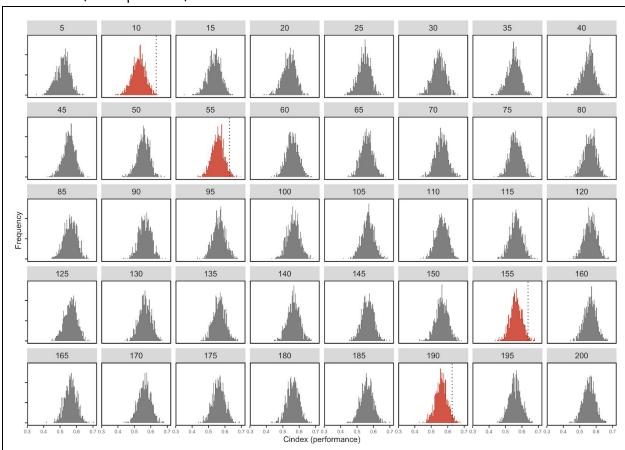
CoRaS: Cox proportional hazards and random survival forest methods for identifying drug repurposing candidates to treat breast cancer.

Pradeep Mangalath¹, Stan Finkelstein², Artem Sokolov³

Supporting Information.

LINCS results - cox-proportional hazards (CPH) approach.

Combined (1097 patients)



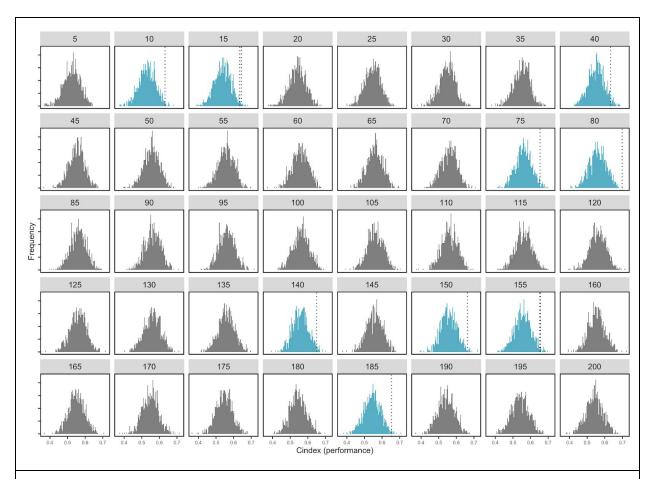
CPH combined results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

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² Massachusetts Institute of Technology, Cambridge, Massachusetts, United States

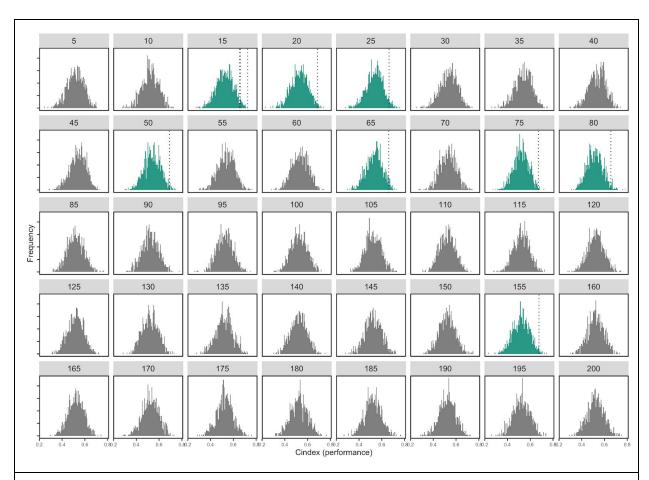
³ Laboratory of Systems Pharmacology, Harvard Medical School, Boston, Massachusetts, United States

ER+ (808 patients)



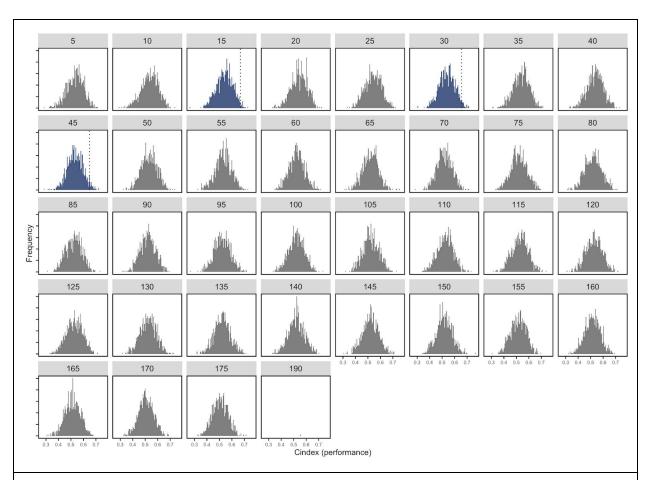
CPH ER+ results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

Her2 (343 patients)



