

Figure 1.





Figure 3.



intrinsic subgroup coding

# b

GADD45 Signaling

ATM Signaling

BER pathway

p53 Signaling

HIPPO signaling

■ positive z-score □ z-score = 0 ■ negative z-score ■ no activity pattern available



=indicates the threshold of adjusted p-value < 0.05

### Figure 4



- \* Number of patients' data used in the study from
- no presurgical treatment group, n = 56
- 2 week's presurgical AI group : n = 198 (157 + 41)

Additional file 3: Figure S1





POETIC HER2- (n=155) baselineESR1 residual Ki67



	baseline.ESR1
	VS.
	residual.Ki67
Spearman r	
r	-0.1558
95% confidence interval	-0.3102 to 0.00657
P value	
P (two-tailed)	0.0529
P value summary	ns
Exact or approximate P value?	Approximate
Significant? (alpha = 0.05)	No
Number of XY Pairs	155

С POETIC HER2+ (n=23) baselineESR1 change in Ki67



	vs. change.Ki67									
Spearman r										
r	-0.6077									
95% confidence interval	-0.8199 to -0.2487									
P value										
P (two-tailed)	0.0021									
P value summary	**									
Exact or approximate P value?	Approximate									
Significant? (alpha = 0.05)	Yes									
Number of XY Pairs	23									

POETIC HER2+ (n=23) baselineESR1 residual Ki67

d



	baseline.ESR1
	VS.
	residual.Ki67
Spearman r	
r	-0.6156
95% confidence interval	-0.824 to -0.2605
P value	
P (two-tailed)	0.0018
P value summary	**
Exact or approximate P value?	Approximate
Significant? (alpha = 0.05)	Yes
Number of XY Pairs	23

### Additional file 5: **Figure S3**



Additional file 6: Figure. S4





Additional file 7: Figure S5b

Pathways		CONEI	BRCA1	CCNE2	CDK1	<b>CCND1</b>	<b>CHEK1</b>	E2F2 F2F5	CCNB1	CDC25A	CDC25C	CDK6 skp3	CCNR2	CCND2	PPM1J sen	TGFB3	MYC	PCNA	PLKI	CCNA2	FANCD2	RFC5,SLC19A1	AURKA	BLM CDKN2D	FAS	FOX01	FZD7	HDAC11	KA12B MSH6	PKMYT1	PRKCA Smarca4	SUV39H1	TGFBR2
	<b>⊯</b> 1	6 12	2 11	11	11	10	9	99	8	7	7	7 7	6	6	6 6	6	5	5 :	5 5	4	4	4	3	3 3	3	3	3	3 3	3 3	3	3 3	3	3
Cell Cycle: G2/M DNA Damage Checkpoint Regulation	12																																
Cell Cycle: G1/S Checkpoint Regulation	17																		H														
Cyclins and Cell Cycle Regulation	19										1					-								-			1						1
Estrogen-mediated S-phase Entry	11			_																-													
Mitotic Roles of Polo-Like Kinase	8												L					14															
Cell Cycle Control of Chromosomal Replication	2					_								_		_																	
Aryl Hydrocarbon Receptor Signaling	12																									1							
GADD45 Signaling	9																						_						_				
Role of BRCA1 in DNA Damage Response	10									_							_																
Role of CHK Proteins in Cell Cycle Checkpoint Control	12																												_	-	_		
Hereditary Breast Cancer Signaling	14													_																			
DNA damage-induced 14-3-3σ Signaling	8		_		_																		_										
ATM Signaling	10											_							_														
Antiproliferative Role of TOB in T Cell Signaling	8								_	_																				_			
Molecular Mechanisms of Cancer	24																												_				
Mismatch Repair in Eukaryotes	3																																
BER pathway	1			_												_												_					
p53 Signaling	9																																
Cell Cycle Regulation by BTG Family Proteins	7																																
Small Cell Lung Cancer Signaling	12																																
Breast Cancer Regulation by Stathmin1	8																																
Glioblastoma Multiforme Signaling	9																																
Factors Promoting Cardiogenesis in Vertebrates	8											_																					
Role of Oct4 in Mammalian Embryonic Stem Cell Pluripotency	1																																
HIPPO signaling	4																																
					-	-	-			_	_		-				-									_	_		_				_

## Additional file 9: Figure S6



### Additional file 10: Figure S7a



0

-1

### Additional file 10: **Figure S7b**

2



0

-1

### Additional file 11: **Figure S8a**

1



### Additional file 11: Figure S8b



### Additional file 12: Figure S9