SUMMARY STATEMENT

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(Privileged Communication)

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Application Number: 1 F32 CA180655-01

Boehm, Frederick J MD The Board of Regents of the University of Wisconsi **5730 Medical Sciences Center** 1300 University Ave Madison, WI 53706-1510

Review Group: ZRG1 F09A-L (20)

Center for Scientific Review Special Emphasis Panel

Fellowship: Oncology

Meeting Date: 03/14/2013

Council: MAY 2013 PCC: 1STR

Requested Start:

Project Title: Sample size planning methods in post-GWAS breast cancer eQTL studies

Requested: 1 year 7 months

Sponsor: NEWTON, MICHAEL A Department: Biostatistics/Med Info

Organization: UNIVERSITY OF WISCONSIN-MADISON

City, State: MADISON **WISCONSIN**

SRG Action: Impact Score: 33 Percentile: 37

Next Steps: Visit http://grants.nih.gov/grants/next_steps.htm

Human Subjects: 10-No human subjects involved

Animal Subjects: 10-No live vertebrate animals involved for competing appl.

1F32CA180655-01 Boehm, Frederick

RESUME AND SUMMARY OF DISCUSSION: The research project will explore sample size planning for eQTL studies in post-GWAS SNP characterization, where high dimensionality of genomic data complicates the eQTL study design. The training plan is comprehensive and includes regular meetings with the mentor and co-mentor. The applicant is strong; however, he has no first author research publications. The research strategy showed a number of weaknesses, which reduces enthusiasm for the fellowship. The proposal is written from a biostatistics viewpoint and validation of the initial results is not considered and the translational impact is not discussed. The research plan lacks details, which makes it difficult to evaluate. Overall, reviewers agree that the fellowship would have a moderate impact on the research career of the applicant.

DESCRIPTION (provided by applicant): Frederick Boehm, M.D. seeks to translate genome-wide association study (GWAS) results into clinical applications and to become an independent biomedical investigator. To achieve these goals, he will undertake a rigorous postdoctoral training program in biostatistical genomics under the mentorship of Michael Newton and Michael Gould. The research component of Boehm's program requires him to: 1) develop methods for expression quantitative trait locus (eQTL) study sample size planning with false discovery rate control in the setting of post-GWAS SNP characterization, and 2) develop sample size planning methods for pathway-based inference in eQTL studies. Boehm's approach uses hierarchical mixture modeling of variations in gene expression levels. To assess the statistical models, Boehm will apply his methods both in simulation studies and in analysis of pilot genomics data from 62 breast tissue samples. The proposed research will enable post-GWAS SNP characterization studies. Such studies are a critical intermediate step between GWAS results and their clinical application. These post-GWAS studies have the potential for major public health impacts by enabling personalized genomic medicine. Boehm's training plan includes courses in statistical theory and methods. Course topics include Bayesian methods, high-dimensional inference, survival analysis, and statistical consulting. These courses also will fulfill a requirement for the statistics Ph.D. program in which Boehm is a student. Boehm will supplement formal coursework with independent study of key texts and journal articles in multivariate and high-dimensional statistics. Professor Newton will meet regularly with Boehm to discuss questions that arise from his readings. To facilitate professional development throughout the fellowship period, Boehm will enroll in courses that address skills such as grant proposal writing, manuscript writing, research team management, and research mentoring. To refine mentoring skills, he will train a summer undergraduate researcher during his second year. Boehm also will undertake an ambitious program for training in responsible conduct of research that includes a capstone research ethics presentation to the university community.

PUBLIC HEALTH RELEVANCE: Our project will develop biostatistical tools to advance public health by enabling 1) discovery of novel biological targets for drugs to treat breast cancer and 2) development of markers for early detection of breast cancer. In this manner, our research is a translational bridge between recent advances in genetics technology and improvements in breast cancer diagnosis and treatment.

CRITIQUE 1:

Fellowship Applicant: 2

Sponsors, Collaborators, and Consultants: 1

Research Training Plan: 3

Training Potential: 1

Institutional Environment & Commitment to Training: 1

Overall Impact/Merit: Proper integration of GWAS findings with other genomic data will bridge the chasm between GWAS findings and personalized medicine implementation. A natural extension of

GWAS is studies that address whether disease-associated SNPs impact gene expression in relevant tissues. We use the term "eQTL", an abbreviation of "expression quantitative trait locus", to refer to SNPs whose genotypic variation associates with difference in mean expression level of one or more transcripts. While many investigators have undertaken eQTL studies in post-GWAS SNP characterization, high dimensionality of genomic data complicates eQTL study design. Sample size planning for eQTL studies remains an open question.

