BIOGRAPHICAL SKETCH

Provide the following information for the key personnel and other significant contributors in the order listed on Form Page 2. Follow this format for each person. **DO NOT EXCEED FOUR PAGES.**

NAME Robert Martin Plenge eRA COMMONS USER NAME RPLENGE	POSITION TITLE Assistant Professor (Harvard Medical School) Director, Genetics and Genomics (BWH, Division of Rheumatology, Immunology and Allergy)
	Associate Member (Broad Institute, Inc.)

EDUCATION/TRAINING (Begin with baccalaureate or other initial professional education, such as nursing, and include postdoctoral training.)

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INSTITUTION AND LOCATION	DEGREE (if applicable)	YEAR(s)	FIELD OF STUDY
University of California, San Diego (UCSD)	B.S.	1988-1992	Biology (cum laude)
Case Western Reserve University (CWRU)	Ph.D., M.D.	1992-2000	Human Genetics
University of California, San Francisco (UCSF)		2000-2002	Internal Medicine
Harvard Medical School (HMS) and Brigham and Women's Hospital (BWH)		2002-2006	Clinical Rheumatology
Broad Institute of MIT and Harvard		2003-2008	Human Genetics

A. PERSONAL STATEMENT

For the current project, I will use my expertise in performing functional studies of the *CD40* rheumatoid arthritis (RA) risk allele and high-throughput drug screens. Dysregulation of the immune system is a fundamental disease mechanism, yet is incompletely understood. My laboratory applies human genetics and genomics tools to understand dysregulation of the immune system in the context of human disease [and rheumatoid arthritis (RA) in particular]. Our ultimate goal is to improve the lives of patients with immune-mediated diseases. Towards this end, my laboratory has three key research areas: (1) Discovery genetics/genomics: we use genome-wide approaches (genotyping and sequencing) to find DNA variants that are associated with immune-mediated traits, including disease susceptibility, response to therapy, and subclinical immune phenotypes (e.g., inflammation, autoantibody production); (2) Functional translation: once associated DNA variants are discovered, we use "humans as the model organism" to understand function and key biological pathways, with a long-term goal of disease prevention and drug discovery; we use genomic discoveries to guide high-throughput, small molecule drug screens; and (3) Clinical translation: we use clinical data from the electronic medical records (EMR) linked with biospecimens to enable clinical translational research.

B. POSITIONS AND HONORS

Positions and Employment

1994-1995	Howard Hughes Medical Student Fellow (CWRU, advisor = Dr. Huntington Willard)
1995-2000	Medical Scientist Training Program (CWRU, advisor = Dr. Huntington Willard)
2000-2002	Internal Medicine Resident, Department of Medicine (UCSF)
2002-2006	Clinical Rheumatology Fellow, Division of Rheumatology, Immunology and Allergy (BWH)
2003-2008	Post-doctoral Research Fellow, Broad Institute of MIT and Harvard (Dr. David Altshuler)
2006-2008	Instructor of Medicine, Division of Rheumatology, Immunology and Allergy (BWH)
2008-current	Assistant Professor of Medicine, Division of Rheumatology, Immunology and Allergy (BWH)
2008-current	Director, Genetics and Genomics, Division of Rheumatology, Immunology and Allergy (BWH)
2011-current	Associate Member. Broad Institute of MIT and Harvard

Other Experience and Professional Memberships

2000-2002	Molecular Medicine Research Fellowship Program (UCSF)
2002-current	Massachusetts Medical License
2004-current	Internal Medicine Board Certification
2006-current	Rheumatology Board Certification
2007-2012	Editorial Board, Arthritis Research and Therapy
2011-2012	Advisory Editor, Arthritis & Rheumatism
2012-current	Associate Editor, Arthritis & Rheumatism

2012-current Scientific Advisory Council, Research and Education Foundation of the ACR

2003-current Ad hoc manuscript reviewer: Nature, New England Journal of Medicine, Nature Genetics,

JAMA, PLoS Genetics, Am. Journal of Human Genetics, Arth & Rheum and >15 other journals

2009-current Ad hoc grant reviewer: NIH Genetics of Human Disease (GHD) Study Section, Psoriasis

Foundation, American College of Rheumatology "Within Our Reach" campaign, Alliance for

Lupus Research, EULAR "Orphan Disease Program", and other.

