

Supplementary Materials for

Systematic Analysis of Challenge-Driven Improvements in Molecular Prognostic Models for Breast Cancer

Adam A. Margolin,* Erhan Bilal, Erich Huang, Thea C. Norman, Lars Ottestad, Brigham H. Mecham, Ben Sauerwine, Michael R. Kellen, Lara M. Mangravite, Matthew D. Furia, Hans Kristian Moen Vollan, Oscar M. Rueda, Justin Guinney, Nicole A. Deflaux, Bruce Hoff, Xavier Schildwachter, Hege G. Russnes, Daehoon Park, Veronica O. Vang, Tyler Pirtle, Lamia Youseff, Craig Citro, Christina Curtis, Vessela N. Kristensen, Joseph Hellerstein, Stephen H. Friend,* Gustavo Stolovitzky, Samuel Aparicio, Carlos Caldas, Anne-Lise Børresen-Dale

*Corresponding author. E-mail: margolin@sagebase.org (A.A.M.); friend@sagebase.org (S.H.F.)

Published 17 April 2013, *Sci. Transl. Med.* **5**, 181re1 (2013) DOI: 10.1126/scitranslmed.3006112

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SUPPLEMENTARY MATERIALS

Materials and Methods

OsloVal data generation

Patient material: The OsloVal cohort consists of fresh frozen primary tumors from 184 breast cancer patients collected from 1981 to 99 (148 from 1981-89 and 36 from 1994-99) at the Norwegian Radium Hospital. Tumor material was collected for hormone receptor analysis by DCC (dextran coated charcoal) and excess tumor material was stored in a biobank (at -80°C). Frozen tissue samples were first cut in three parts, and tissue sections made from each (6 μm) for later evaluations using IHC. Tissue-Tek was removed from the specimens and all tissue were sliced, mixed and homogenized, and tissue aliquots stored at -80°C. Two tissue fractions of approximately 15 mg each were used for DNA and RNA isolation and analyses.

mRNA extraction: Extractions were performed on the QIA symphony SP robot from Qiagen. A total of 400 μl RLT buffer was added to the samples while on dry ice, followed by homogenization (Tissuelyzer). The QIA symphony RNA Kit cat# 931636 from Qiagen was used. Extracted RNA was quantified using Nanodrop 1000 and the RNA integrity determined using the Agilent Bioanalyzer.

DNA extraction: Extractions_were performed on the QIAsymphony SP robot from Qiagen. A total of 20μ l Proteinase K and 180μ l ATL buffer was added to the tissue aliquot and lysed for three hours at 56 degrees on a Thermomixer with 900 rpm, followed by addition of 4μ l RNase A and incubation at RT for 2 min. The extraction kit used was the QIAsymphony DNA mini Kit cat#931236 from Qiagen,

CNA analyses: A total of 500 ng Genomic DNA was used for the Affymetrix Genome-Wide Human SNP 6.0 Assay, and processed according to the manufacturer's protocol using the Affymetrix GW Human SNP Nsp/Sty 6.0 assay kit (prod. # 901015). Fragmented and end labeled PCR product were hybridized to the Affymetrix Human SNP 6.0 array (prod. # 901150) followed by incubation for 16 h at 50^oC at constant rotation (60 rpm). The washing and staining procedure was performed in the Affymetrix Fluidics Station 450. The arrays were scanned at 560 nm using a confocal laser-scanning microscope (Affymetrix Scanner 3000 7G). The SNP calls and Copy number analysis were done using the Affymetrix Genotyping Console Software ver 4.0.

Expression analysis: A total of_100 ng total RNA was used for cRNA preparation using the Ambion Illumina Total Prep kit (prod. # AMIL1791). The samples were hybridized to Illumina HT-12 v4 BeadChips, incubated 16h at 580C and subsequently scanned using the Illumina BeadStation.

