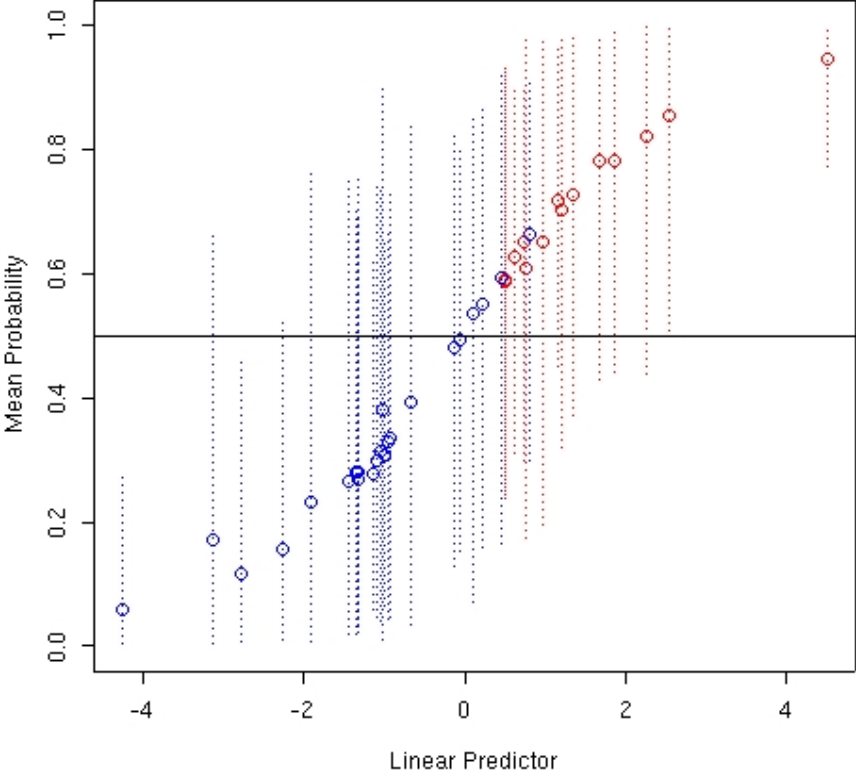
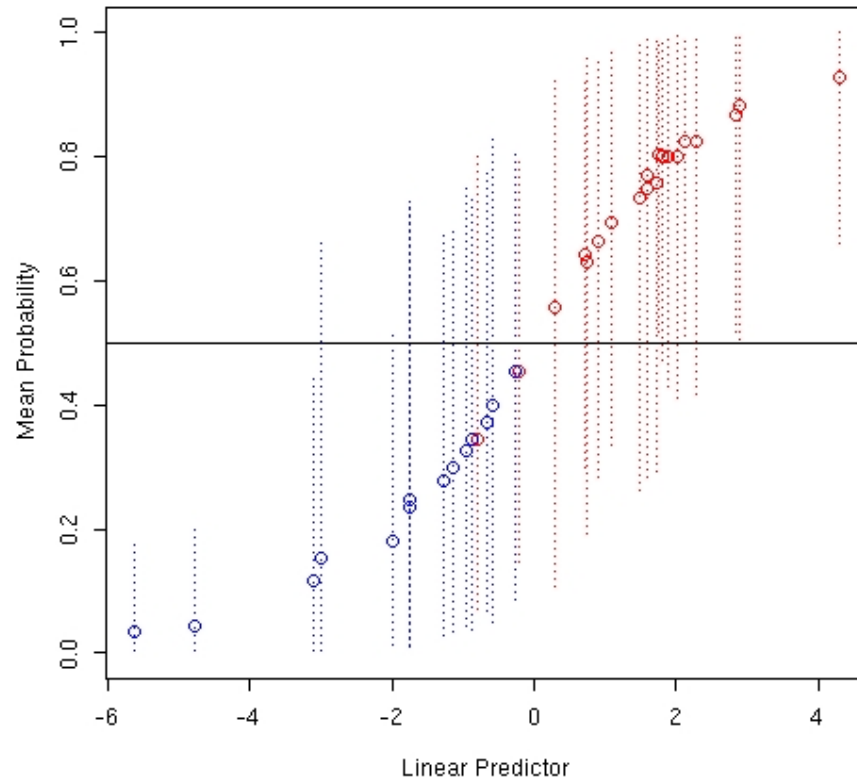


