

From: [Young, Daniel W.](#)
To: ["Greg Porreca"; Waller, Patrick](#)
Cc: [Eric Boutin](#)
Subject: RE: G0871.70000US00; METHOD FOR REDUCING REPRESENTATIONAL BIAS IN MULTIPLEX AMPLIFICATION OF NUCLEIC ACIDS (Porreca)
Date: Thursday, April 30, 2009 10:13:34 AM

Greg,
Below is a revised claim based on our discussion yesterday. Are you available to talk with Patrick and me at 11AM today?
Thanks
Dan

1. A method of analyzing a plurality of genetic loci, the method comprising:
 - (i.) obtaining a plurality of molecular inversion probe capture products, each comprising a molecular inversion probe sequence, a polymorphic sequence and a source sequence,
 - (ii.) amplifying the plurality of molecular inversion probe capture products,
 - (iii.) determining numbers of occurrence of combinations of polymorphic sequence and source sequence of molecular inversion probe capture products in the amplified plurality, and
 - (iv.) if the number of occurrence of a specific combination of polymorphic sequence and source sequence exceeds a predetermined value, detecting bias in the amplification of the molecular inversion probe capture product comprising the specific combination.

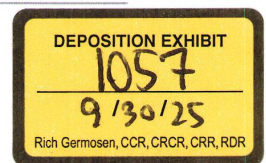
From: Greg Porreca [mailto:gporreca@gsgenetics.com]
Sent: Thursday, April 30, 2009 8:12 AM
To: Waller, Patrick
Cc: Young, Daniel W.; Eric Boutin
Subject: Re: G0871.70000US00; METHOD FOR REDUCING REPRESENTATIONAL BIAS IN MULTIPLEX AMPLIFICATION OF NUCLEIC ACIDS (Porreca)

Hi Patrick-
I'm available for a call any time. When is best for you?
-Greg

On Thu, Apr 30, 2009 at 7:53 AM, Patrick Waller <Patrick.Waller@wolfgreenfield.com> wrote:

Hi Greg,
Yes - we can file this application today. We were planning to file these applications as provisionals with the USPTO and this is probably the best strategy at this stage. However, we will call you to discuss this morning.
Patrick

From: Greg Porreca [mailto:gporreca@gsgenetics.com]
Sent: Wednesday, April 29, 2009 9:41 PM
To: Young, Daniel W.; Waller, Patrick
Cc: Eric Boutin



ML-IPR-000479

Subject: Re: G0871.70000US00; METHOD FOR REDUCING REPRESENTATIONAL BIAS IN MULTIPLEX AMPLIFICATION OF NUCLEIC ACIDS (Porreca)

Hi Dan and Patrick-

We've decided the disease gene list in this application is OK as is.

We think all other content is fine, and would like to proceed immediately to get this application on file in its current form. Could this go in to (I assume) WIPO on Thursday 4/30?

Thanks-
Greg

On Fri, Apr 24, 2009 at 4:37 PM, Daniel Young <Daniel.Young@wolfgreenfield.com> wrote:

Greg,

Attached is an updated version of the above-mentioned application for you to review. I will forward you a revised set of claims for Uri's application later today. In the meantime, could you send me a document containing the final figures for Uri's application.

Thanks

Dan

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ML-IPR-000480