Clinical and pathology variables: Routine tissue sections were retrieved for all patient samples (hematoxylin and eosin stained). Histological type was assessed according to WHO classification (*45*) and then histological grade were determined (*46*). ER and PR status were scored using mRNA expression data. HER2 amplification status was scored from SNP6 arrays. Survival time was collected from the national registries

Data availability

We provide the METABRIC and OsloVal data described in the paper via two mechanisms. Each mechanism is under different Terms of Use described in the links, be cognizant that these are patient data and therefore fall under data governance rules:

(A) For a 6-month post-Challenge "Validation Phase", we provide registered participants access to the data to re-evaluate their models

https://synapse.prod.sagebase.org/#!Synapse:syn1710250

(B) For those interested in accessing these data for independent research, please use these links for further information on how to obtain access to these protected health data
METABRIC: <u>https://synapse.prod.sagebase.org/#!Search:syn1688369</u>
OsloVal: <u>https://synapse.prod.sagebase.org/#!Search:syn1688370</u>

Table S1: Univariate and multivariate Cox regression statistics for the clinical covariates in the METABRIC data set.

	Multivariate Cox regression				Univariate Cox regression					
Covariate	coef	exp(coef)	se(coef)	Z	Pr(> z)	coef	exp(coef)	se(coef)	Z	Pr(> z)
Age at diagnosis	0.03946	1.040249	0.00349	11.3074	0	0.028595	1.029008	0.002929	9.764013	0
Tumor size	0.010088	1.010139	0.001815	5.557924	2.73E-08	0.014921	1.015032	0.001511	9.872802	0
Lymph nodes	0.053953	1.055435	0.006378	8.459218	0	0.06636	1.068612	0.004943	13.42513	0
Intermediate grade	0.217714	1.243232	0.150131	1.450162	0.147013	0.294219	1.342078	0.148386	1.982796	0.04739
High grade	0.313008	1.367533	0.152654	2.050437	0.040322	0.604817	1.830917	0.144895	4.174163	2.99E-05
ER negative	0.145647	1.156787	0.115006	1.266429	0.205359	0.320903	1.378372	0.076893	4.173364	3.00E-05
PR negative	0.145485	1.1566	0.08454	1.720888	0.085271	0.369708	1.447311	0.067268	5.496077	3.88E-08
HER2 negative	-0.25087	0.778123	0.103818	-2.41646	0.015672	-0.46318	0.62928	0.093925	-4.93135	8.17E-07
CT/HT	-0.08126	0.921953	0.334094	-0.24323	0.807829	-0.22345	0.799753	0.319135	-0.70018	0.483813
CT/HT/RT	-0.77431	0.461022	0.22377	-3.4603	0.00054	-0.55967	0.571396	0.215676	-2.59497	0.00946
CT/RT	-0.35825	0.698896	0.21013	-1.70491	0.088211	-0.27657	0.758382	0.204537	-1.35217	0.176322
HT	-0.906	0.404138	0.215142	-4.21115	2.54E-05	-0.56977	0.565655	0.187623	-3.03678	0.002391
HT/RT	-1.0773	0.340514	0.210783	-5.11095	3.21E-07	-0.76561	0.465052	0.18511	-4.13596	3.53E-05
RT	-1.24311	0.288487	0.226251	-5.49438	3.92E-08	-1.32218	0.266553	0.207634	-6.36785	1.92E-10
No treatment	-0.89846	0.407194	0.217971	-4.12194	3.76E-05	-0.86439	0.421307	0.193406	-4.46932	7.85E-06

Treatment:

CT - Chemo therapy

RT - Radio therapy

HT - Hormonal therapy

Table S2: Univariate and multivariate Cox regression statistics for the clinical covariates in the OsloVal data set.