Supplementary Figure 1



Supplementary Figure 2



# Supplementary Figure 3

## IBC

	# Genes	False Disc. Rate	p-value	Bayes Factor
1. GO:0030205 [8]: dermatan sulfate metabolism	1 <a href="#">[show]</a>	1.91e <sup>-2</sup>	1.11e <sup>-8</sup>	4
2. GO:0030208 [9]: dermatan sulfate biosynthesis	1 <a href="#">[show]</a>	1.91e <sup>-2</sup>	1.11e <sup>-8</sup>	4
3. GO:0030206 [8]: chondroitin sulfate biosynthesis	1 <a href="#">[show]</a>	2.44e <sup>-2</sup>	1.64e <sup>-7</sup>	3
4. GO:0050651 [8]: dermatan sulfate proteoglycan biosynthesis	1 <a href="#">[show]</a>	2.44e <sup>-2</sup>	1.64e <sup>-7</sup>	3
5. GO:0030204 [7]: chondroitin sulfate metabolism	1 <a href="#">[show]</a>	2.44e <sup>-2</sup>	1.64e <sup>-7</sup>	3
6. GO:0050650 [7]: chondroitin sulfate proteoglycan biosynthesis	1 <a href="#">[show]</a>	2.44e <sup>-2</sup>	3.49e <sup>-7</sup>	2
7. GO:0050655 [7]: dermatan sulfate proteoglycan metabolism	1 <a href="#">[show]</a>	2.44e <sup>-2</sup>	3.49e <sup>-7</sup>	2
8. GO:0050654 [6]: chondroitin sulfate proteoglycan metabolism	1 <a href="#">[show]</a>	2.68e <sup>-2</sup>	5.90e <sup>-7</sup>	2
9. GO:0007516 [5]: hemocyte development	1 <a href="#">[show]</a>	3.66e <sup>-2</sup>	1.85e <sup>-6</sup>	2
10. GO:0008202 [6]: steroid metabolism	2 <a href="#">[show]</a>	3.66e <sup>-2</sup>	2.36e <sup>-6</sup>	2
11. GO:0042386 [4]: hemocyte differentiation (sensu Arthropoda)	1 <a href="#">[show]</a>	3.66e <sup>-2</sup>	2.50e <sup>-6</sup>	2
12. GO:0006024 [8]: glycosaminoglycan biosynthesis	1 <a href="#">[show]</a>	3.79e <sup>-2</sup>	4.07e <sup>-6</sup>	1
13. GO:0006023 [7]: aminoglycan biosynthesis	1 <a href="#">[show]</a>	3.79e <sup>-2</sup>	5.17e <sup>-6</sup>	1
14. GO:0046209 [5]: nitric oxide metabolism	1 <a href="#">[show]</a>	3.79e <sup>-2</sup>	6.43e <sup>-6</sup>	1
15. GO:0006809 [5]: nitric oxide biosynthesis	1 <a href="#">[show]</a>	3.79e <sup>-2</sup>	6.43e <sup>-6</sup>	1
16. GO:0030166 [7]: proteoglycan biosynthesis	1 <a href="#">[show]</a>	3.79e <sup>-2</sup>	6.43e <sup>-6</sup>	1
17. GO:0042133 [5]: neurotransmitter metabolism	1 <a href="#">[show]</a>	5.29e <sup>-2</sup>	1.59e <sup>-5</sup>	1
18. GO:0006029 [7]: proteoglycan metabolism	1 <a href="#">[show]</a>	5.80e <sup>-2</sup>	2.25e <sup>-5</sup>	1
19. GO:0044272 [6]: sulfur compound biosynthesis	1 <a href="#">[show]</a>	6.57e <sup>-2</sup>	3.39e <sup>-5</sup>	1
20. GO:0030203 [7]: glycosaminoglycan metabolism	1 <a href="#">[show]</a>	7.16e <sup>-2</sup>	4.75e <sup>-5</sup>	0