CPH Her2 results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

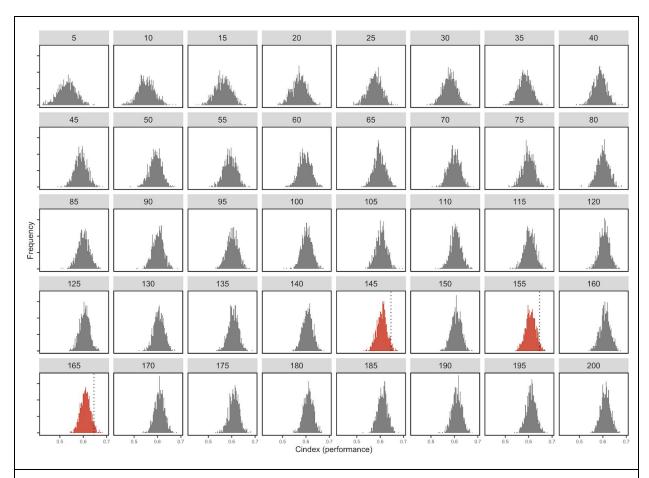
Triple Negative (247 patients)



CPH Triple negative results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

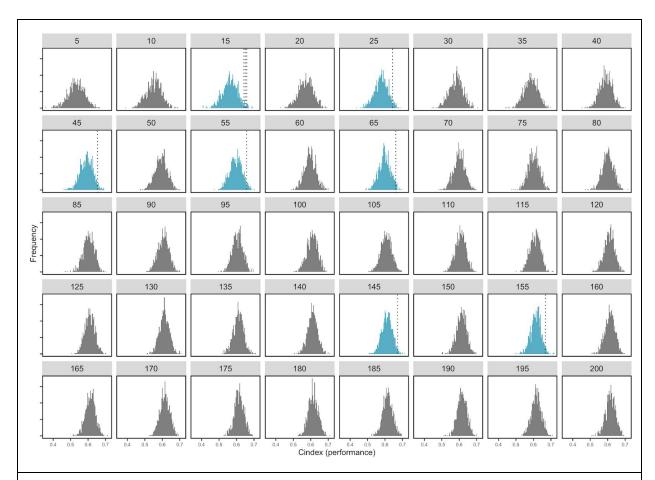
LINCS results - Random survival forests (RSF) approach.

Combined (1097 patients)



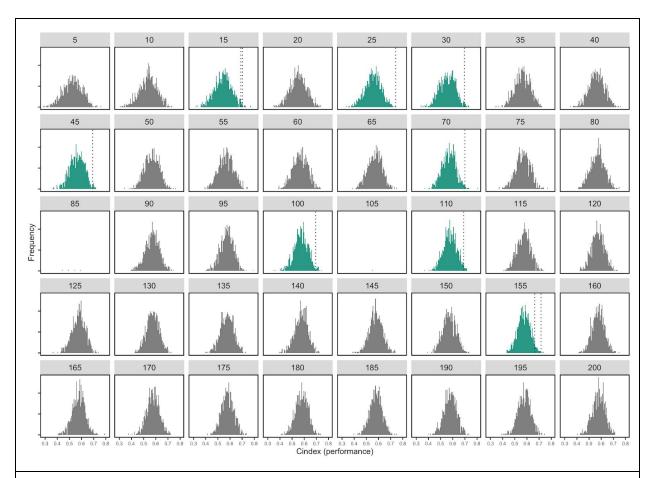
RSF combined results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

ER+ (808 patients)



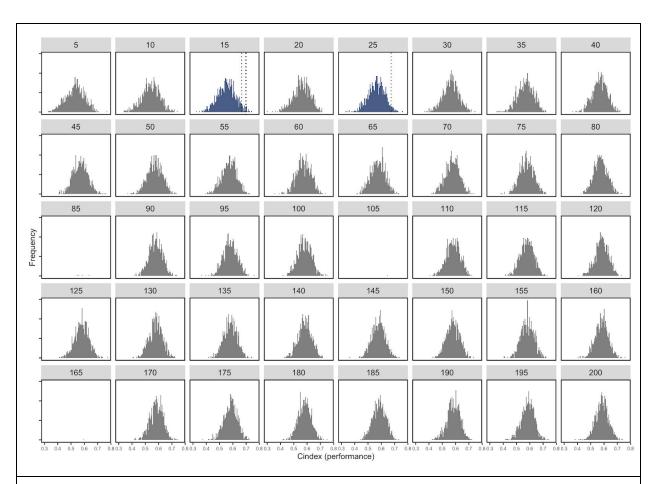
RSF ER+ results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

Her2 (343 patients)



RSF Her2 results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

Triple Negative (247 patients)



RSF Triple negative results | Significant drugs were identified by comparing 1000 random gene signature models of signature lengths n by bins sized in increments of 5 genes, to drug models of equal signature lengths. Significant drugs were identified by considering performance (c-index) below 5% threshold of empirical p-values, computed by comparing to the background distribution. Bins with significant drugs are highlighted in color. Dotted lines indicate drug performance (c-index).