	Multivariate Cox regression				Univariate Cox regression					
Covariate	coef	exp(coef)	se(coef)	Z	Pr(> z)	coef	exp(coef)	se(coef)	Z	Pr(> z)
Age at diagnosis	0.038351	1.039096	0.015439	2.483999	0.012992	0.034676	1.035285	0.007749	4.475184	7.63E-06
Tumor size	0.086823	1.090704	0.119527	0.726388	0.467601	0.166058	1.180641	0.060727	2.73449	0.006248
Lymph nodes	0.145669	1.156814	0.034743	4.192749	2.76E-05	0.132847	1.142075	0.017272	7.69146	1.45E-14
Intermediate grade	0.296181	1.344713	0.838911	0.353054	0.724048	0.775138	2.170891	0.470543	1.647326	0.099491
High grade	0.568608	1.765807	0.848445	0.670177	0.502745	0.846257	2.330906	0.474803	1.782331	0.074695
ER negative	0.947331	2.578818	0.409886	2.311205	0.020822	0.298188	1.347415	0.177244	1.682354	0.0925
PR negative	-0.22781	0.796275	0.406875	-0.5599	0.575546	0.311458	1.365415	0.219119	1.421411	0.155197
HER2 negative	0.221261	1.24765	0.46046	0.480522	0.630856	-0.19023	0.826768	0.254062	-0.74876	0.454002
СТ/НТ	0.677686	1.969316	0.43508	1.557614	0.119325	0.902983	2.466951	0.294647	3.064628	0.002179
CT/HT/RT	0.346296	1.413821	0.566452	0.611342	0.540973	1.040813	2.831519	0.317192	3.281334	0.001033
CT/RT	-0.43961	0.644287	0.812231	-0.54124	0.588343	-0.09935	0.905427	0.609823	-0.16291	0.870586
HT	0.240398	1.271756	0.616252	0.390097	0.696464	1.240686	3.457985	0.370471	3.348945	0.000811
HT/RT	-16.0705	1.05E-07	4503.05	-0.00357	0.997153	-0.49426	0.610025	1.019176	-0.48496	0.627707
RT	0.168161	1.183127	1.12635	0.149297	0.881319	0.13017	1.139022	0.73491	0.177123	0.859412

Treatment:

CT - Chemo therapy

RT - Radio therapy

HT - Hormonal therapy

Table S3: The Breast Cancer Challenge Consortium: Challenge participants who submitted amodel to phase 3 of the BCC.