## Hypoxia

	# Genes	False Disc. Rate	p-value	Bayes Factor
1. GO:0006263 [7]: DNA-dependent DNA replication	2 <a href="#">[show]</a>	4.05e <sup>-3</sup>	1.47e <sup>-9</sup>	5
2. GO:0006262 [7]: DNA-dependent DNA replication	2 <a href="#">[show]</a>	4.05e <sup>-3</sup>	1.47e <sup>-9</sup>	5
3. GO:0006261 [7]: DNA-dependent DNA replication	2 <a href="#">[show]</a>	4.05e <sup>-3</sup>	2.79e <sup>-9</sup>	4
4. GO:0006988 [4]: cellular defense response	2 <a href="#">[show]</a>	4.14e <sup>-3</sup>	5.69e <sup>-9</sup>	4
5. GO:0051085 [9]: chaperone cofactor dependent protein folding	1 <a href="#">[show]</a>	6.30e <sup>-3</sup>	3.81e <sup>-8</sup>	3
6. GO:0051084 [8]: posttranslational protein folding	1 <a href="#">[show]</a>	6.30e <sup>-3</sup>	3.81e <sup>-8</sup>	3
7. GO:0006260 [6]: DNA replication	2 <a href="#">[show]</a>	6.60e <sup>-3</sup>	6.25e <sup>-8</sup>	3
8. GO:0006268 [8]: DNA unwinding	1 <a href="#">[show]</a>	7.88e <sup>-3</sup>	1.63e <sup>-7</sup>	3
9. GO:0006265 [6]: DNA topological change	1 <a href="#">[show]</a>	7.88e <sup>-3</sup>	1.63e <sup>-7</sup>	3
10. GO:0000067 [6]: DNA replication and chromosome cycle	2 <a href="#">[show]</a>	8.00e <sup>-3</sup>	2.15e <sup>-7</sup>	3
11. GO:0006955 [4]: immune response	3 <a href="#">[show]</a>	1.32e <sup>-2</sup>	8.70e <sup>-7</sup>	2
12. GO:0006270 [8]: DNA replication initiation	1 <a href="#">[show]</a>	1.32e <sup>-2</sup>	1.04e <sup>-6</sup>	2
13. GO:0009611 [5]: response to wounding	2 <a href="#">[show]</a>	1.35e <sup>-2</sup>	1.60e <sup>-6</sup>	2
14. GO:0010986 [5]: cellular defense response (sensu Vertebrata)	1 <a href="#">[show]</a>	1.35e <sup>-2</sup>	1.67e <sup>-6</sup>	2
15. GO:0006952 [5]: defense response	3 <a href="#">[show]</a>	1.35e <sup>-2</sup>	1.83e <sup>-6</sup>	2
16. GO:0009607 [4]: response to biotic stimulus	3 <a href="#">[show]</a>	1.81e <sup>-2</sup>	4.17e <sup>-6</sup>	1
17. GO:0009613 [5]: response to pest, pathogen or parasite	2 <a href="#">[show]</a>	2.83e <sup>-2</sup>	1.51e <sup>-5</sup>	1
18. GO:0000082 [7]: G1/S transition of mitotic cell cycle	1 <a href="#">[show]</a>	2.83e <sup>-2</sup>	1.52e <sup>-5</sup>	1
19. GO:0043207 [5]: response to external biotic stimulus	2 <a href="#">[show]</a>	2.99e <sup>-2</sup>	1.96e <sup>-5</sup>	1
20. GO:0006259 [5]: DNA metabolism	2 <a href="#">[show]</a>	3.19e <sup>-2</sup>	2.55e <sup>-5</sup>	1