Selected Honors and Awards

2012

Phi Beta Delta International Honor Society (UCSD) 1992 1995 Alpha Omega Alpha Research Award (CWRU) 1995 Pre-doctoral Clinical Award, The American Society of Human Genetics 2007, 2008 Excellence in Tutoring Award, Human Genetics (HMS) Young Investigator Award, Department of Medicine (BWH) 2008 2008-current Burroughs Wellcome Fund Career Award for Medical Scientists

C. SELECTED PEER-REVIEWED PUBLICATIONS (from >70 PubMed references) Most relevant to the current application

Member, The American Society for Clinical Investigation

- Plenge RM*, Seielstad M*, Padyukov L, Lee AT, Remmers EF, Ding B, Liew A, Khalili H, Chandrasekaran A, Davies LRL, Li W, Tan AKS, Bonnard C, Ong RTH, Thalamuthu A, Pettersson S, Liu C, Tian C, Chen WV, Carulli J, Altshuler D, Alfredsson L, Criswell LA, Amos CI, Seldin MF, W, Kastner DA, Klareskog L*, and Gregersen PK*. (2007) Genome-Wide Search Identifies TRAF1-C5 as Rheumatoid Arthritis Risk Locus, New Eng J Med. Vol. 357 (12): 1199-209. *contributed equally. PMCID: PMC2636867
- 2. Plenge RM, Cotsapas C, Davies L, Price AL, et al (2007) Two independent alleles at 6q23 associated with risk of rheumatoid arthritis. Nat Genet, Vol 39 (12): 1477-82. PMCID: PMC2652744
- 3. Raychaudhuri S, Remmers EF, Lee AT, Hackett R, Guiducci C, ... Klareskog L, Gregersen PK, Daly MJ, Plenge RM (2008) Common variants at CD40 and other loci confer risk of rheumatoid arthritis. Nat Genet, 40(10):1216-23. PMCID: PMC2757650
- 4. Stahl EA, Raychaudhuri S, Remmers EF, Xie G, Eyre S, Thomson BP, ... Gregersen PK, Klareskog L, Plenge RM (2010) Genome-wide association study meta-analysis identifies seven new rheumatoid arthritis risk loci. Nat Genet 42(6):508-14. PMC Journal – In Process.
- 5. Stahl EA, Wegmann D, Kraft P, Chen R, Kallberg H, Kurreeman FAS, Gregersen PK, Alfredsson L, Siminovitch KA, Worthington K, de Bakker PlW, Raychaudhuri S*, Plenge RM* (2012) Bayesian inference reveals polygenic architecture of four common disease, Nat Genet. Mar 25. doi: PMID: 22446960

Additional recent publications of importance to the field (in chronological order)