First@Name	Last [®] Name	Affiliation	Email		
Miika🛛	Ahdesmäki	Almac@iagnostics,@Almac@roup,@craigavon,@T6355QD,@united%Kingdom	miika.ahdesmaki@almacgroup.com		
Robert	Atlas	Department®ftComputertSciences,IUniversity®ftBVisconsin,IMadison,IWIES3717,IUSA	ratlas@cs.wisc.edu		
Nikolay	Balov	Department®fBiostatistics®ndIComputationalBiology University®fBrochesterIMedicalICenter,IRochester,INYII.4642,IUSA	nikibalov@gmail.com		
Bonnie	Berger	Departments @filMathematics @nd Electrical Engineering Computer Science Computer Science @nd Artificial Intelligence Laboratory, Massach usetts Institute Df Intelligence Laboratory, Intelligence	bab@csail.mit.edu		
Archit	Bhise	MassachusettsInstituteIIIfTechnology,ICambridge,IMAII2139-4307,IUSA	archit@MIT.EDU		
Eric	Bonnet	(1) (1) বাগ্য যৈ যে যে হৈ বিৰুদ্ধে বিশেষ সিমান সিমান প্ৰায় বিশেষ সিমান প্ৰ প্ৰথম সিমান প্ৰ প্ৰ প্ৰ প্ৰ প্ৰ স্ট প্ৰ	eric.bonnet@curie.fr		
Aaron	Boudreau	Department@fllaboratory@Medicine,llniversity@fllalifornia,lSan@rancisco,lSan@ Francisco,lLa@4143,llSA	aaron.boudreau@ucsf.edu		
Chunhui	Cai	Department Bf Biomedical Informatics, University Bf Pittsburgh, PA 25206-3701, USA	chunhuic@pitt.edu		
Yifei	Chen	Department@ffComputertScience,BUniversity@ffCalifornia,Brvine,Brvine,CA1926972 USA	yifeic@uci.edu		
Jie	Cheng	QuantitativeEsciences,EGlaxoSmithKline,Ecollegeville,IPAE9426,IUSA	jcheng88@gmail.com		
Sean	Cory	Goodman ICancer Research ICentre, IMcGill University, II.160 Pine Avenue West, Montreal, IQuebec, IH3AIA3, ICanada	sean.cory@gmail.com		
Edmund3.	Crampin	Melbourne怎chool@fi歪ngineering,ITheIUniversity@fIMelbourne,IParkville,IVictoria,I Australia	e.crampin@auckland.ac.nz		
Chad 🖪 🛛	Creighton	Division@fBiostatistics,@Danal.@DuncanaCanceraCenter,BayloraCollege@fBMedicine,2 Houston,Zexas	creighto@bcm.edu		
Benjamin	Haibe-Kains	Institut@le@Recherches@liniques@le@Montréal,@RCM,@Montreal,@Quebec,@Canada	bhaibeka@ircm.qc.ca		
Thomas	Kelder	TNO, Microbiology and Systems Biology, Zeist, The Netherlands	thomaskelder@gmail.com		
Jeff	Knisley	Institutefor @uantitativeBiology East&ennessee & tate@university, @ohnson & ity, @N&7614, @USA&	knisleyj.etsu@gmail.com		
Vincenzo	Lagani	Bioinformatics型.aboratory,≣nstitute™f⊠Computer™cience理∓oundation™or®research型 and@technology@FORTH),IN.IPlastira型.003/assilika型/outon,IGR-70021.33Heraklion,I2 Crete,IGGreece	vlagani@ics.forth.gr		
Kai∄Yeung	Lau	AmgenInc,ISeattle,I98119BWA,IUSA	kaiyeung.lau@gmail.com		
Xinghua	Lu	Department@fiBiomedicalInformatics,IUniversity@fiPittsburgh,IPAII5206-3701,IUSA	xinghua@pitt.edu		
Songjian	Lu	Department@fiBiomedicalInformatics,IUniversity@fiPittsburgh,IPAII5206-3701,IUSA	songjian@pitt.edu		
Jian	Peng	Computert&ciencetandtantitikiallantelligencetaboratory,aMassachusettsanstitutetaffa Technology,,aCambridge,aMAt02139,aUSA	jpeng@csail.mit.edu		
Robert ³ .	Prill	IBM @Almaden@Research@Center,@San@ose,@CA:95120,@USA	rjprill@us.ibm.com		
Markus₽	Ringnér	Department®f10ncology,1Clinical®ciences,1Lund1University,1Lund,1Sweden12	markus.ringner@med.lu.se		
Richard 8.2	Savage	SystemsBiologyCentre,IIniversityCflWarwick,IInitedKingdom	r.s.savage@warwick.ac.uk		
Ben	Silva	Department the statistics, #lorida State University, Tallahassee, #L B2306-4330, USA	bsilva@stat.fsu.edu		
Piotr	Sobczyk	ICM,@university@bf@Warsaw,@Poland	sobbombo@gmail.com		
Artem	Sokolov	Baskin&chool@fEngineering,IUniversity@ffCalifornia,ISantaCruz,ISantaCruz,ICA2 95064,IUSA	sokolov@soe.ucsc.edu		
Emmett	Sprecher	Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept Dept 	emmett.sprecher@yale.edu		
Ioannis	Tsamardinos	(1) Bioinformatics狙aboratory,通nstitute函f重Computer低cience改革oundation值or집 Research馌nd团echnology④FORTH)函、即lastira狙00回/assilika鄧/outon,亟R-700週3団 Heraklion,亚Crete,亟reec眞2) Department찝f重Computerเዄcience,迅niversity亟f重Crete P.O.Box遼208,亟R-714釦9础eraklion,亚Crete,亟reece	tsamard@ics.forth.gr		
Jean-Philipp	Vert	(1)℃entreੴor℃omputational।避iology週℃BIO,弧Mines理arisTech,還5當ue巻aint-Honoré,還 Fontainebleau,正-77300運rance電2)個nstitut電Curie,涩6蹭ue置''Ulm,理aris,正-75248運rance (3)個NSERM,펞900,理aris,正-75248運rance	jean-philippe.vert@mines.org		
Yi⊠Kan⊠	Wang	Auckland Bioengineering Institute, I hell niversity Bf Auckland, New Zealand	yikan.wang@auckland.ac.nz		
ICharlesID.	Warden	BioinformaticsICore,IDepartmentIDfIMolecularIMedicine,ICityIDfIHopeINationalI MedicalICenter,IDuarte,ICA,ID1010,IUSA	cwarden@coh.org		
Xiaohui	Xie	Department@ftComputertScience,@university@ftCalifornia,Irvine,@rvine,#CA1926972 USA	xhx@ics.uci.edu		