywhzau@gmail.com)]
Time: 2014/2/18 21:36
To: ywhzau@gmail.com
Subject: Read functions for new tools

Dear Wen,

First of all, thank you for creating the intansv package, it has been most hopeful in my research thus far.

I was wondering whether you are still maintaining the package.

If you are it would be great if you could include compatibility for new algorithms such as;

Lumpy; <http://arxiv.org/abs/1210.2342>

SoftSearch; <https://code.google.com/p/softsearch/>

These algorithms are more sensitive as they use multiple lines of evidence.

A good interface to combine and compare these datasets would be greatly helpful.

Kind regards,

Joep de Ligt

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