- Plenge RM, Hendrich BD, Schwartz C, Arena JF, Naumova A, Sapienza C, Winter RM, Willard HF (1997) A promoter mutation in the XIST gene in two unrelated families with skewed X chromosome inactivation, Nat Genet Vol. 17: 353-356. PMID: 9354806
- 2. Plenge RM, Padyukov L, Remmers EF, et al. (2005) Replication of putative candidate gene associations with rheumatoid arthritis in over 4,000 samples from North America and Sweden: association of susceptibility with PTPN22, CTLA4 and PADI4, Am J Hum Genet Vol. 77: 1044-1060. PMCID: PMC1285162
- 3. Remmers EF*, Plenge RM*, Lee AT, Graham RR, Hom G, Behrens TW, de Bakker PIW, Le JM, Lee H, Batliwalla F, Li W, Masters SL, Booty MG, Carulli JP, Padyukov L, Alfredsson L, Klareskog L, Chen WV, Amos CI, Criswell LA, Seldin MF, Kastner DL, and Gregersen PK (2007) STAT4 and the risk of rheumatoid arthritis and systemic lupus erythematosus, New Eng J Med. Vol. 357 (10): 977-86. *contributed equally. PMCID: PMC2630215
- 4. Raychaudhuri S, Plenge RM, Rossin EJ, Ng AC; International Schizophrenia Consortium, Purcell SM, Sklar P. Scolnick EM, Xavier RJ, Altshuler D, Daly MJ (2009) Identifying Relationships Among Genomic Disease Regions: Predicting Genes at Pathogenic SNP Associations and Rare Deletions. PLoS Genetics 5(6):e1000534. [Epub 2009 Jun 26] PMCID: PMC2694358

- 5. Raychaudhuri S, Thomson BP, Remmers EF, Eyre S, Hinks A, Guiducci C, Catanese JJ, Xie G, Stahl EA,... **Plenge RM** (2009) *Genetic variants at CD28, PRDM1, and CD2/CD58 are associated* with rheumatoid arthritis risk. Nat Genet 41(12):1313-8. PMCID: PMC3142887
- 6. Karlson EW, Chibnik LB, Kraft P, Cui J, Keenan BT, Ding B, Raychaudhuri S, Klareskog L, Alfredsson L, Plenge RM (2010) *Cumulative association of 22 genetic variants with seropositive rheumatoid arthritis risk.* Ann Rheum Dis 69(6):1077-85. PMCID: PMC2933175
- 7. Cui J, Saevarsdottir S, Thomson B, Padyukov L, van der Helm-van Mil AH, Nititham J, Hughes LB, de Vries N,...**Plenge RM** (2010) *Rheumatoid arthritis risk allele PTPRC is also associated with response to anti-tumor necrosis factor alpha therapy.* Arthritis Rheum 62(7):1849-6. PMC Journal In Process.
- 8. Liao KP, Cai T, Gainer V, Goryachev S, Zeng-Treitler Q, Raychaudhuri S, Szolovits P, Churchill S, Murphy S, Kohane I, Karlson EW, **Plenge RM** (2010) *Electronic medical records for discovery research in rheumatoid arthritis*. Arthritis Care Res (Hoboken) 62(8): 1120–1127. PMCID: PMC3121049
- 9. Zhernakova A, Stahl EA, Trynka G, Raychaudhuri S, Festen EA, Franke L, Westra HJ, Fehrmann RS, Kurreeman FA, Thomson B, Gupta N, Romanos J, ...Huizinga TW, Wijmenga C, **Plenge RM** (2011) *Meta-analysis of genome-wide association studies in celiac disease and rheumatoid arthritis identifies fourteen non-HLA shared loci*. <u>PLoS Genet</u>. Feb;7(2):e1002004. [Epub 2011 Feb 24.] PMCID: PMC3044685
- 10. Kurreeman F, Liao K, Chibnik L, Hickey B, Stahl E, Gainer V, Li G, Bry L, Mahan S, Ardlie K, Thomson B, Szolovits P, Churchill S, Murphy SN, Cai T, Raychaudhuri S, Kohane I, Karlson E, Plenge RM (2011) Genetic Basis of Autoantibody Positive and Negative Rheumatoid Arthritis Risk in a Multi-ethnic Cohort Derived from Electronic Health Records, Am J Hum Genet. Jan 7;88(1):57-69. PMCID: PMC3014362

D. RESEARCH SUPPORT

Ongoing Research Support

<u>Career Award for Medical Scientists</u> Plenge (PI)

9/01/2008-8/31/2013

Burroughs Wellcome Fund

Genotype-phenotype studies of rheumatoid arthritis susceptibility genes

Goal: Career development award to facilitate the transition to an independent investigator; the research pertains to genotype-phenotype correlations of rheumatoid arthritis susceptibility genes

R01 AR056291 Choi (PI)

8/01/2008-7/31/2013

NIH-NIAMS

Genetics of Gout

Goal: Conduct a GWAS of uric acid and gout, incorporating environmental risk factors.