Aim 1: Develop methods to estimate eQTL study sample size with false discovery rate control in the setting of post-GWAS SNP characterization.

Aim 2: Develop sample size planning methods for pathway-based inference in eQTL studies. Completion of these aims will bridge a gap between GWAS results and their clinical applicability through personalized genomic medicine.

This research project is timely and of broad scientific interest. The overall impact is high, both from the perspective of the research project and the likelihood that the applicant will achieve independence.

1. Fellowship Applicant:

Strengths

- The PI Frederick Boehm completed an MD in 2007 and is currently working on a PhD in biostatistics – the mix of clinical knowledge and bio-statistical methodology is desired.
- Excellent academic record with many awards and honors.
- He has 4 publications and 4 clinical book chapters.
- Four very strong reference letters: Weir, Professor Dept. Biostatistics, Univ. Washington; Zhang, Professor, Dept. Statistics, Univ. Wisconsin-Madison; Crosslin, Senior Research Scientist U. Washington; (no name provided for fourth letter).
- Letter from Newton sponsor.

Weaknesses

- Only one abstract/presentation listed in 2000.
- Fourth reference letter not provided with the name of the referee.

2. Sponsors, Collaborators, and Consultants:

Strengths

- Sponsor: Michael Newton, Professor of Statistics with more than 80 publications and very good funding record.
- Co-sponsor: Michael Gould, Professor of Oncology with >190 publications.
- Both have had previous pre-and post-doctoral students who have moved onto independent research positions.
- Good mix of necessary skills biostatistics, molecular genetics and clinical.

Weaknesses

None.

3. Research Training Plan:

Strengths

Very timely topic – investigation of post-GWAS loci.

The sample size required for eQTL analysis is the subject of much debate.

Weaknesses

- Aim 2 was not well described.
- The overall proposal may benefit from access to additional databases which have both genotype and expression data.
- There was not enough description on the validation of the results.

4. Training Potential:

Strengths

- Excellent plan with good mix of coursework, seminars, collaborations with other scientists/physicians, teaching and communication skills.
- Regular meetings with mentor and co-mentor.
- Comprehensive approach.

Weaknesses

None.

5. Institutional Environment & Commitment to Training:

Strengths

Outstanding.

Weaknesses

None noted.

Protections for Human Subjects:

Not Applicable (No Human Subjects)

Data and Safety Monitoring Plan (Applicable for Clinical Trials Only):

Not Applicable (No Clinical Trials)

Vertebrate Animals:

Not Applicable (No Vertebrate Animals)

Biohazards:

Not Applicable (No Biohazards)

Training in the Responsible Conduct of Research:

Acceptable

Comments on Format (Required):

Comprehensive well laid out plan

Comments on Subject Matter (Required):

Adequate

Comments on Faculty Participation (Required):

Adequate

Comments on Duration (Required):

Adequate

Comments on Frequency (Required):

Adequate

Select Agents:

Not Applicable (No Select Agents)

Resource Sharing Plans:

- Acceptable
- They state they will freely share open-source code and accompanying tutorial via the Bioconductor software repository.

Budget and Period of Support:

Recommend as Requested

CRITIQUE 2:

Fellowship Applicant: 4

Sponsors, Collaborators, and Consultants: 1

Research Training Plan: 3 Training Potential: 3

Institutional Environment & Commitment to Training: 1

Overall Impact/Merit: The applicant proposes to develop methods for sample size estimation in eQTL analysis and pathway based eQTL inference. The applicant has had a complicated journey to graduate school which is not 100% reassuring. Enthusiasm is lessened based on confusion of how the applicant's MD benefits the work is biostatistics

1. Fellowship Applicant:

Strengths

- Outstanding undergraduate grades.
- Very good medical school and graduate school grades.

Weaknesses

- Unclear why applicant is leaving medicine.
- The personal statement discusses a program to focus on personalized medicine which is not revealed in the application to determine statistical power for eQTL analyses.

2. Sponsors, Collaborators, and Consultants:

Strengths

• Outstanding advisors and track records.

Weaknesses

None noted.

3. Research Training Plan:

Strengths

• Clear, achievable goals.

Weaknesses

- All method development, for a scientist rather than statistician this is surprising.
- No discussion of GETx.
- Lack of discussion on how a user of the tool would benefit.