## Lymph Node Persistence

	# Genes	False Disc. Rate	p-value	Bayes Factor
1. GO:0050789 [2]: regulation of biological process	11 <a href="#">[show]</a>	7.78e <sup>-3</sup>	2.74e <sup>-10</sup>	5
2. GO:0019222 [4]: regulation of metabolism	9 <a href="#">[show]</a>	7.78e <sup>-3</sup>	1.44e <sup>-9</sup>	5
3. GO:0050791 [3]: regulation of physiological process	10 <a href="#">[show]</a>	7.78e <sup>-3</sup>	1.63e <sup>-9</sup>	5
4. GO:0030155 [4]: regulation of cell adhesion	2 <a href="#">[show]</a>	8.42e <sup>-3</sup>	3.92e <sup>-9</sup>	4
5. GO:0008892 [5]: negative regulation of metabolism	3 <a href="#">[show]</a>	8.42e <sup>-3</sup>	9.33e <sup>-9</sup>	4
6. GO:0007275 [2]: development	8 <a href="#">[show]</a>	8.42e <sup>-3</sup>	1.02e <sup>-8</sup>	4
7. GO:0016322 [6]: neuronal remodeling	1 <a href="#">[show]</a>	8.42e <sup>-3</sup>	1.52e <sup>-8</sup>	4
8. GO:0045449 [6]: regulation of transcription	8 <a href="#">[show]</a>	8.42e <sup>-3</sup>	2.20e <sup>-8</sup>	4
9. GO:0019219 [5]: regulation of nucleobase, nucleoside, nucleotide and nucleic acid metabolism	8 <a href="#">[show]</a>	8.42e <sup>-3</sup>	2.61e <sup>-8</sup>	3
10. GO:0006139 [5]: nucleobase, nucleoside, nucleotide and nucleic acid metabolism	10 <a href="#">[show]</a>	8.42e <sup>-3</sup>	3.13e <sup>-8</sup>	3
11. GO:0006350 [6]: transcription	8 <a href="#">[show]</a>	9.09e <sup>-3</sup>	4.64e <sup>-8</sup>	3
12. GO:0006348 [5]: loss of chromatin silencing	1 <a href="#">[show]</a>	1.02e <sup>-2</sup>	8.21e <sup>-8</sup>	3
13. GO:0009790 [3]: embryonic development	2 <a href="#">[show]</a>	1.02e <sup>-2</sup>	8.99e <sup>-8</sup>	3
14. GO:0006357 [8]: regulation of transcription from Pol II promoter	3 <a href="#">[show]</a>	1.16e <sup>-2</sup>	1.41e <sup>-7</sup>	3
15. GO:0000578 [5]: embryonic axis specification	1 <a href="#">[show]</a>	1.36e <sup>-2</sup>	4.63e <sup>-7</sup>	2
16. GO:0009952 [4]: anterior/posterior pattern formation	1 <a href="#">[show]</a>	1.36e <sup>-2</sup>	4.63e <sup>-7</sup>	2
17. GO:0007350 [4]: blastoderm segmentation	1 <a href="#">[show]</a>	1.36e <sup>-2</sup>	4.63e <sup>-7</sup>	2
18. GO:0007351 [5]: regional subdivision	1 <a href="#">[show]</a>	1.36e <sup>-2</sup>	4.63e <sup>-7</sup>	2
19. GO:0009948 [5]: anterior/posterior axis specification	1 <a href="#">[show]</a>	1.36e <sup>-2</sup>	4.63e <sup>-7</sup>	2
20. GO:0008592 [6]: determination of anterior/posterior axis, embryo	1 <a href="#">[show]</a>	1.36e <sup>-2</sup>	4.63e <sup>-7</sup>	2

**Supplementary Table 1. Summary of clinical properties of KF-SYSCC breast cancer samples.**

<b>Clinical Variable</b>	<b>No. Samples</b>
Age	
< 40	47
> 40	111
Path Stage	
1A	28
2A	29
2B	48
3A	44
3B	9
Tumor Size	
< 2 cm	55
2 – 5 cm	93
> 5 cm	10
Lymph Nodes	
0	41
1 – 3	71
> 4	46
Survival	
Relapse (avg 24.2 mo)	69
No relapse (avg. 60.0 mo)	89
ER	
Positive	110
Negative	48
Treatment	
None	7
Hormone only	31
Chemo only	45
Hormone & Chemo	75

**Supplementary Table 2. Summary of clinical properties of Duke breast cancer samples.**

<b>Clinical Variable</b>	<b>No. Samples</b>
Age	
< 40	10
> 40	91
Path Stage	
1	30
2A	40
2B	27
3A	2
4	1
unk	1
Tumor Size	
< 2 cm	36
2 – 5 cm	62
> 5 cm	3
Lymph Nodes	
0	56
1 – 3	28
> 4	17
Recurrence	
Relapse (avg 42.7 mo)	26
No relapse (avg. 92.5 mo)	75
ER	
Positive	37
Negative	64
Treatment	
None	16
Hormone only	30
Chemo only	36
Hormone & Chemo	19