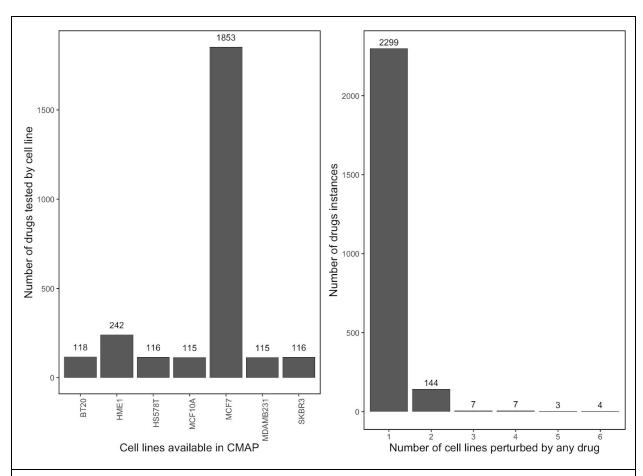
LINCS Results - List of significant drugs using RSF approach

Orug	Target	Approval	Combined	Er	Her2	Tneg
Alvocidib	CDK1	FDA	N	Υ	Y	N
AT9283	AURKA/B-JAK2	Experimental	N	N	Υ	N
AZ-628	BRAF	Experimental	N	N	N	Υ
AZD 5438	CDK2	Experimental	N	N	Υ	N
BMS-536924	IGF1R	Experimental	N	Υ	N	N
CHIR-99021	GSK3B	Experimental	N	N	Υ	N
Crizotinib	c-Met	FDA	Υ	Υ	N	N
Dovitinib	FLT3	FDA	Υ	N	N	N
Epigallocatechin gallate	PP1A	FDA	N	Υ	N	N
Gefitinib	EGFR	FDA	Υ	N	Υ	N
GSK 690693	AKT1	Experimental	N	N	Υ	N
GSK1059615	PIK3CA	Experimental	N	N	Υ	N
HG-6-64-01	ABL1	Experimental	N	N	N	Υ
Motesanib	VGFR1	FDA	N	Υ	N	N
Neratinib	ERBB2	FDA	N	N	Υ	N
PF-06463922	ROS1	Experimental	N	Υ	N	N
PI103	PIK3CA	Experimental	N	N	Υ	N
PRT062607	SYK	Experimental	N	Υ	Υ	N
Regorafenib	c-Kit	FDA	N	N	N	Υ
S-Ruxolitinib		FDA	N	Υ	N	N
Trichostatin A	HDAC1	FDA	N	N	N	Υ
VX-11e	ERK	Experimental	N	Υ	N	N

LINCS Results - List of significant drugs using CPH approach

Drug	Target	Approval	Combined	Er	Her2	Tneg
Afatinib	ERBB2	FDA	N	Υ	N	N
Alvocidib	CDK1	FDA	Υ	Υ	Υ	N
AZD 5438	CDK2	Experimental	N	N	Υ	N
Baicalein	CP2C9	FDA	N	N	Υ	N
BI-2536	PLK1	Experimental	N	Υ	Υ	N
BMS-536924	IGF1R	Experimental	Υ	N	N	N
Bosutinib	Src	FDA	Υ	N	N	N
Cediranib	VEGF	FDA	N	Υ	N	N
Gefitinib	EGFR	FDA	N	Υ	N	N
Go 6976	PKC	Experimental	N	N	Υ	N
GSK1059615	PIK3CA	Experimental	N	N	Υ	N
HG-9-91-01	SIK1	Experimental	N	N	Υ	N
LY2090314	GSK3	Experimental	N	N	Υ	N
Masitinib	c-Kit	FDA	N	N	N	Υ
Olomoucine II	CDK1	FDA	N	Υ	N	N
PF-06463922	ROS1	Experimental	N	Υ	N	N
PI103	PIK3CA	Experimental	N	N	N	Υ
PRT062607	SYK	Experimental	N	N	Υ	N
Regorafenib	c-Kit	FDA	N	N	N	Υ
SB 216763	GSK3B	Experimental	Υ	Υ	N	N
Sotrastaurin	PKC	FDA	N	Υ	N	N
XMD16-144	AURKA	Experimental	N	N	Υ	N
Y-27632	ROCK1	Experimental	N	Υ	N	N
ZM-447439	AURKA	Experimental	N	Υ	N	N

cMAP drug instances by cell line



We considered only drugs that were tested on breast cancer (BRCA) cell lines in cMAP. There were 7 such cell lines. Distribution of drugs tested on each cell line is shown in the figure on the left. MCF7 (ER+) cell line had the greatest number of drugs tested. A distribution of number of drugs tested in multiple cell lines are shown on the figure on the right. Even though 2299 drugs were tested on at least 1 cell line, this number includes replicates. There were 1711 unique drugs in total.

List of LINCS drugs available in CMAP drug instance list.

Link to excel file.

List of FDA approved drugs for breast cancer.

Link to excel file.