4. Training Potential:

Strengths

Very good training plan that show strong commitment.

Weaknesses

None noted.

5. Institutional Environment & Commitment to Training:

Strengths

Outstanding.

Weaknesses

• None noted.

Protections for Human Subjects:

Acceptable Risks and Adequate Protections

Inclusion of Women, Minorities and Children:

- G2A Only Women, Acceptable
- M4A Minority Representation Unknown, Acceptable
- C4A Children Representation Unknown, Acceptable

Vertebrate Animals:

Not Applicable (No Vertebrate Animals)

Biohazards:

• Not Applicable (No Biohazards)

Training in the Responsible Conduct of Research:

Acceptable

Budget and Period of Support:

Recommend as Requested

CRITIQUE 3:

Fellowship Applicant: 3

Sponsors, Collaborators, and Consultants: 1

Research Training Plan: 2 Training Potential: 2

Institutional Environment & Commitment to Training: 1

Overall Impact/Merit: This F32 application is to develop biostatistical methods to apply GWAS results in clinical usage. The applicant is a Ph.D. student with an M.D. degree in population health sciences. The primary mentor is an expert in biostatistics, and the co-mentor is an established clinical oncologist. The research plan combines the strengths of both co-mentors, and is to develop biostatistical methods with strong potential in clinical usage. The training plan and career development plan are nicely laid out. This is an application with very high impact and training potential.

1. Fellowship Applicant:

Strengths

- The candidate obtained an M.D. degree in population health sciences at U Wisconsin Madison, and is now pursuing Ph.D. on Statistics at U Wisconsin Madison.
- The candidate is highly motivated to become a biostatistician.
- The applicant has a strong academic record in college, and has won numerous fellowships/awards.
- The applicant has written a number of review articles and book chapters.

Weaknesses

- The applicant has not published first author papers.
- The academic performance has declined in med/grad school.

2. Sponsors, Collaborators, and Consultants:

Strengths

- The sponsor, Dr. Michael Newton, is a full professor in the Department of Statistics and Biostatistics & Medical Informatics at the U Wisconsin Madison. Dr. Newton is an established investigator with expertise in biostatistics.
- Dr. Newton has a strong publication record and funding support, and has strong training experience.

BOEHM, F

• The co-sponsor, Dr. Michael Gould, is a full professor in the Department of Oncology. He is an expert in medical oncology of breast cancer, and has a strong publication and funding record.

Weaknesses

Not noted.

3. Research Training Plan:

Strengths

- The research is to develop biostatistical methods to apply genome-wide association study (GWAS) results into clinical applications, with focus on breast cancer.
- The research plan combines the expertise from both co-mentors.

Weaknesses

Not noted.

4. Training Potential:

Strengths

- The training plan and career development plan are very well laid out.
- The two co-mentors will make significant contributions to the applicant's growth.
- The applicant's medical background will largely facilitate his development in the field of biostatistics.

Weaknesses

Not noted.

5. Institutional Environment & Commitment to Training:

Strengths

• U Wisconsin Madison provides an outstanding environment.

Weaknesses

None noted.

Budget and Period of Support:

Recommended budget modifications or possible overlap identified:

THE FOLLOWING RESUME SECTIONS WERE PREPARED BY THE SCIENTIFIC REVIEW OFFICER TO SUMMARIZE THE OUTCOME OF DISCUSSIONS OF THE REVIEW COMMITTEE ON THE FOLLOWING ISSUES:

COMMITTEE BUDGET RECOMMENDATIONS: The budget was recommended as requested.

NIH has modified its policy regarding the receipt of resubmissions (amended applications). See Guide Notice NOT-OD-10-080 at http://grants.nih.gov/grants/guide/notice-files/NOT-OD-10-080.html.

The impact/priority score is calculated after discussion of an application by averaging the overall scores (1-9) given by all voting reviewers on the committee and multiplying by 10. The criterion scores are submitted prior to the meeting by the individual reviewers assigned to an application, and are not discussed specifically at the review meeting or calculated into the overall impact score. Some applications also receive a percentile ranking. For details on the review process, see http://grants.nih.gov/grants/peer_review_process.htm#scoring.

MEETING ROSTER

Center for Scientific Review Special Emphasis Panel CENTER FOR SCIENTIFIC REVIEW Fellowship: Oncology ZRG1 F09A-L (20) L March 14, 2013 - March 15, 2013

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