From: <u>Mason, Richard Esten</u>

To: <u>Bill Rooney</u>

Subject: Re: Fixed or random

Date: Monday, October 05, 2009 1:15:57 PM

That is about the same answer I have come up with. QTLs are usually treated as fixed, I believe, but there is justification to treat them as random also.

Lately I have seen more people starting to publish RILs as random, especially since PROC MIXED has become so popular.

Esten

---- Original Message -----

From: "Bill Rooney" < wlr@tamu.edu>

To: "Richard Esten' 'Mason' < richard-esten-mason@neo.tamu.edu>

Sent: Monday, October 5, 2009 12:23:53 PM GMT -06:00 US/Canada Central

Subject: RE: Fixed or random

Esten

I've seen people run and publish them as either, just depends on how you justify the analysis. I think you can easily justify a random effect, unless you start to focus on particular lines and particular regions of the genome.....

How's that for an answer?

Regards,

bill

-----Original Message-----

From: Mason, Richard Esten [mailto:richard-esten-mason@neo.tamu.edu]

Sent: Monday, October 05, 2009 11:18 AM

To: Bill Rooney

Subject: Fixed or random

Bill,

Would you say recombinant inbred lines have a random or fixed effect?

Esten

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