Role: co-investigator

R01 AR056768

Plenge (PI)

5/15/2009-4/30/2014

NIH-NIAMS

Dissection of two rheumatoid arthritis risk loci from genome-wide studies

Goal: Sequence two RA risk loci (TRAF1-C5 and TNFAIP3) for common and rare variants, and to conduct pilot functional studies of the risk alleles

U01 GM092691

Plenge (PI)

7/01/2010-6/30/2015

NIH-NIGMS

Genetic predictors of response to anti-TNF therapy in rheumatoid arthritis

Goal: GWAS genotype new samples and develop new statistical methodologies for analysis of response to anti-TNF therapy in patients with RA

Role: PI

R01_AR059648

Plenge (PI)

8/01/2010-6/30/2012

NIH-NIAMS

Deep Re-Sequencing of Rheumatoid Arthritis Risk Loci

Goal: Next-generation sequencing of coding exons from 30 genes identified by GWAS

Role: PI

U54 LM 008748 Kohane (PI) 9/15/10-9/14/14

NIH National Center for Biomedical Computing

Create an informatics infrastructure to mine electronic medical records for discovery research purposes.

Role: PI on sub-contract

<u>U54 HL 117798</u> FitzGerald (PI) 5/01/2012-4/30/2014

NIGMS

Personalization of therapeutic efficacy and risk

Goal: Conduct a 2-year pilot project linking clinical data from the EMR to discarded biospecimens.

Role: PI on sub-contract

Harvard Catalyst Child Health Plenge (PI) 5/01/2012-4/30/2013

Harvard Medical School

Novel autoantigen discovery with a synthetic human peptidome in juvenile idiopathic arthritis

Goal: Pilot project to discover autoantigens and correlate with clinical features in JIA.

Role: PI

Broad SPARC/Arthritis Foundation Plenge (PI) 9/01/2012-8/31/2013

Broad Institute and the Arthritis Foundation

Genomic analysis in the arthritis internet registry (AIR) cohort

Goal: Pilot project to perform GWAS and expression analysis in 300 AIR patients

Role: PI

Recently Completed Research Support (selected)

K08 Al55314 Plenge (PI) 5/01/04-1/31/09

NIH-NIAID

The goal of this mentored training grant was to study the genetic basis of the immune response.

U54 RR020278 Plenge (PI) 9/01/06-08/31/07

National Center for Research Resources (NCRR)

The goal was to perform subsidized genotyping of a GWAS of rheumatoid arthritis risk.

Gate's Foundation, #43307 Walker (PI) 9/01/07-09/30/08

Gate's Foundation

The goal of this study was to conduct a GWAS to search for genetic predictors of HIV viral load

Role: co-investigator on the subcontract

2008A053376 Plenge (PI) 5/1/08-4/30/09

Hearst Young Investigator Award:

The goal of this career award was to study the clinical and biological significance of RA risk alleles.

<u>U54 LM 008748</u> Kohane (PI) 8/01/07-7/31/10

NIH National Center for Biomedical Computing

Create an informatics infrastructure to mine electronic medical records for discovery research purposes.

Role: PI on sub-contract

Research and Education Foundation Plenge (PI) 7/01/08-6/31/10

American College of Rheumatology

Conduct a candidate gene and pilot GWAS of response to anti-TNF therapy in rheumatoid arthritis.

R01 AR057108 Plenge (PI) 5/12/2009-4/30/2011

NIH-NIAMS

Meta-analysis of GWAS data to identify novel rheumatoid arthritis risk loci.

Goal: Perform a meta-analysis of GWAS data to identify novel rheumatoid arthritis risk loci